

# Enzyme-assisted extraction of phlorotannins from Sargassum and biological activities

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# Enzyme-assisted extraction of phlorotannins from Sargassum and biological activities



# A Dissertation Submitted by: Maya Puspita 26010112510005

Presented to the Doctoral Program of Coastal Resources

Management in partial fulfillment of the requirement for the

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Extraction assistée par enzyme provenant d'algues brunes du genre *Sargassum* et les activités biologiques

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#### **DECLARATION OF ORIGINALITY**

I, undersigned below, declare that this doctoral dissertation entitled "Enzyme-assisted extraction of *Sargassum* phlorotannins and biological activities" is my own work, is not a copy from any scientific writing and has not been published elsewhere. Information derived from the published and unpublished work of others has been acknowledged in the text and a list of references is given in the bibliography.

This dissertation has been written based on the directions of my supervisors and co-supervisors as one of the requirements to obtain the doctoral degree of the Double Degree Program at Doctoral Program of Coastal Resources Management, Faculty of Fisheries and Marine Science, Diponegoro University. This work was under the joint supervisor between University of South Brittany, Vannes and Diponegoro University, Semarang. If a part of this dissertation has been proven to be a copy from other works or the originality of my work is questioned, I am willing to accept any sanctions according to the applicable regulations.

Semarang, ...... 2017

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# Extraction assistée par enzyme de phlorotannins provenant d'algues brunes du genre *Sargassum* et les activités biologiques

#### Résumé:

En France et en Indonésie, *Sargassum* est une macroalgue abondante mais cette ressource n'est pas encore exploitée. Les composés de l'algue *Sargassum* sont de nature chimique très diverse et certains possèdent des activités biologiques parmi lesquels les phlorotannins. Il est donc intéressant des voies de valorisation dans des domaines tels que l'industrie alimentaire, pharmaceutique, et cosmétique. Par conséquent, la connaissance de la composition biochimique de *Sargassum* qui représente aussi sa valeur nutritionnelle est importante. L'extraction solide-liquide est la technique la plus utilisée pour l'extraction des phlorotannins. L'efficacité de l'extraction est néanmoins limitée par l'accessibilité aux composés, difficulté due à la complexité structurale de la paroi des algues. En conséquence, il est important de considérer la dégradation de la paroi pour favoriser l'extraction des phlorotannins. L'utilisation d'enzymes comme outil technologique est un processus qui pourrait faciliter cette dégradation.

Cette étude a eu trois objectifs principaux; (1) la caractérisation de la composition biochimique de *Sargassum* et l'étude de la variabilité saisonnière, (2) l'extraction solide-liquide des phlorotannins et (3) l'optimisation des conditions d'extraction assistée par enzymes des phlorotannins et l'étude des activités biologiques des extraits produits. Les activités analysées concernent les activités antioxydants (piégeage de radicaux DPPH et pouvoir réducteur FRAP), les activités cosmétiques de blanchiment de la peau (anti-tyrosinase), les activités anti-âge (anti-élastase), les activités antibactériennes, l'inhibition de la formation de biofilm, la mesure de la cytotoxicité et des activités antivirales.

Dans cette étude, les minéraux représentent d'une valeur importance avec une teneur qui atteint 33% suivi par les protéines avec 24% de la matière sèche. Les sucres ne représentent que 13% de la matière sèche. L'étude de la variabilité saisonnière montre que la composition biochimique de *Sargassum*, comme pour la plupart des algues, est fortement affectée par la saison de collecte et semble donc liée au cycle de vie. Ce résultat est également confirmé pour les composés phénoliques.

Cependant des méthodes d'extraction adaptées permettent d'augmenter le rendement. L'extraction solide-liquide conduit à des rendements plus faibles (5 à 24% de la matière sèche de l'algue) comparée à l'extraction solide-liquide assistée par enzyme (26 à 38% de la matière sèche de l'algue). En ce qui concerne la teneur en polyphénols, l'extraction assistée par enzymes permet aussi d'obtenir un meilleur rendement 6% contre 4% de la matière sèche si l'extraction est réalisée sans enzyme.

Après le criblage de l'action de huit enzymes pour extraire les polyphénols de *S. muticum*, Viscozyme, un cellulase, libère une plus grande quantité de composés phénoliques d'après les teneurs déterminées par la méthode de Folin Ciocalteu, résultat confirmé par l'activité anti-radicalaire de l'extrait obtenu

 $(IC_{50}\ 0.6 \pm 0.1\ mg\ mL^{-1})$ . Cet extrait montre la plus forte activité anti-biofilm avec  $43.7 \pm 4.2\%$  d'inhibition de la formation de biofilm de *P. aeruginosa*.

Tous les extraits produits montrent des activités biologiques. L'activité FRAP la plus importante a été montrée par l'extrait produit avec l'enzyme AMG, un exo-amylase, ( $60.8 \pm 1.8 \, \mu M$  of Fe<sup>2+</sup>). L'extrait obtenu avec l'Ultraflo – un xylanase, ( $24.5 \pm 2.6\%$  d'inhibition) possède la plus forte activité de blanchiment (anti tyrosinase), tandis que l'extrait de Shearzyme, cellulase et xylanase, ( $30.9 \pm 5.9\%$  d'inhibition) présente une activité anti-âge plus élevée que les autres extraits.

L'utilisation d'enzymes pour l'extraction des composés phénoliques augmete le rendement d'extraction. De plus, les extraits enzymatiques de *Sargassum* démontrent des activités biologiques intéressantes. Il est important de considérer qu'en utilisant l'extraction assistée par enzyme, d'autres composés que les polyphénols sont co-extraits. En conséquence, il sera intéressant de purifier les extraits pour déterminer la nature des molécules actives. En outre, la caractérisation complète des composés phénoliques de *Sargassum* pourrait aider à mieux comprendre les mécanismes d'action liés aux activités biologiques observées.

Mots-clés: composition biochimique, variation saisonnières, polyphénols, Sargassum, activités biologiques

# Enzyme-assisted of phlorotannins from Sargassum and biological activities

#### **Resume:**

In France and Indonesia, Sargassum is quite abundant yet this alga has not been optimally exploited. Sargassum contains macro and micro molecules and also bioactive compounds, the most known is phlorotannins, showing a wide range of pharmacological properties. Therefore, it is interesting to further valorize this alga for various industrial applications, such as food, pharmacology and cosmetic. Knowledge on biochemical composition of this alga, which also represents its nutritive value, is essential for food industries. Concerning the bioactive compound, solid-liquid extraction is the most commonly used technique in the extraction of phlorotannins from brown algal biomass. However, phlorotannins of brown algae are known to be bound with algal cell wall and the high content of various cell wall polysaccharides in macroalgae limit the active compound accessibility. Thus, these cell walls strongly reduced the extraction efficiency application of conventional extraction. As a consequence, it is important to consider the degradation of algal cell wall as a way to liberate more phlorotannins. Application of enzymes is considered as one process that might facilitate the degradation of algal cell wall leading to the recovery of phlorotannins from brown algae.

This study had three main subjects of interest; they were (1) the characterization of *Sargassum* biochemical composition and its seasonal variation, (2) solid-liquid extraction and (3) enzyme-assisted extraction of *Sargassum* phlorotannins and their bioactivities. The bioactivities analyzed in this study included antioxidant (DPPH radical scavenging and FRAP reducing power), antibacterial, tyrosinase, elastase, biofilm inhibition, cytotoxicity and antiviral activity.

In this study, minerals represented a significant value with a content reaching 33% followed by proteins with 24% of algal dry material. In the contrary, sugars showed only 13% of algal dry material. The study of seasonal variability demonstrated that the biochemical composition of *Sargassum*, as for most algae, is strongly affected by the season and it appears to be related to the life cycle of the algae. Same cases happened with the polyphenol compound of *Sargassum*.

Nevertheless, suitable extraction method is important to choose in order to increase the yield. Solid-liquid extraction yielded lower (5-24% of algal dry material) compared to enzyme-assisted solid-liquid extraction (21-38% of algal dry material). Regarding the polyphenol content, enzyme-assisted extraction also yielded a better yield, 32%, compared to 26% of algal dry material if the extraction is carried out without enzyme.

After the screening of eight enzymes to extract the polyphenols from *S. muticum*, Viscozyme - a cellulase, released a greater quantity of phenolic compounds according to Folin Ciocalteu method, result was also confirmed by the anti-free radical activity of (IC<sub>50</sub>  $0.6 \pm 0.1$  mg mL<sup>-1</sup>). This extract also managed to

prevent  $43.7 \pm 4.2\%$ , higher than the other extracts, of biofilm formation generated by *P. aeruginosa*.

Other than antiradical activity, it seems that each enzyme showed different potential. In reducing power, the most important activity was shown by AMG (an exo-amylase) extract ( $60.8 \pm 1.8 \, \mu M$  of Fe<sup>2+</sup>). Ultraflo (a xylanase) extract had promising potential in inhibiting the tyrosinase at  $24.5 \pm 2.6\%$ , while Shearzyme (cellulose and xylanase) extract showed higher inhibition elastase activity than the other extracts,  $30.9 \pm 5.9\%$ . Viscozyme extract was able to hamper  $43.7 \pm 4.2\%$  of film formation generated by *P. aeruginosa*. However, Shearzyme extracts exhibited strong biofilm inhibition activity against the *E. coli*,  $60.3 \pm 10.3\%$  of inhibition.

The application of enzymes in the extraction of the Sargassum phenolic compound is quite effective. In addition, *Sargassum* enzyme extracts demonstrate a wide range of bioactivity. It is important to consider that, using enzymes, the phenolic compound is probably not the only compound extracted from the algal matrix by the enzymes. As a result, it will be interesting to continue working on the purified samples pretreated by enzyme-assisted extraction in the future. In addition, the characterization of the *Sargassum* phenolic compound could help to better understand the mechanisms of their bioactivity.

Keywords: biochemical composition, seasonal variation, polyphenols, Sargassum, biological activities

#### Ekstraksi enzimatik senyawa florotannin dari alga coklat Sargassum dan aktivitas biologi

#### Ringkasan:

Di Perancis dan Indonesia, Sargassum merupakan spesies yang cukup melimpah namun alga ini belum dimanfaatkan secara optimal. Sargassum mengandung makro dan mikromolekul serta senyawa bioaktif, yang paling dikenal adalah florotanin, yang memiliki beragam sifat farmakologi. Oleh karena itu, Sargassum menjadi salah satu spesies alga coklat yang menarik untuk divalorisasi lebih lanjut untuk beragam aplikasi seperti industry makanan, farmakologi, dan kosmetik. Pengetahuan tentang komposisi biokimia dari Sargassum, yang juga mewakili kandungan nutrisi, merupakan satu hal yang penting bagi industri makanan. Selanjutnya, aplikasi senyawa bioaktif, ekstraksi padat-cair telah menjadi teknik yang paling umum digunakan dalam ekstraksi florotanin dari biomassa alga coklat. Akan tetapi, florotanin alga coklat dikenal dengan ikatan kompleksnya dengan polisakarida di dinding sel alga dan tingginya kandungan polisakarida membatasi aksesibilitas senyawa aktif dari matriks dinding sel. Oleh karena itu, keberadaan dinding sel ini mengurangi efisiensi ekstraksi senyawa bioaktif dari alga. Alhasil, penting untuk mempertimbangkan degradasi dinding sel alga sebagai cara untuk mengekstraksi florotanin lebih banyak. Aplikasi enzim dianggap sebagai salah satu proses yang dapat menguraikan dinding sel alga sehingga meningkatkan efisiensi ekstraksi florotanin.

Penelitian ini memiliki tiga tujuan utama, yaitu (1) karakterisasi komposisi biokimia *Sargassum* dan variasi musiman, (2) ekstraksi padat-cair (3) dan enzimatik senyawa florotanin *Sargassum* dan bioaktifitas. Bioaktifitas senyawa florotanin yang dianalisis dalam penelitian ini meliputi aktivitas antioksidan (antiradikal DPPH radikal dan FRAP), antibakteri, tirosinase, elastase, penghambatan biofilm, sitotoksisitas dan antivirus.

Selain dari polisakarida, tingginya kandungan abu menjadi komposisi dominan kedua di *Sargassum* karena mewakili hingga 35% dari berat kering. Selanjutnya, komposisi biokimia dari *Sargassum* sangat dipengaruhi oleh musim dan berhubungan dengan siklus hidup. Musim mungkin menjadi faktor yang paling mempengaruhi kandungan polifenol dalam *Sargassum*, tetapi metode ekstraksi yang sesuai menentukan efisiensi ekstraksi polifenol dari *Sargassum*.

Ekstraksi padat-cair menghasilkan berat kering lebih rendah (berkisar antara 5-24% dari berat kering) dari ekstraksi enzimatik (21-38% dari berat kering). Kandungan polifenol dari ekstrak enzimatik *Sargassum* memiliki kandungan yang lebih tinggi dari ekstraksi padat-cair.

Berdasarkan evaluasi penapisan enzim untuk mengekstrak polifenol *S. muticum*, di antara delapan enzim, Viscozyme mengekstrak senyawa polifenol lebih banyak dan mempunyai aktivitas antiradikal yang kuat. Hal ini mungkin berkaitan dengan kondisi optimum yang digunakan selama proses ekstraksi, yaitu 50 ° C selama 3 H. Oleh karena senyawa polifenol bersifat sensitif terhadap panas dan mudah teroksidasi oleh paparan cahaya, sangat disarankan untuk menerapkan

suhu yang sedang antara 40 sampai 60 ° C. Selain itu, ekstrak enzimatik dari polifenol dari *Sargassum* dianggap memiliki farmakologis menjanjikan, nutraceutical, dan sifat cosmeceutical.

Dalam studi ini, kandungan mineral dalam *Sargassum* mencapai 33% diikuti oleh protein dengan 24% dari total berat kering alga. Sebaliknya, kandungan gula menunjukkan hanya 13% dari total berat kering alga. Studi tentang variabilitas musiman menunjukkan bahwa komposisi biokimia dari *Sargassum* sangat dipengaruhi oleh musim dan berkaitan dengan siklus hidup dari alga, sama halnya dengan kandungan polifenol pada *Sargassum*.

Namun demikian, metode ekstraksi yang cocok penting untuk dipertimbangkan dalam rangka meningkatkan efisiensi ekstraksi. Ekstraksi padatcair menghasilkan lebih rendah (5-24% dari bahan kering alga) berat kering dibandingkan dengan ekstraksi enzimatik (26-38% dari bahan kering alga). Mengenai kandungan polifenol, ekstraksi enzimatik juga mengekstrak lebih banyak, 6%, dibandingkan dengan 4% dari total berat ekstrak jika ekstraksi dilakukan tanpa enzim.

Setelah skrining dengan menggunakan delapan enzim untuk mengekstrak polifenol dari *S. muticum*, Viscozyme - karbohidrase, membebaskan lebih banyak senyawa polifenol, dan menunjukan aktivitas antiradikal bebas yang kuat (IC<sub>50</sub> 0.6  $\pm$  0.1 mg mL<sup>-1</sup>). Ekstrak ini juga berhasil mencegah 43.7  $\pm$  4.2%, lebih tinggi dari ekstrak lainnya, pembentukan biofilm yang dihasilkan oleh *P. aeruginosa*. Ekstrak AMG mempunyai aktifitas reduksi besi atau FRAP pada 60.8  $\pm$  1,8 $\mu$ M dari Fe<sup>2+</sup>. Ekstrak Ultraflo memiliki potensi yang menjanjikan dalam menghambat tirosinase di 24,5  $\pm$  2,6%, sedangkan ekstrak Shearzyme menunjukkan aktivitas antielastase yang lebih tinggi dari ekstrak lainnya, 30,9  $\pm$  5,9%.

Aplikasi enzim dalam ekstraksi senyawa polifenol Sargassum cukup efektif. Selain itu, ekstrak enzimatik dari *Sargassum* menunjukkan bioaktivitas yang sangat luas. Namun demikian, sangatlah penting untuk diêrhatikan bahwa, dengan menggunakan enzim, senyawa polifenol mungkin bukan satu-satunya senyawa yang diekstrak dari matriks alga. Oleh karena itu, sangat disarankan untuk melanjutkan penelitian dengan menggunakan sampel yang dimurnikan. Selain itu, karakterisasi senyawa polifenol *Sargassum* bisa membantu untuk lebih memahami mekanisme bioaktivitas dari senyawa tersebut.

Kata kunci: komposisi biokimia, variasi musiman, polifenol, Sargassum, aktivitas biologi

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#### Introduction

Brown algae (Phaeophyceae) consist of 2038 species from 20 order in which order of Fucales contributes 597 species. *Sargassum*, known as gulf-weed or sea holly belonging to family Sargassaceae, contains 353 species that have been accepted taxonomically (Guiry and Guiry, 2015). *Sargassum* species are spread throughout the tropical and subtropical regions attached on the substrates with its holdfast. This species is characterized by its short stipe differentiating to numerous branches that have many leaf-like laterals and spherical vesicles as tool for flotation. Due to the presence of this floating vesicle, *Sargassum* often detaches from the substrates and forms vast pelagic mats as massive floating biomass as in the Sargasso Sea (Mattio and Payri, 2011). In Indonesia, there are 58 species recorded by Atmadja and Prud'homme van Reine (2014). One of *Sargassum* species found in France is *S. muticum*. However, this is not the native species from this region as it was imported unintentionally in European waters in 1970s (Josefsson and Jansson, 2011). *Sargassum* has been acknowledged for its massive influx of biomass along the coast (Milledge and Harvey, 2016).

Interesting fact of *Sargassum* species is that communities in some regions, for example in China and Japan, have noticed the important role of this species in pharmacology and food industry. It has been mentioned that *Sargassum*, particularly *S. fulvellum* (Turner) C. Agardh, *S. henslowianum* C. Agardh, *S. thunbergii* (Mertens ex Roth) Kuntze, and *S. horneri* (Turner) C. Agardh, has been part of Chinese traditional medicine from the ancient times and has been formulated in classical Chinese prescription. Meanwhile in Japan, *S. fusiforme* becomes a popular side dish boiled with water and served with the soy sauce (Liu *et al.*, 2012). Supposedly, the finding of ancient manuscript concerning medicinal uses of *Sargassum* has drawn the attention and made this species compelling for further study.

Recognition of *Sargassum* pharmacological properties with wide range of potential derived from its extracts or isolated pure components have unveiled the remarkable prospect of this species for modern medicine. Moreover, the high content of polysaccharides in *Sargassum* cell wall highlights even more the potential application of this species for hydrocolloids and biomaterial industries (Mirshafiey and Rehm, 2009; Rhein-Knudsen *et al.*, 2015). For this reason, *Sargassum* has been considered as sustainable resources with appealing bioactivities related to its primary and secondary metabolites (Liu *et al.*, 2012; Yende *et al.*, 2014). Previous studies have reported that *Sargassum* produces diverse metabolite of structural class such as terpenoids, polysaccharides, polyphenols, sargaquinoic acids, sargachromenol, plastoquinones, steroids, glycerides etc. (Yende *et al.*, 2014).

Among the arsenal metabolite compound synthesized by *Sargassum*, polyphenols are one of the most extensively studied as promising bioactive compounds with potential health benefits. Polyphenols in brown algae are mostly phlorotannins, an oligomer of phloroglucinol (1,3,5-trihydroxybenzene). Phlorotannins have been recognized as the most significant group of biologically active substances determining the pharmacological value of algae (Kim and Himaya, 2011). The potential health benefits of phlorotannins comprise antioxidant, antibacterial, anti-HIV, anticancer, anti-arthritis activities and enzyme inhibitory effects (Li *et al.*, 2011).

The successful of phlorotannins extraction is determined by the appropriate selection of extraction method. Rate and yield of extraction are two important factors contributing to the economic feasibility of an extraction process. There are several techniques of extraction available for obtaining polyphenol compounds from brown algae biomass, *i.e.* extraction with water, acid and alkaline hydrolysis, alternative and conventional solvent extraction (CSE) (Michael and Chojnacka, 2014). Up until now, the CSE, consisting of liquid-liquid (LLE), solid-liquid (SLE) and Soxhlet apparatus (Michael and Chojnacka, 2014) extraction, is the most commonly used technique in the extraction of

phenolic compound of algal biomass. Type of solvents, solid/liquid ratio, temperature, time, and agitation are mentioned as crucial factors to increase efficiency of extraction by improving the solubility and mass transfer (Puri et al., 2012; Li et al., 2017). Nevertheless, this extraction technique is often characterized by applications of organic solvents in high volume that are often toxic, long extraction time, low extraction yield of bioactive and present low selectivity (Ibanez et al., 2012). Phlorotannins are easily oxidized due to light and heat-sensitive. As complex polysaccharides linkages constitute the brown algal cell wall in high content, solid-liquid extraction can be a long and exhausting process. It could lead to the degradation of phlorotannins during the process resulting low yield of target compound (Franco et al., 2008; Wijesinghe and Jeon, 2012). Considering that phlorotannins are cell-wall-bound compounds, they cannot be easily extracted using typical extraction methods with aqueous solvents. Thus, it limits the accessibility of solvents to target compound. In other word, the complexity of polysaccharides in algal cell becomes the main physical barriers causing the decrease of extraction efficiency. Therefore, degradation of algal cell wall has been considered as important step so that the active compounds become more accessible (Wijesinghe and Jeon, 2012). To disrupt the integrity of algal cell wall, enzymes are required as they can hydrolyze the components of cell wall thereby increasing the permeability of cell wall that can result higher extraction yields (Puri et al., 2012). A successful use of this technique to enhance the recovery of phenolics compounds from several terrestrial plants has been reported. This enzyme-assistant extraction method on macroalgae has also been reported in several studies (Wijesinghe and Jeon, 2012).

Sargassum might have promising potential as sustainable resources for diverse applications. In order to announce algae for human consumption either directly as vegetables or integrated in foods, the details of their nutritional compositions as well as variations among species are required to acknowledge (Gomez-Ordonez et al., 2011). However, not all algae have been thoroughly studied related to their nutritional values and one of them is Sargassum. In

addition, to our knowledge, there have not been any studies that compare between solid-liquid and enzyme-assisted extraction of *Sargassum* polyphenols, in this case phlorotannins. Moreover, characterization of *Sargassum* phlorotannins extracted with enzymes is still scarce.

#### **Objectives**

This work is part of a search for active molecules within species of Indonesian and French Sargassaceae (Phaeophyceae, Fucales) especially through phlorotannins. Furthermore, this work is conducted within joint supervisor framework between two universities, Université Bretagne-Sud (UBS) in France and Diponegoro University (UNDIP) in Indonesia.

In summary, the objectives of this study will be:

- 1. To evaluate the biochemical composition of different species of Sargassum and its seasonal variation represented by French S. muticum,
- 2. To determine the phenolic compound of different species of *Sargassum* extracted with different type of solvents in the SLE technique followed by its seasonal profile in the French *S. muticum*,
- 3. To evaluate and to compare the efficiency and the biological activity of *Sargassum* extracted with two different techniques, i.e. the SLE and enzyme-assisted extraction (EAE),
- 4. To evaluate the biological activities of *Sargassum* enzyme-assisted extracts.

#### Phase I

In this phase, we studied both the biochemical composition of different species of *Sargassum* and its seasonal variation represented by one species of *Sargassum*, i.e. the French *S. muticum*. The biochemical compositions that will be evaluated are protein, neutral sugar, sulfates groups, polyphenol, uronic acid, nitrogen, and ash.

Later, the seasonal phenolic content of *S. muticum* S/L extract will also be analyzed as well as its seasonal profile. For the S/L extraction, four different solvents (water, methanol, acetone and methanol 50%) were applied during the extraction of phenolic extraction.

#### Phase II

In this phase, *Sargassum* polyphenol was extracted using solid-liquid extraction in four different solvents, namely water, methanol, methanol 50%, and ethanol 75%. Later, the antibacterial and antioxidant activity represented by DPPH radical scavenging activity and reducing power capacity of *Sargassum* polyphenol were evaluated.

#### Phase III

In this phase, focus of our work will be more about the biological activity of *Sargassum* enzyme-assisted extracts both the cure and fractionated samples. Therefore, in this part, the biological activities analysis were extended to reveal the possibility potential in nutraceutical, pharmaceutical and cosmeceutical industry.

The biological activities evaluated were the antibacterial, antioxidant represented by DPPH (1,1-diphenyl-2-picrylhydrazyl) free radical scavenging activity and FRAP (Ferric Reducing Ability of Plasma), inhibition of tyrosinase and elastase as well as antiviral activity and cytotoxicity based upon cell viability of Vero Cell/HSV-1 and fibroblast model.

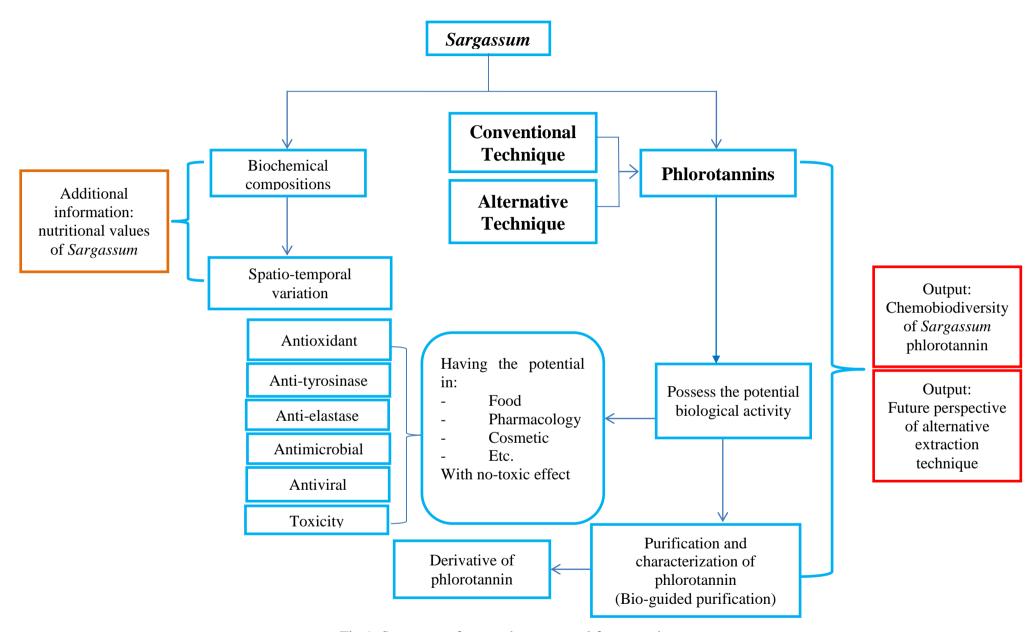


Fig 1. Summary of research conceptual framework

### **Chapitre 1 Literature Review**

#### 1.1 General introduction

Definition of algae classes is very complex and many other criteria are involved, such as the nature and locations of the reserve glucanes, the nature of their cell walls, the number of periplastidial membranes, the shape of mitochondrial crystal, the presence and ultra structure of flagella, the type of mitosis, the mode of nutrition, reproduction and habitat (de Reviers, 2002). In the biological sense of the word, the word algae (algae) is used to designate a large, varied, and heterogeneous group of organisms that, at present, do not have a clear cut, formal taxonomic status since it keep on up-dating and is under constant and rapid revision at all levels (Barsanti and Gualtieri, 2006; Mouritsen *et al.*, 2013).

Algae might resemble to plant yet they are more likely to be saltwater dwelling, simple, multicellular, macro thallic organisms that fall into the rather outdated general category of 'plants'. They are the oldest members of the plant kingdom, extending back to many hundreds of millions years. They are typically plant without leaves and stems as well as root system to take in water or nutrients (like the terrestrial plant) as their cells are in direct contact with the surrounding water from which they derive their nourishment. They do not form embryos as their reproductive structures consist of cells that are potentially fertile and lack of sterile cells covering or protecting them. For the multicellular algae, they are generally attached with so-called *holdfast* serving as anchorage functions (Mouritsen *et al.*, 2013).

Most macroalgae come from either brown algae (phylum Ocrophyta, class Phaephyceae) or red algae (phylum Rhodophyta). A few marine macroalgae occur in the phylum Charophyta since about 600 large green algal taxa are in the phylum Chlorophyta (Dawes, 2016). Macroalgae in these four groups have more fundamental differences regarding cytological, chemical and life histories, between one another than with the vascular plants (Dawes, 1998). Dring (1982)

listed 900 species of green algae, 997 brown algae, and 2540 red algae worldwide. Since then, field, culture, and molecular studies have identified many new and cryptic species and taxonomic relationships. These advancements in phycology have resulted in major changes in the classification of algae (Schneider and Wynne, 2007, 2013; Wynne and Schneider, 2010). Thus the number of seaweed species has increased. Guiry and Guiry (2015) listed 4422 species in the Charophyta and 6055 in the Chlorophyta, 2041 species in the class Phaeophyceae, and 7107 species in the Rhodophyta. In W.R. Taylor's 1957 flora *Seaweeds of the Northeastern Coast of North America*, he listed 170 genera (37 green, 57 brown, 76 red algae) and 401 species (95 green, 142 brown, and 164 red algae), while Mathieson and Dawes (2016) have described 248 genera (57 green, 89 brown, 102 red) and 535 species (146 green, 180 brown, and 209 red taxa) for the same geography (Dawes, 2016).

For centuries, coastal communities have harvested a wide variety of marine macroalgae within the entire algal group, *i.e.* brown, green and red algae. At first, they were frequently used for domestic human consumption; later, industrial uses were discovered. Macroalgae is a lucrative business, driven by economic rather than environmental conditions (Delaney *et al.*, 2016). Along with the growth of human population, the need to look for alternatives natural resources raises. Moreover, the scarcity of land for cultivation, infertility of soil, limitation of various crops, requirement for more fodder, requisition of specific compounds for life-saving medicines, and other specialized requirements have compelled mankind to search more into the depth of the ocean (Murty and Banerjee, 2012). Aside from consumptive forms, macroalgae also include in the non-consumptive forms as medicine, fertilizer and animal feed, inputs into industrial process and for other purposes such as building materials (Delaney *et al.*, 2016).

#### 1.2 Brown algae (Phaeophyceae)

The kingdom Chromista has over 21,500 taxa and includes the phyla Dinophyta (Dinoflagellates), and the Ochrophyta. The latter phylum contains 16 classes and

over 3700 species including the brown algae in the class Phaeophyceae (Guiry and Guiry, 2015). Based on nucleotide sequences, it appears that the Phaeophyceae diverged from the diatom phylum Bacillariophyta that is also in the kingdom Chromista around 200 million years ago (Hori and Osawa, 1987). The majority of species in the Ochrophyta occur in the classes Chrysophyceae (613 species that range from unicellular to filamentous freshwater taxa), Xanthophyceae (559 species including the coenocytic marine genus *Vaucheria*), and the Phaeophyceae (with about 265 genera and 2040 species; (Guiry and Guiry, 2015)). Over 95% of the brown algae are marine (Wang *et al.*, 2013).

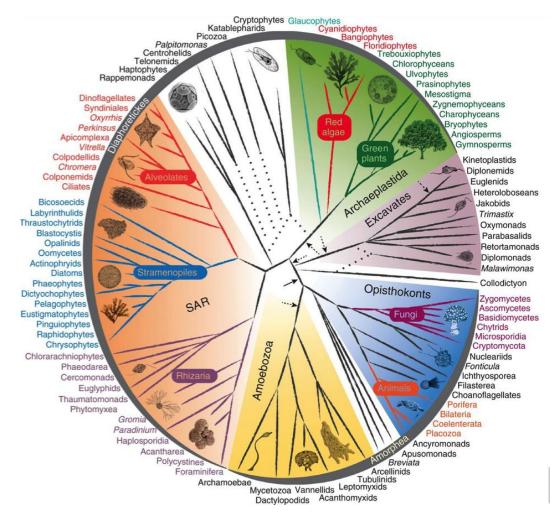


Fig 1-1 Phylogenetic tree showing the position of Phaeophyceae within the Eucaryotic group (Burki, 2014).

Fig 1-1 illustrates the phylogenetic tree showing the position of Phaeophyceae within the eukaryotic group based on Burki (2005). As presented in this figure, brown algae are included in the SAR group – an acronym of its constituents, *i.e.* stramenopiles, alveolates, and rhizaria, particulary in stramenopiles (also known as heterokonts). This super-group, most recently recognized, embraces a very large diversity of protists, including diatoms, multicellular algae and heterotrophic, often parasitic species such as oomycetes and *Blastocystis* (Burki, 2014).

In addition to Fig 1-1, Fig 1-2 illustrates the phylogenetic tree of eukaryotic based on the major component of cell wall. CesA\* represents members of the cellulose synthase family whose proteins assemble into rosette terminal complexes, CesA is the ancestral form of cellulose synthases, and CslA/C is a single gene that is most similar to the land plant CslA and CslC gene families. The arrows indicate the origin of the plastids: PE (solid arrow), primary endosymbiosis; SE1 (dotted arrow), secondary endosym- biosis scenario 1 in which the Alveolata, Cryptophyceae, Haptophyta, and Stramenopiles originate from a common ancestor, which acquired its plastids by a secondary endosymbiosis event with a red alga (Cavalier-Smith, 1999), which Rhizaria subsequently lost and the Chlorarchniophyta regained (Keeling, 2010); SE2 (dashed arrows), secondary endosymbiosis scenario 2, separate acquisitions of rhodobionts (Baurain *et al.*, 2010).

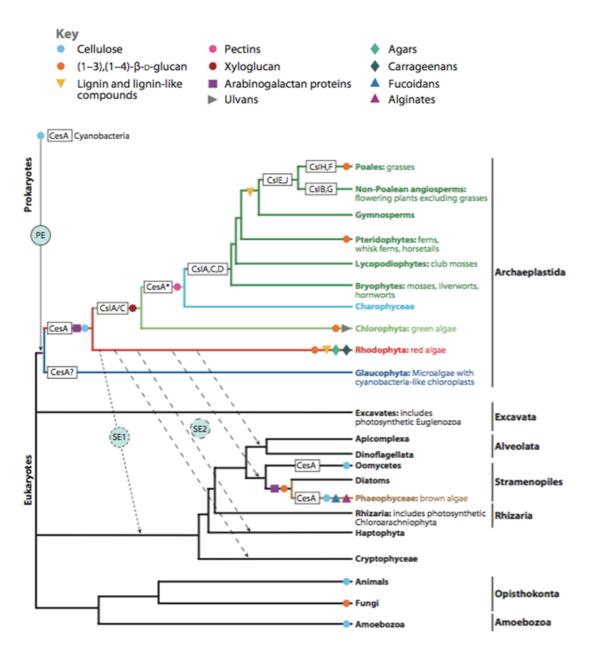


Fig 1-2Simplified phylogenetic tree representing the major cell wall component (Popper *et al.*, 2011).

Members of brown algae are multicellular, and their primary photosynthetic pigments are chlorophylls 'a' and 'c' but derive their brownish colour characteristic from a mixture of xanthophyll pigments ( $\beta$ -carotene, neofucoxanthin, and other carotenoids), of which fucoxanthin is the most important (Dring, 1982; Dawes, 2016). Primary reserve of brown algae is chysolaminaran, a  $\beta$ -1,3-linked glucan occurring in solution in special vacuoles as well as mannitol and lipid. Cell wall of brown algae have a felt-like weft of

cellulose microfibrils stiffened by calcium alginate, which is a commercially important phycocolloid (Dawes, 1998). In general, brown algal life histories are either haplodiplontic or diplontic with morphological appearance ranging from filamentous thalli (Ectopearpus fasciculatus Harvey) to large complex with blades, stipes and floats as with Sargassum filipendula C. Agardh. Most of larger algae, belonging to the 17 orders in the class Phaeophyceae, occur in the orders Desmarestiales, Dictyotales, Fucales, and Laminariales. Brown algae are most inhabited the cold (arctic to boreal) to temperate waters, while most species of Dictyoles and Sporochnales are mainly found in subtropical or tropical waters. Many brown algae are lithophytes and grow attached to hard substrate (Dawes, 2016), and some of them, such as members of the order Fucales (the rockweeds), are commonly found along rocky seashores (Venugopal, 2011). The major brown algae, which include such genera as Macrocystis, Laminaria, Sargassum, Pterygophora, and Nereocystis grow upward like trees and spread their blades at the surface of the water, where they obtain the maximum amount of light. The Pacific coast of both North and South America is dominated by Macrocystis, whereas Laminaria is dominant in Atlantic waters and in Japan (Venugopal, 2011).

Several methods of reproduction occur in algal reproduction, they are vegetative by the division of a single cell of fragmentation of a colony, asexual by the production of motile spore, or sexual by the union of gametes. Vegetative and asexual modes allow stability of an adapted genotype within a species from a generation to the next. Sexual mode involves plasmogamy (union of cells), karyogamy (union of nuclei), chromosome/gene association, and meiosis, resulting in genetic recombination. In this mode, variation is less yet possible as it is mostly because of the waste of gametes that fails to mate (Barsanti and Gualtieri, 2006).

In Fucales, the reproduction occurs by releasing the haploid gametes and the biological life cycle is monogenetic diplophase. Adult individuals are monoic or dioecious; they are meiosporpohytes (2n chromosomes), producing gametes by

chromatic reduction. Gametes are produced in conceptacles. The conceptacles are grouped into receptacles at apexes of fertile lateral in the Genera of *Cystoseira* and *Bifurcaria*, at the base of pleuridia along the axis in the Genera *Turbinaria* and along the fertile laterals in the Genera of *Sargassum* (Le Lann, 2009).

The plant body of Sargassum is a diploid sporophyte, reproducing both by vegetative and sexual methods. It does not multiply asexually by means of spores. Instead it reproduces by vegetative method and the formation of sex organs. Fragmentation is the only known method of vegetative reproduction in this alga. Their vegetative reproduction involves fragmentation. Each fragment grows into new plant. The separation commonly occurs accidentally or by death and decay of older parts separating the younger parts of the branches. In the Sargasso Sea, huge masses of floating Sargassum are formed by continuous vegetative reproduction. Their conceptacles can be seen occasionally, however they are functionless since the maintenance of the population is entirely by vegetative means (Graham and Wilcox, 2000). The sexual reproduction is oogamous and involves the fusion of motile sperm or male gamete with a non-motile ovum or female gamete. They are respectively borne inside antheridia and oogonia. They are also termed as microsprangia and megasporangia. The spores germinate inside the sporangia to produce gametes. The sex organs are produced inside special cavity called conceptacles. Conceptacles are found inside small finger-like branchlets called receptacles and usually bear only one type of sex organs (Ogawa, 1976; Critchley et al., 1991).

A large number of antheridia develop in each male conceptacle. They are borne on the lower branches of paraphyses. The first antheridium is borne at the tip of a 3-celled filament which arises from a cell of the fertile layer. The basal cell remains at the level of the fertile layer. The middle one is known as stalk cell. The upper or antheridial cell develops into an antheridium. Antheridium is ovoid in outline. It has a thick wall which becomes mucilaginous towards maturity. Internally the antheridium contains initially a single diploid nucleus and a few chromatophores. The diploid nucleus divides meiotically and then mitotically to

form 64 haploid nuclei. The chromatophores also divide. They ultimately get placed near the nuclei in one to one ratio. The cytoplasm cleaves and the contents of the antheridium get divided into 64 haploid protoplasts. Each such protoplast transforms itself into a single sperm or antherozoid. Thus 64 sperm develop inside an antheridium (Ogawa, 1976; Critchley *et al.*, 1991).

The antherozoids are then expelled into the medium while the oosphere, after rupture of the oogone wall, remains on the surface of the design. The antherozoids move towards the oosphere by chemotaxis, attracted by fucoserratene, pheromone excreted by the oospheres (De Reviers 2003a). After gametes melt, zygotes form on the surface of receptacles (Yoshida, 1983). The zygotes germinate to give seedlings bearing 8 to 16 rhizoids, which once released, sediment rapidly and attach to a hard substrate, usually near the parental thallus (Stiger and Payri 1999).

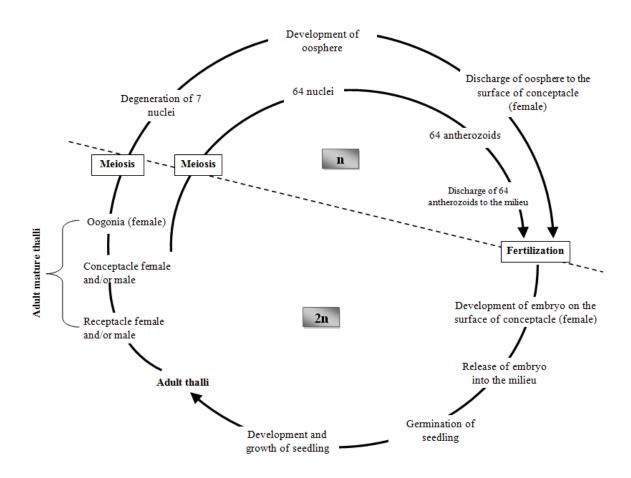


Fig 1-3Life cycle of Sargassaceae according to Stiger (1997).

# 1.2.1 Brown algae Sargassum

Classification of brown algae *Sargassum* according to Lee (2008) and Guiry and Guiry (2015) is:

Phylum: Ocrophyta

Class: Phaeophyceae

Order: Fucales

Family: Sargassaceae

Genus: Sargassum

The thallus of *Sargassum* is differentiated into root-like form called holdfast; their main cylindrical axis into stem-like form called stipe and their strand into leaf-like form called blade (Fig 1-4). In addition, they have air bladder in every corner of their thallus. These bladders are responsible for their floating ability. As a result, they are considered as a disturbing community especially around the harbor and coastal. Furthermore, by having bladders, *Sargassum* is well known by its fast and long-distant distribution. Other factors that influence their distribution are their fast growth rate, first year fertility and monoecious reproductions – eggs and sperms come from the same gametophyte. Main axis of *Sargassum* is perennial and slow growth rate; therefor their main lateral can grow annually (Graham and Wilcox, 2000).

Both the growth and the distribution of *Sargassum* are influenced by their surroundings, which is their environmental. Each environmental factor gives different effect to different species of *Sargassum* especially related to their metabolism. There are several environmental factors considered as the most influenced variables based on previous researches. The most important parameters regulating algal growth are nutrient quantity and quality, light, pH, salinity, temperature, and turbulence. The most optimal parameters as well as the tolerated ranges are species-specific and the various factors may be inter-independent and a parameter optimal for one set condition is not necessarily optimal for another (Barsanti and Gualtieri, 2006).

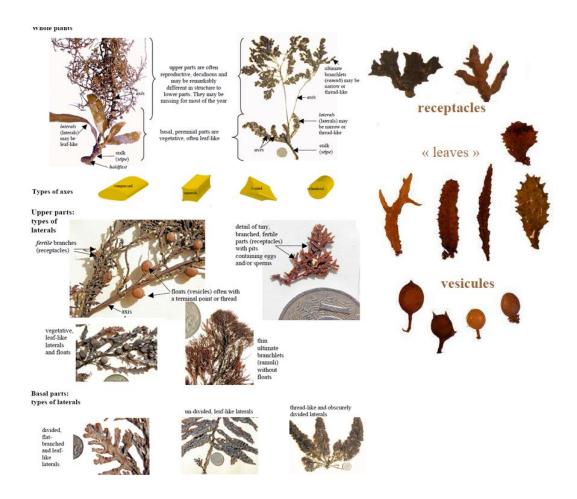


Fig 1-4 General morphology of Sargassum (Baldock, 2009; Mattio et al., 2009).

# 1.2.1.1 S. aquifolium

Taxonomic synonym of *S. aquifolium* is *S. echinocarpum* (Bold and Wynne, 1985). In general, the morphology of this alga is similar with other *Sargassum* species. The difference lays on its flattened branches, with lance-like leaves, vesicles which are mucronate with winged pedicles (Low and Chou, 2013).



Fig 1-5 S. aquifloium (pers. documentation).

## 1.2.1.2 S. ilicifolium

S. ilicifolium is generally brown or dark green having a taxonomic synonym as S. duplicatum. Oogonia and antheridia occur in conceptacles embedded in receptacles on special branches. Their extensions are modified pedicles on which berry like bladders are borne. Linear-lanceolate fronds possess a dentate sinuate margin, distinct midrib and small numerous irregularly placed cryptostomata. The shapes of their vesicles are globular often without prolongation at the top. Then shapes of their receptacles are cylindrical, filiform and irregularly ramified (Sumithra and Arunachalam, 2014).



Fig 1-6 S. ilicifolium (pers. documentation).

### 1.2.1.3 S. muticum

S. muticum brown color varies from dark brown to pale, yellowish brown depending on season and growing conditions. They have regularly alternating lateral shoots or branches on a central perennial stem. They attach to substrate with their disc-shaped holdfast. They have numerous small air bladders of 2-3 mm round lying on the stems. These bladders help them to stand upright in the water or float if there is any part detached from the basal.

S. muticum has a 75-120 cm long frond, but it reaches 1.5-2 m long in Swedish waters, 6-7 m in French waters and up to 8.5 m in Norwegian waters. Their lateral branches detach in summer or autumn, leaving a short perennial basal steam to over winter (Wallentinus, 1999). During the summer, their receptacles develop in the areas where the annual shoot attaches to the stem but may also sit on top of the branch.



Fig 1-7 S. muticum Yendo (Fensholt) (pers. documentation).

Sheltered hard-bottoms are their habitat preferences. However, they may also attach to hard substrate on soft-bottom, such as stone or shells. *S. muticum* tolerates water currents but they are not fond to direct wave exposure. They are often found in shallows waters, in lower intertidal and upper sub tidal zones. *S. muticum* stands for a wide range of abiotic conditions and is thus considered highly invasive (Wallentinus, 1999). Their optimal salinity level for growth is 34 ‰. They need a temperature of more than 8 °C at least four month to reproduce (Steen, 2004).

# 1.2.1.4 S. polycystum

The morphology of *S. polycystum* is not different from other Phaeophyceae. They have a cylindrical small spiny thallus with a disc-shaped holdfast and air bladders in every corner. Typical characteristics of this species are their small, pointed and jagged blades. *S. polycystum* can be found at the tide pool area, the reef flats, the reef slope, the ridge and the lagoon. They grow mostly in the tropical region with temperature ranged from 27.5-29.3 °C and water salinity 32-33.5 ‰. In general, *Sargassum* needs of light intensity is higher compared to the red algae (Kadi, 2005). They need 6500-7000 lux of light intensity (Boney, 1965). Consequently, they inhabit mostly along the coast with 0.5-5 m.



Fig 1-8 S. polycystum (pers. documentation).

### 1.3 Abiotic factors

### 1.3.1 Temperature

The ideal temperature for algae varies depends on where they live. In general, for algae inhabit the polar zone they can tolerate temperature < 10 °C, the temperate zone between 10-25 °C and tropical > 20 °C. Temperate algae can tolerate cold water at least down to -1.5 °C (seawater freezing point) (Lobban and Harrison, 1997).

Temperatures have fundamental effects on the chemical-reaction rates related to the metabolic rates of algae. The effects of a given temperature change will not be same on all metabolic processes, because of differing temperatures sensitivities of enzymes and the influences of other factors, including light, pH, and nutrients (Lobban and Harrison, 1997).

Temperatures optima vary among species and among strains as well as between heteromorphic life-history stages. Changes with the age of the thallus have also been noted. Norton (1977) has suggested that *S. muticum* shows a high growth rate at high temperatures and it is what makes them so invasive in relatively warm waters.

S. muticum optima temperature for maximum growth is 25 °C. He also reported that the temperature range of S. muticum germlings is able to tolerate a temperature between 10-30 °C. This species is able to tolerate an annual temperature range spanning about 10 °C to 15 °C (Hales and Fletcher, 1989). S. polycystum is intolerant to seawater temperature > 30 °C, thus they have a lower optimum growth temperature between 20-25 °C and tolerance limit between 25-30 °C (Hwang et al., 2004). Rao et al. (2002) reported that their rapid increase in growth and maximum development were shown during the low period of low temperature (22 to 28 °C). For S. ilicifolium, its highest growth rate was recorded during the cold season at mean temperature 28 to 30 °C. In regions where maximum temperatures are lower, S. ilicifolium maintains relatively high biomass all through biomass through the year (Ateweberhan et al., 2005), for example at the Philippines, with the mean annual temperature 29.1 to 32.7 °C (Hurtado and Ragaza, 1999). S. aquifolium has an optimum temperature for its growth at 23 – 30 °C with low tolerant limit between 19-23 °C. Its germlings attained maximum growth at 35 °C and its maximum growth at 35 °C (Hamza et al., 2015).

In Indonesia, as tropical characterized with its two season, i.e dry and rainy season, there are not much researches related specifically to the seasonal variation or effects of temperature towards the growth rate of *Sargassum*. Apparently, the optimum temperature ranges from 27.25 to 29.30 °C (Kadi, 2005).

### 1.3.2 Salinity

Algae found mostly in the spray zone (supralittoral zone) are subjected to the open air for certain period of time frequently experience salinity fluctuation. Evaporation causes an increase in the salinity. The most important effects of salinity are the osmotic consequences of the movements of water molecules along water-potential gradients and the flow of ions along electrochemical gradients. These processes take place simultaneously and both are regulated in part by the semipermeable membranes that surround cells, chloroplasts, mitochondria and vacuoles.

In the brown algae, changes of salinity as well as temperature and irradiance might induce the polyphenol production (Ragan and Glombitza, 1986). Connan and Stengel (Connan and Stengel, 2011) have demonstrated that the reduction of salinity decreases the polyphenol content and increase in polyphenol exudation in the *Fucus vesiculosus* and *Ascophyllum nodosum*. The increase of polyphenol exudation was assumed to be a mechanism that prevents the copper from entering and damaging the photosynthesis apparatus of algae. Thus, this mechanism gives the ability of algae to tolerate high copper contamination in the surrounding – also known as a detoxification mechanism (Connan and Stengel, 2011).

Algae are extremely tolerant to changes in salinity. One of them is *S. muticum* who has been well known by their tolerance towards salinity ranges, *i.e.* between 25 to 34 ‰. Pacific Northwestern *S. muticum* can tolerate different range of salinities, which is 24 to 25.5 ‰, 27 to 35 ‰ and 21 to 30 ‰ (Josefsson and Jansson, 2011). In Indo-Pacific Ocean, *Sargassum* tolerates the change of salinity range between 32–33.5 ‰ (Kadi, 2005).

### 1.3.3 Nutrients

The nutrient availability will affect algae's growth rate. C, H, O, P, Mg, Cu, Mn, Zn and Mo are considered to be required by all algae (O'Kelley, 1974; DeBoer and Ryther, 1977). Among these possible limiting elements, nitrogen appears to be the most frequently limit the growth of algae (Topinka and Robbins, 1976), followed by phosphorus as the second most limiting nutrient (Manley and North, 1984). Nitrate, nitrite, ammonium and phosphate concentrations vary from 0 to 30, 1, 3, and 2  $\mu$ M, respectively, for most pristine temperature areas (Lobban and Harrison, 1997). The concentrations of these elements vary depends on the physical and biological parameter of the surrounding environment where the algae live.

### **1.3.4 Light**

Light intensity plays important role since it is the source of energy during photosynthetic reactions in algae. However, the requirements greatly vary depends on the depth and the algae density. If the light is too high, it might lead to photo-inhibition.

Light intensity also affects the production of polyphenol contents found in the brown algae. *Fucus spiralis*, *F. vesiculosus* and *A. nodosum* that grow under high daily irradiation inhabited the intertidal and eulittoral zone had a higher polyphenol content than algae grow in the low intertidal or sublittoral zone (Abdala-Diaz *et al.*, 2006).

Most often employed light intensities range between 100 and 200 μE sec<sup>-1</sup>, which corresponds to about 5 to 10 % of full daylight (2000 μE m<sup>-2</sup> sec<sup>-1</sup>) (Barsanti and Gualtieri, 2006). Kadi (2005) also reported that *Sargassum* needed light intensity ranged between 6000–7500 lux equivalents to 100-200 μE m<sup>-2</sup> sec<sup>-1</sup>. Nevertheless, the optimum light intensity for maximum growth required by marine algae is species-specific. For example, *S. muticum*, their maximum germling growth was recorded at high light intensity (200 μE m<sup>-2</sup> sec<sup>-1</sup>).

### 1.3.5 Wave and currents

These two physical factors are an incredible source of energy to the coastal environment. It can contribute to ten times more energy than does the solar radiation. This tremendous energy input explains the high productivity of coastal macroalgae communities. In other words, this energy gives a positive effect on algae. By constantly moving the frond of algae, waves and currents reduce the shading and bring a constant resupply of inorganic nutrients to the algae. These factors also reduce the nutrient depletion by reducing the thickness of the boundary layer of unmoving water associated with algae surfaces. In addition,

waves and current can remove sessile organism that compete with algae for attachment in the intertidal zone (Lobban and Harrison, 1997).

Along with wave and currents, tidal pattern also affects the algae especially to the structural communities. However, tidal patterns based on the observation by Pavia and Brock (Pavia and Brock, 2000) seems to regulate the polyphenol content in the tissue of *A. nodosum*. Greater polyphenol concentration was found during the low tides than during immersion. In addition, this variation also correlated to the air temperature (case for *P. canaliculata*) and to the tidal regime as in the case of *A. nodosum* (Connan, 2004).

#### 1.3 Biotic factor

Impact of herbivores on macroalgae have been extensively studied and considered as one of crucial factor affecting the metabolic system of algae. It is believed that herbivores of macroalgae have a number of counter adaptations, including feeding specialization and sequestration of defenses. Such adaptation represents potential steps toward a top-down control of macroalgal communities (Pereira and da Gama, 2008).

To counter attack the herbivores, algae employ a number of chemical, morphological, and structural defense mechanisms. These defense mechanisms vary according to spatial and temporal patterns of both algae and their associated herbivores. Moreover, such mechanisms are metabolically costly for algae. Previous study on the grazing of *F. vesiculosus* by isopod showed a temporally variable defense pattern that improve the algae's costs/benefit ratio since the deterrence of herbivore involves redistribution of metabolism from growth and primary towards defense (Flöthe *et al.*, 2014). Grazing activity caused by fish, crabs, sea urchins, gastropods, ampiphods, and isopods on algae as their primary nutrient source, might result in the alteration of algal communities. If the grazing activity is so intense, it can destroy a kelp forest occupying several square kilometers. Biomass, coverage, and composition of *Sargassum* sp in central Japan

was altered by the presence of sea urchin *Diadema setosum* (Ishikawa and Kurashima, 2016).

Other than grazing, microbial colonization and epibiosis on algae also contribute to algal community (Pereira and da Gama, 2008). As marine macroalgae are continuously immersed in seawater they are regularly colonized by many microbial populations forming surface biofilm which exert significant physiological health pressures by changing the hydrodynamics, increasing resistance to heat transfer, and enhancing subsequent colonization by other fouling organisms and corrosion and destruction of underlying surfaces leading to death of the host (Stern *et al.*, 1996). The encrusting bryozoan *M. membranacea* is an invasive species along the Atlantic coast of Nova Scotia, to which a disruption in sea urchin-kelp dynamics has been attributed (Scheibling *et al.*, 1999). By overgrowing kelp fronds, it causes thallus fragmentation and loss, decimating kelp beds.

Algal defense mechanisms have evolved a variety of strategies including: (1) temporal and special escape, (2) chemical defense, (3) morphological or structural defense, and (4) tolerance (Pereira and da Gama, 2008). Herbivore pressures fluctuate as a function of available habitats and seasonal patterns. Abundant populations of unpalatable algae coexist with attractive, palatable algae; fish (as example) rapidly consume the slow growing *Hypnea musciformis* when cultured in isolation, but when the unpalatable brown algae *Sargassum* grows adjacent to *H. musciformis* it grows well (Hay, 1986). Another interesting defense performed by macroalgae is the production of secondary metabolites such as aromatic compounds, terpenes, and many more, as a mechanism to deter epibions and herbivores (Kubanek *et al.*, 2004; Wiencke *et al.*, 2007; Blunt *et al.*, 2012). In the case of brown algae, they are well known for its polyphenols (phlorotannins) productions (Jormalainen *et al.*, 2003).

### 1.4 Biochemical composition of Sargassum

Algae have been subjected to many researches these past years and some of them have been introduced and used in food, cosmeceutical and pharmaceutical industries. Biochemical constitutions of algae have been widely conducted worldwide. Polysaccharides, proteins, minerals, vitamins, dietary fiber, and lipids from macroalgae have been reported frequently (Peng *et al.*, 2013). These constituents of algae, including major nutrients, like proteins and minerals, and micronutrients, like lipids and witamins, provide high nutritional values that are beneficial for human (Marinho-Soriano *et al.*, 2006; Patarra *et al.*, 2011). Because of their high nutritional value, algae are traditionally consumed as food or as herbal medicine. Furthermore, they are also used as fodder, fertilizer, fungicides, herbicides, dietary supplements, and as a resource of phycocolloides such as agar, alginate, and carrageenan for various industrial applications (Peña-Rodríguez *et al.*, 2011; Kilinç *et al.*, 2013).

Biochemical composition of algae varies, depending on species, time of collection, geographic habitat and on many external conditions, like water temperature, light intensity, and nutrient concentration in water (Marinho-Soriano *et al.*, 2006; Sudatti *et al.*, 2011). Additionally, chemical composition of algae demonstrates great differences of concentration of nutritional factors such as proteins, lipids, minerals, or dietary fiber within diverse genera of algae (Marinho-Soriano *et al.*, 2006; Leal *et al.*, 2008; Stengel *et al.*, 2011).

### 1.5.1 Cell wall and sugars

Algae are widely known with their large amount of polysaccharides as constituting their cell wall and reserved as their storage. In general, algal cell wall is distinguished as skeletal and amorphous matrix. The skeletal phase is mainly composed of cellulose in the form of crystalline microfibrils. As for the amorphous matrix phase, separates the cell wall from the cells and serves as cement, mostly contains a network of anionic polysaccharides called phycocolloides (Jouanneau, 2015).

Brown algal cell walls (Fig 1-9) are generally composed of two layers with cellulose making up the main structural skeleton (Kloareg and Quatrano, 1988) and its content ranges from 5.7 to 14 % (Kloareg and Quatrano, 1988; Park *et al.*, 2000). Amorphous component of brown algal cell wall is made up of alginic acid and fucoidan (Lee, 2008), whereas their mucilage and cuticle polysaccharides are mainly composed of fucoidans, alginate and laminarin with total ratio 3:1:1 (Balboa *et al.*, 2013). Alginate generally makes up to 10-30% of the algae dry weight (Park *et al.*, 2000) as for fucoidan, their content range from 1-25% (Obluchinsjaya *et al.*, 2002; Kusaykin *et al.*, 2008). Beside structural polysaccharides, algae also contain storage polysaccharides, notably laminaran and mannitol (Stiger-Pouvreau *et al.*, 2016).

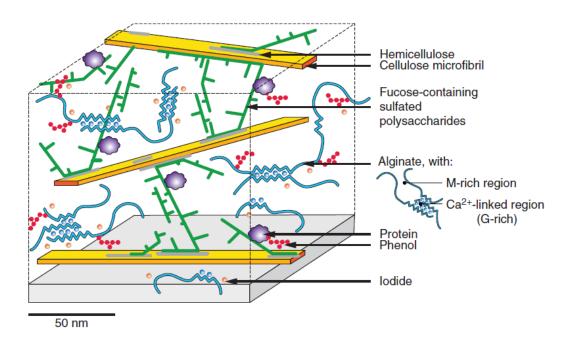


Fig 1-9 Cell wall model of brown algae from the order Fucales (Deniaud-Bouet et al., 2014).

Table 1-1 Content of total polysaccharides in some species of Algae (% of dry weight) (<sup>a</sup>Ahmad *et al.*, (2012); <sup>b</sup>Gomez-Ordonez *et al.*, (2010); <sup>c</sup>Haque *et al.*, (Haque *et al.*, 2009); <sup>d</sup>Marinho-Soriano *et al.* (2006); <sup>e</sup>Patarra *et al.*, (2011); <sup>f</sup>Burtin, (2003a).

Genera	Total polysaccharides (% of dry weight)		
Fucus	63.88 <sup>e</sup>		
Padina	85 <sup>a</sup>		
Sargassum	83°; 47.43°; 68 <sup>d</sup>		
Turbinaria	83 <sup>a</sup>		
Laminaria	30.23°; 37.3 <sup>f</sup>		

Figure 1-9 presents the cell wall of brown algae proposed by Deniaud-Bouet (2014). Cellulose microfibrils are sparse having a ribbon-shaped like form. Alginates and fucose-containing sulfated polysaccharides form the greater part of cell wall polymers, the latter acting as cross-linkers between cellulose microfibrils. Phenols are likely associated with alginates and proteins. Large amounts of iodide are also present in the wall (Deniaud-Bouet *et al.*, 2014).

### 1.5.1.1 Crystalline microfibrils form

The skeletal phase is mainly composed of cellulose in the form of crystalline microfibrills, which can be replaced by xylan in the green and red algae, in addition to mannan in the green algae. Brown algal embedding matrix is predominantly alginic acid or alginate (salt of alginic acid) with smaller amounts of sulfated polysaccharide (fucoidan). In red algae, this matrix contains sulfated galactans, *i.e.* carrageenan, agar, poryphyran (Stiger-Pouvreau *et al.*, 2016).

Algal cellulose is rarely a pure  $\beta$ -1,4 glucan; more frequently it contains sugar than glucose, commonly xylose (Mackie and Preston, 1974). Algal cellulose is homopolysaccharides; whereas their hemicellulose is heteropolysaccharides whose constituents are hexoses, pentoses, and uronic acids. Cellulose belongs to polymers that are very difficult to dissolve and this is determined by compact packing – mainly crystalline, with single armophous region – of cellulose chains in microfibrils. Adjacent chains are linked by mutiple hydrogens bonds instead of covalents. This linkage is the reason why cellulose is insoluble in water, dilute

acid and alkalies, and even in organic solvents (Tarchevsky and Marchenko, 1991).

Unlike cellulose, hemicelluloses are more easily soluble to water due to the looser packing of component molecules; hence they are more accessible to solvents. Hydrolysis of hemicelluloses into sugars components is usually carried out with 2 M hydrochloric acid at 100 °C. Hemicelluloses can be subdivised into several groups on the basis of monosaccharide: 1) arabinans; 2) xylans (homoand heteropolysaccharides including arabinoxylans, glucuronoxylans, arabinoglucuronoxylans, galactose, arabinose, rhamnose, glucuronic and galacturonic acids, etc.); 3) glucans (homopolysaccharides including β-1,3-Dglucan callose, and heteropolysaccharides such as arabinoglucans); 4) galactans (arabinogalactans); 5) mannans (glucomannans, galactomannans, glucogalactomannans; 6) fructosans and glucofructosans; 7) galacturonans, and 8) mannuronans (Tarchevsky and Marchenko, 1991).

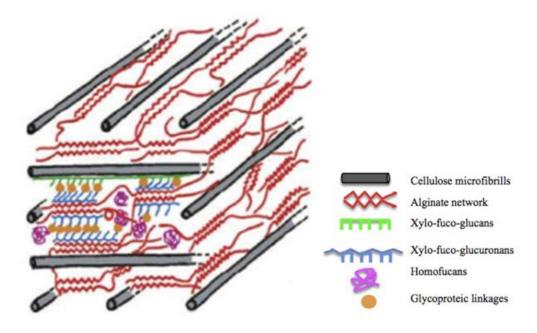


Fig 1-10 Detail of brown algal cell wall composition (Michel et al., 2010).

# 1.5.2.1 Charged matricial form

As for the amorphous matrix phase, separating the cell wall from the cells and serving as cement, mostly contains a network of anionic polysaccharides called phycocolloides (Jouanneau, 2015).

Alginic acid consists of (1-4) linked β-D-mannuronic acid (M) and α-L-guluronic acid (G) residues (Figure 15). In the mixed salts form – known as alginates – they contain several cations such as Na<sup>+</sup>, K<sup>+</sup>, Mg<sup>2+</sup> and Ca<sup>2+</sup> (Percival and McDowell, 1967). Brown macroalgae contain alginates ranging up to 40% of the dry weight (Stiger-Pouvreau *et al.*, 2016).

To isolate alginic acids, the algal biomass is pretreated with dilute acid to remove polyvalent cations, and the polymer is extracted in the form of readily soluble sodium salt by treatment with sodium carbonate solution at elevated temperature. Alginic acids are precipitated after acidification of the extract, whereas sodium alginate may be precipitated with ethanol; alternatively, insoluble calcium alginate may be obtained by addition of calcium chloride ((Haug, 1962; Hernandez-Carmona et al., 1999a, 1999b). Alginic acids isolated from various sources may differ in the M/G ratio depends on the species, the part of the thallus, their surrounding conditions and the harvest period (Haug et al., 1974; Apoya et al., 2002; Chandia et al., 2004; Skriptsova et al., 2004). Moreover, the molecules may differ in block composition, since uronic acid residues are usually arranged in two homopolymeric blocks, poly-M and poly-G, divided by approximately alternating sequences, poly-MG. Such a blockwise distribution of monomeric units is the result of specific biosynthetic pathway, where the formation of linear polymannuronan precursors is followed by epimerization of some  $\beta$  -Dmannuronic acid residues at C5 under the action of mannuronan-C(5)-epimerase, giving rise to the appearance of  $\alpha$  -L-guluronic acid residues, single or contiguous, within the polymeric molecule (Skjak-Braek, 1992; Ertesvag et al., 1996).

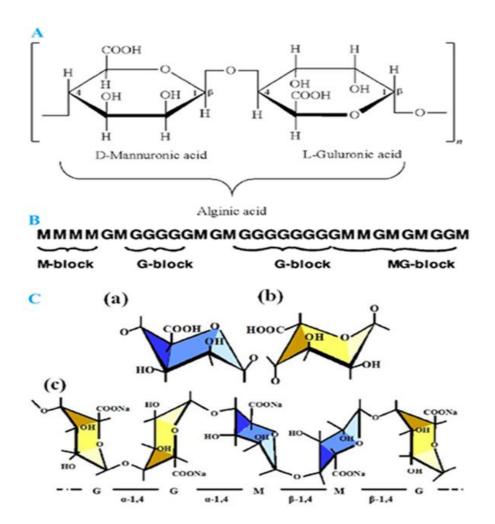


Fig 1-11 Chemical structure of alginates from brown algae; (A) Alginic acid; (B) Distributions of blocks; (C) Ring conformation (Draget *et al.*, 2005; Kashima and Imai, 2012).

Physical properties of alginate including solubility depend on the M/G ratio and the molecular mass ranging from 150 to 1700 kDa depending on the source. Mass molecular of *Padina gymnospora* alginate was about 5000 kDa (Salgado *et al.*, 2007). Molecular weight of alginate extracted from brown algae *Costaria costata* ranged from 394 to 714 kDa (Imbs *et al.*, 2009). Molecular weight distribution of alginate can have implications for the uses of alginate itself. Low molecular weight containing only short G-blocks may not take part in gelnetwork formation; as a consequent, it do not contribute to the gel strength (Draget *et al.*, 2005). Therefore, to have great gel strength, an alga should have

more G block units than M blocks. Alginates provide flexibility and strength for algal thallus (Milani and Maleki, 2012);

Various industries related to food processing, pharmaceuticals, feed and cosmetics use alginic acid extracted from *Laminaria hyperborean*, *L. digitata*, *L. japonica*, *Ascophyllum nodosum*, *Macrocystis pyrifera*, *Sargassum*, *Saccharina* and *Undaria* (Jayasankar, 1993; Sartori, 1997; Lee and Mooney, 2012). It has been reported that alginic acid leads to a decrease in the concentration of cholesterol, exerts an anti-hypertension effect, can prevent absorption of toxic chemical substances, and plays a major role as dietary fiber for the maintenance of animal and human health (Murata and Nakazoe, 2001; Nishide and Uchida, 2003; Kim and Lee, 2008). These dietary polysaccharides are not found in any land plants (Balboa *et al.*, 2013). Alginates have been considered only as thickening, gel forming, and stabilizing agents, however, recent pharmacological research discovered that alginates isolated from *Sargassum* demonstrated anticancer (*in vivo*), antiviral (*in vitro*), and hypolipidemic (*in vivo*) activity (Mao *et al.*, 2004; Sousa *et al.*, 2007; Sinha *et al.*, 2010).

Fucoidans or fucans or fucanoids are complex sulfated polysaccharides primarily composed of sulphated L-fucose (Holdt and Kraan, 2011; Balboa *et al.*, 2013) with small proportions of galactose, mannose, xylose, glucose, rhamnose and uronic acid (Pomin and Mourau, 2008; Usov and Bilan, 2009). Figure 1-12 presents the general structure of fucoidan and proposed fucoidan structure isolated from one of *Sargassum* species, *S. myriocystum*.

Molecular weight of fucoidan varies according to species, geographic habitat, season, and even extraction method. Therefore, the exact molecular weight of fucoidan is under debates (Kim, 2012). *Sargassum* sp. extracted by water had molecular weight of fucoidan at 57.7 and 2.7 kDa, while HCl extract of this species had molecular weight at 46.6 and 3.38 kDa (Junaidi, 2013). Regarding the extraction method, fucoidans are easily extracted from the algal cell wall with hot water or acid solution. Mannitol, alginic acid, and laminaran are often

occurred in fucoidans extraction, thus, ethanol precipitation is employed to isolate the fucoidans from brown algae. In addition, elevated temperature of more than 70°C is frequently selected as the extraction condition. Even though, the ambient or slightly elevated temperature is still served as a preferred step for isolating fucoidans prior to any fractionation and purification process (Ale and Meyer, 2013).

Fucoidans account for 10-20% of dry weight and are not found in other divisions of algae or in terrestrial plants. They are very soluble and do not develop highly viscous solutions (Balboa *et al.*, 2013). The chemical composition of fucoidan depends on the algal species, part of the thallus and the harvesting period (Skriptsova *et al.*, 2010). Table 1-2 describes the chemical composition of fucoidan in several species of brown algae.

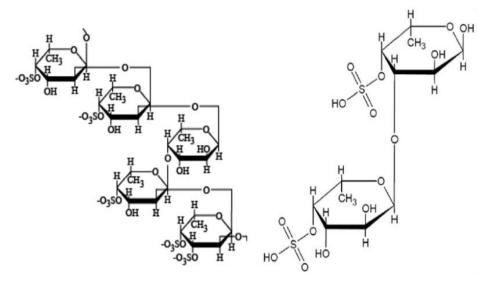


Fig 1-12 Fucoidan structure – a branched sulfated polymer with L-fucose building blocks (Left) Probable structure of fucoidan from *S. myriocystum* as proposed by Badrinathan *et al.* (2012).

Many previous studies have revealed the wide range potential showed by fucoidans especially related their bioactivity. They are known to show anti-tumor (Maruyama *et al.*, 2006; Alekseyenko *et al.*, 2007), anticoagulant (Zoysa *et al.*, 2007; Rodrigues *et al.*, 2011), anti-inflammatory, antithrombotic (Cumashi *et al.*, 2007; Patel, 2012; Dore *et al.*, 2013), antioxidant (Yuan *et al.*, 2005; Lim *et al.*, 2014; Wang *et al.*, 2015), antiviral (Bouhlal *et al.*, 2011; Elizondo-Gonzalez *et al.*,

2012; Wang *et al.*, 2012; Rabanal *et al.*, 2014) and anticancer properties (Villarreal-Gómez *et al.*, 2010; Fedorov *et al.*, 2013; Marudhupandi *et al.*, 2014; Wang *et al.*, 2015).

Table 1-2 Chemical composition of fucoidan from several brown algae.

Brown Algae	Chemical Composition	References		
Fucus vesiculus	Fucose, sulfate	Black et al. (1952),		
		Nishino <i>et al.</i> (1994)		
F. evanescens	Fucose/sulfate/acetate	Bilan <i>et al.</i> , (2002)		
	(1/1.23/0.36)			
	Fucose, xylose, galactose,	Cumashi et al., (2007)		
	uronic acid and sulfate			
F. distichus	Fucose/sulfate/acetate	Bilan <i>et al.</i> , (2004)		
	(1/1.21/0.08)			
	Fucose, xylose, galactose,	Cumashi et al., (2007)		
	uronic acid and sulfate			
F. serratus	Fucose/sulfate/acetate (1/1.12)	Bilan et al., (2006)		
Lessonia vadosa	Fucose/sulfate (1/1.12)	Chandi et al., (2008)		
Macrocytis pyrifera	Fucose/galactose (18/1), sulfate	Black et al., (1952)		
Turbinaria ornate	Fucose/galactose (3/1), sulfate	Than et al., (Thanh et		
		al., 2013)		
Undaria pinnatifida	Fucose/galactose (1/1.1), sulfate	Lee et al., (2004)		
(Mekabu)				
Ascophyllum nodosum	Fucose (49%), xylose (10%),	Percival <i>et al.</i> , (1968)		
	GlcA (11%), sulfate			
Himanthalia lorea and	Fucose, xylose, GlcA, sulfate	Mian <i>et al.</i> (1973)		
Bifurcaria bifurcate				
Padina pavonia	Fucose, xylose, mannose,	Hussein et al., (1980a,		
	glucose, galactose, sulfate	1980b)		
Laminaria angustata	Fucose/galactose/sulfate (9/1/9)	Kitamura <i>et al.</i> , (2014)		
Ecklonia kurome	Fucose, galactose, mannose,	Nishino <i>et al.</i> , (1994)		
	xylose, GlcA, sulfate			
Sargassum stenophyllum	Fucose, galactose, mannose,	Duarte <i>et al.</i> , (2001)		
	GlcA, glucose, xylose, sulfate			
S. muticum	Glucose, xylose, galactose,	Balboa et al., (2013)		
	mannose, fucose			
S. polycycstum	Glucose, xylose, rhamnose,	Sugiono et al., (Sugiono		
	fucose, galactose, mannose	et al., 2014)		
S. wightii	Fucose, sulfate	Marudhupandi and		
		Kumar, (2013)		

Marinova – a Tasmanian company – has managed to isolate fucoidans from a range of species and provide characterized fractions for either investigational research or as ingredients for nutraceutical and cosmetic applications. They have successfully isolated and supplied commercial volumes of

fucoidan extracted from *Undaria, Lessonia, Macrocystis, Cladosiphon, Durvillea, Laminaria, Ecklonia, Fucus, Sargassum, Ascophyllum* and *Alaria* with Halal- and Kosher-certified (Holdt and Kraan, 2011).

# 1.5.3 Storage metabolites of brown algae

Laminaran, also called laminarin, is carbohydrate reserve of brown algae located in plastids of each cells. Amount of laminaran varies up to 34% of dry weight; it is preferably present in algae of the order Laminariales. Nevertheless, this carbohydrate reserve was also discovered in *S. fusiforme* and *S. trichophyllum* in low amounts, less than 1% (Rioux and Turgeon, 2015).

Laminaran is a polymer made of glucose units, varying in structure and composition that depens on the species. It is basically composed of chains of 1,3-β-D-glucose units (Holdt and Kraan, 2011; Rioux and Turgeon, 2015), a linear polysaccharide made up of 25-50 glucose units. The terminal reducing end of laminaran is diverse, corresponding to a glucose residue in G-type laminarins and to a mannitol residue in M-type laminarins. This diversification generates two forms of solubility: soluble and insoluble, for each type of laminaran. First characteristic of laminaran is a complete solubility in cold water, while second is only soluble in hot-water (Ruperez *et al.*, 2002). Laminarans have an average molecular weight of 5 kDa. The molecular weight of soluble laminaran, extracted from *L. digitata*, is about 5.3 kDa, while the insoluble form of laminaran, obtained from *Laminaria hyperborea*, is about 3.5 kDa (Chapman, 1980).

Other carbohydrate reserve in brown algae aside from laminaran is mannitol, a six-carbon alcohol. Mannitol represents up to 30% of algal dry weight and it is accumulated mainly in *Laminaria* and *Saccharina* species (Holdt and Kraan, 2011). In *A. nodosum* and *Fucus* species, mannitol was produced with content lower than 16% of dry weight. Meanwhile, mannitol content content in *S. pacificum* (formerly known as *S. mangarevense*) was 14% of dry weight as reported by Zubia et al. (Zubia *et al.*, 2008). As reviewed by same authors, the mannitol content of *Sargassum* ranged from 1% to 12% of dry weight.

Laminaran and mannitol content vary considerably according to algal species, age and part of the thallus, and seasons. Meanwhile, the extraction of these reserve polysaccharides is usually with acid solution (Rioux and Turgeon, 2015). Laminaran has been reported to possess various biological activities. Laminarin of *A. nodosum* and *L. hyperborean* demonstrated an antioxidant and antibacterial activities (Kadam *et al.*, 2015c). Antiviral, anticoagulant, and antiproliferative activities were also reported from the laminaran (Hoffman *et al.*, 1995; Miao *et al.*, 1999; Devillé *et al.*, 2007). As for mannitol, the bioactivity reported was its efficacy in protecting the photosynthetic apparatus from low-salinity damage (Lobban and Harrison, 1997).

#### 1.5.2 Proteins

Proteins of marine algae have attracted less attention and their structure and biological properties are not as widely documented as their polysaccharides (Chojnacka *et al.*, 2012). However, recently, they have gained increasing attention due to their functional properties and bioactivity that is potential for further application (Michalak and Chojnacka, 2015).

The content of brown algae protein is generally low ranging between 6-13% of dry weight compared to the green (15-25%) and red algae (15-30%) (Kim, 2012; Balboa *et al.*, 2013). However, their content in different species is dependent on season and is linked to their surrounding variables including nutrient supply. They are present in a diverse range of forms and cellular locations, i.e as a component of cell wall, enzymes and bound to pigments and carbohydrates (Stengel *et al.*, 2011). Table 3 displays protein content of algae from different groups.

Table 1-3 Protein content (% of dry weight (w/w)) of algae according to Harnedy and Fitzgerald (2011) (modified).

Algae	Protein content (% of dry weight)	Reference
Undaria pinnatifida	11-24	Fleurence, (2004)
Laminaria digitata	15.90	Marsham <i>et al.</i> , (2007)
L. saccharina	6-11	Morrissey et al., (2001)
Fucus vesiculosus	5-10	Morrissey et al., (2001)
F. serratus	17.4	Marsham et al., (2007)
Ascophyllum nodosum	5-12	Morrissey et al., (2001)
	3-15	Fleurence, (2004)
Himanthalia elongata	6-11	Morrissey et al., (2001)
Sargassum variegatum	3.0	Ambreen et al., (Ambreen et al.,
		2012)
S. corrifolium	16.07	Haque et al., (Haque et al., 2009)
S. filipendula	5.6	•
S. vulgare	13.6	Marinho-Soriano et al., (2006)

# **1.5.3 Lipids**

Brown algae contain relatively lower contents of lipids having less than 5% of dry weight. Quantitative analysis revealed that total lipid content of a major brown algae family, Sargassaceae, was higher in subarctic zones (approximately 5% of dry weight) than tropical zones (0.9-1.8% of dry weight). In spite of their low total lipid content, brown algae are rich source of polyunsaturated fatty acids (PUFAs) (van Ginneken *et al.*, 2011). Fatty acids are the main constituents of fat raw material in living organisms distinguished into saturated fatty acids (SAFA) and unsaturated fatty acids. The unsaturated fatty acids are later known as two major groups, namely monounsaturated fatty acids (MUFAs) and PUFAs (Kulikova and Khotimchenko, 2000). PUFAs generally contain high levels of omega-3 (ω3) and omega-6 (ω6) (Holdt and Kraan, 2011). *Sargassum* is rich in fatty acids with 20 carbon atoms, such as eicosapentaenoic acid (EPA, ω3 C20:5) and arachidonic acid (AA, ω6 C20:4). Certain species shows different content of these fatty acids. As an example, *S. kjellmanianum* is a rich source of ω6; whereas *S. thunbergii* has high content of ω3 (Burtin, 2003b).

Table 1-4 Total lipid content in some brown algae.

Algae	Total lipid content (% of dry weight)	Reference
U. pinnatifida	5	(Billakanti et al., 2013)
S. miyabei	0.7	(Kulikova and
		Khotimchenko, 2000)
S. horneri	0.82-0.96	(Hossain et al., 2003)
Alaria esculenta	1.3-1.5	
L. digitata	0.8-1.1	
L. hyperbora	1.1-1.4	(Maehre et al., 2014)
F. vesiculosus	2.6-3.5	
Pelvetia canaliculata	3.7-5.8	
A. nodosum	4.5	(man Cinnalson et al
F. serratus	3.7	(van Ginneken <i>et al.</i> ,
S. natans	1.0	2011)

### 1.5.4 Minerals

Marine algae have the capacity to absorb inorganic substances from their surroundings, because of this capacity, they are acknowledged as a significant source of essential minerals (Na, K, Ca, and Mg) and trace elements (Fe, Zn, Mn, and Cu) (Misurcova, 2012; Peng *et al.*, 2015). Algae's absorption capacity is regarded with the presence of polysaccharides in algal cell wall. Different type of polysaccharides also becomes the reason why various group of algae demonstrate differ ability to uptake minerals. Capacity of brown algae to absorb metal is higher than green and red algae due to the high presence as a result of high presence of alginic acid, alginate and salt of alginic acids. These polysaccharides of brown algae have strong ion-exchange properties. Moreover, cellulose and carboxyl groups coexisted in brown algal cell wall contribute to the accumulation of metals as well (Hu *et al.*, 1996). It is highly possible that contents of accumulated minerals in marine algae are higher than those of in the environment by several orders of magnitude (Peng *et al.*, 2015).

Minerals content in algae vary, depending on diverse genera of algae, season, and geographic location. Brown algae, particularly from the genera *Laminaria*, are a great source of iodine. The highest concentration of iodine is in *L. japonica*, 5.6 and 3.0 mg/kg from all investigated algae (Misurcova, 2012).

Table 1-5 Mineral contents in some brown algae (Fajardo *et al.*, 1998; Kolb *et al.*, 2004; Santoso *et al.*, 2006; Manivannan *et al.*, 2008; Misurcova *et al.*, 2009; Misurcova, 2012).

Algae	Ca	Mg	P	Na	Fe	K
L. japonica	8.8 g/kg	5.5 g/kg	3.0 g/kg		73.8 mg/kg	
U. pinnatifida	9.5 g/kg	4.1 g/kg	4.5 g/kg	36.1 mg/kg	133 mg/kg	71.2 g/kg
S. polycystum	18.7 g/kg	5.7 g/kg			277 mg/kg	
P. australis	28.3 g/kg	4.0  g/kg			446 mg/kg	

### 1.5.5 Polyphenols

Phlorotannins, are structural classes of polyketides found exclusively in brown algae and classified into six group based upon variations in their assemblage from the polymerization of phloroglucinol units (Arnold and Targett, 2000, 2002). Stored within cells in vessels called physodes, phlorotannins have been reported from almost all brown algae orders. According to Vo *et al.* (2012), phlorotannins are classified in three primary groups based on their inter-phloroglucinol linkages. They are fucols, phlorethols and fucophlorethols.

Monomeric unit of phlorotannins, *i.e.* phloroglucinol, is proposed to be formed via the acetate-melanote (polyketide) pathway. Two molecules of acetyl coenzyme A are converted into malonyl co-enzyme A through the addition of carbon dioxide. A polyketomethylene chain is formed by three malonyl CoA units, and the polyketide species thus undergoes a Claisen-type cyclization reaction to yield a hexacyclic ring system. This hexacyclic ring, a triketide, undergoes tautomerisation to give phloroglucinol, which is thermodynamically more stable (Fig 1-13). An enzyme complex that converts acetyl CoA and malonyl CoA into the final product without giving any intermediates carries out this whole sequence of reactions. This enzyme complex is known to combine the polyketide synthase and polyketide cyclase activities (Achkar *et al.*, 2005). Study on *Ectocarpus siliculosus* reveals that type III pollyketide synthase (PKS III) catalyzes major step in the biosynthetic pathway of phlorotannins (Meslet-Cladière *et al.*, 2013).

Fig 1-13 Biosynthetic pathway of phlorotannins.

There is specificity within genera as seen with the fucols in *Fucus* and eckols in *Ecklona*. Fucols (Geiselman and McConnell, 1981), phlorethols, fucophlorethols, fuhalols, isofuhalols (Grosse-Damhues and Glombitza, 1984) and eckols differ in the number of the hydroxyl groups present and in their bond linkages. Among these structures, the fucols are attached by ether bonds. The phloroglucinol units are often esterified or acylated, and can dimerize or polymerize into larger units. These compounds also exhibit a broad range of size, thought to polymerize as they age and thereby grow larger. They are typically 10 – 100 kDa, although their range spans from as small as 126 Da to as large as 650 kDa (Boettcher and Targett, 1993; Targett and Arnold, 2001). Figure 1-14 illustrates the chemical structure of main groups of phlorotannins and Table 1-6 presents a summary of several brown algae derived-phlorotannins.

Ecklonia cava, E. stolonifera, E. kurome, Eisenia bicyclis, Ishige ookamurae, Sargassum thunbergii, Hizikia fusiformis, Undaria pinnatifida and Laminaria japonica are some marine brown algae that have been most reported for their phlorotannins as well as its health beneficial biological activities (Li et al. 2011). From Ecklonia cava, several studies managed to isolate their different type of phlorotannins, i.e. phloroglucinol, eckol, fucodiphloroethol G, phlorofucofuroeckol A, 7-phloroeckol, dieckol, and 6,6'-Bieckol (Jung et al., 2006; Yoon et al., 2009b; Wijesekara et al., 2010; Li et al., 2011a).

Phlorofucofuroeckol A was also found in *E. stolinifera* and *E. kurome* (Nagayama *et al.*, 2003; Yoon *et al.*, 2009b). *E. stolinifera* also contained triphloroethol B and 2-phloroeckol as reported by Nagayama *et al.* (Nagayama *et al.*, 2003). 6,6'-Bieckol was also isolated from *Ishige okamurae* (Zou *et al.*, 2008; Yoon *et al.*, 2009a). Eckol, phlorofucofuroeckol B, 6,8'-Bieckol, and 8,8'-Bieckol were successfully isolated from the *Eisenia bicyclis*. Ferreres *et al.* (2012) found 22 different phlorotannins belonging to eckol and fucophlorethol main groups from *Cystoseira nodicaulis*, *C. tamariscifolia*, *C. usneoides* and *Fucus spiralis*.

Fig 1-14 Chemical structure of main groups of phlorotannins characterized from several brown algae; (a) phlorethols, (b) fucols, (c) fucophlorethols, (d) eckols, (e) carmamols, and (f) fuhalols.

Table 1-6 Main group and sub-group of phlorotannins families in different species of brown algae.

Main group of phlorotannins	Sub-group		Species	Interphloroglucinol linkages and Molecular size	Reference
Fucols	Tetrafucol-A and B		Fucus vesiculosus L	With only phenyl linkages	Glombitza <i>et al.</i> , (1975); Truus <i>et al.</i> , (2004)
Phlorethol	r	and	Laminaria ochroleuca	With only arylether bonds	Koch et al., (1980)
	tetraphlorethols A & B Tetraphlprethol C		Ecklonia maxima (Osbeck) Papenfuss	-	Glombitza and Vogel (1985)
	Pentaphlorethol B Hexaphlorethol A	and	Cystophora retroflexa (Labilladiere) J. Argadh	-	Sailler <i>et al.</i> , (1999)
Fuhalols				Phlorethol with additional hydroxyl group	
	Isotrifuhalol		Chorda filum	-	Grosse- Damhues et al., (1984)
	Hydroxytrifuhalol Hydroxypentafuhalol	В, А,	Sargassum spinuligerium	-	Glombitza <i>et al.</i> , (1997a); Glombitza and Keusgen, (1995);
	hydroxyheptafuhalol Hydroxynonafuhalol A	В,			Keusgen <i>et al.</i> , (1997) Glombitza and Shmidt, (1999)
Eckols	J 1 J 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1			Phlorethols includes 1,4-dibenzodioxin system	
	Trimers eckol Dioxinodehydroeckol	and	Ecklonia maxima & Esenia arborea Areschoug	-	Glombitza and Vogel, (1985); Glombitza and Gerstberger (1985)

Table 1-6 Main group and sub-group of phlorotannins families in different species of brown algae (Continued).

Main group of phlorotannins	of Sub-group S	pecies	Interphloroglucinol linkages and Molecular	Reference
			size	
	Eckol, Phlorofucoeckol-A Dieckol, 6,6'-bieckol and 8,8' bieckol	,	-	Kim <i>et al.</i> , (Kim <i>et al.</i> , 2013); Fuji <i>et al.</i> (2013)
	Tetramers 2-phloroeckol and 7	- Eisenia bicyclis (Kjellman	) -	Shibata et al (2004, 2008); Kim et
	phloroeckol	Setchell, E. kurome Okamur		al, (2011); Kita et al., (1990);
		and <i>E. stolonifera</i> Okamura		Nagayama et al., (Nagayama et al., 2003); Kang et al., (Kang et al., 2012); Kim et al., (2005); Goo et al., (2010)
	Diphlorethohydroxycarmalol	Carpophyllum maschalocarpum		Li and Glombitza (1991)
	Triphlorethohydroxycarmalol	Ishige okamurae Yendo		Zou et al., (2008); Yoon et al., (2009a)
Fucophlorethols			With arylether and phenyl linkages	· · · · · · · · · · · · · · · · · · ·
	Fucophlorethols	Himanthalia elongat (Linnaeus)	e ~ 994 Da	Glombitza and Grosse, (1985)
	Quaterfucononaphlorethol	Carpophyllum maschalocarpum (Turner)	1.738 Da	Li and Glombitza (1991)
	Fucophlorethols	Sargassum spinuligerum and Cystophora torulosa (R Brown)		Glombitza <i>et al.</i> ,(1997a)
	Fucophlorethols-B	Laminaria ochroleuca		Koch et al., (1980);
	Fucodiphlorethol-C and G	Eisenia bicyclis, Ecklonic cava	a -	Martinez and Castaneda, (2013); Ham <i>et al.</i> , (2007);

Table 1-6 Main group and sub-group of phlorotannins families in different species of brown algae (Continued).

Main group of phlorotannins	Sub-group		Species	Interphloroglucinol linkages and Molecular size	Reference
	Trifucodiphlorethol A		Fucus vesiculosus		Parys <i>et al.</i> , (2010)
	Bisfucotriphlorethol	A,	Cystophora torulosa	-	Glombitza et al. (1997b)
	Bisfucoetraphlorethol	A,			
	Bisfucopentahlorethol	A,			
	Hydroxyfucodiphlorethol	В,			
	Hydroxyfucotriphlorethol	В,			
	Bisfucotriphlorethol,				
	Hydroxybisfucopentaphloret	hol			
	A and Fucodifucotetraphlore				

## 1.5.5.1 Role of phlorotannins in brown algae

Phlorotannins are commonly characterized as chemical protective defense, which also in the case of this algae, exhibit primary functions for example in growth and the development of the cell wall (Schoenwaelder and Wiencke, 2000). Koivikko (2008) stated that the links between cell wall (alginic acid) and phlorotannins are the ester bond and the hemiacetal bond, both of which are the covalent bonds. Therefore, it makes them hard to degrade.

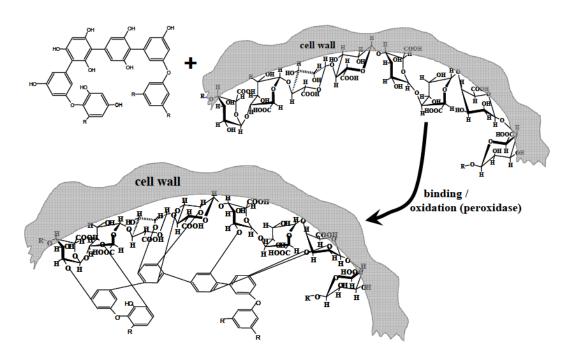


Fig 1-15 Illustration of phlorotannin polymer covalent bond at the alginic acid cell wall of brown algae as proposed by Koivikko (2008).

Apparently, phlorotannins also play another role in the defense mechanisms of algae towards any stress or changing in their surroundings, thus they are also known as stress compound (Stengel *et al.*, 2011). Phlorotannins are involved in a protective mechanism for examples against grazing (Koivikko *et al.*, 2005; Coleman *et al.*, 2007; Ank *et al.*, 2013), attachment and settlement of bacteria or other epibiosis effect (Mokrini *et al.*, 2008; Ank *et al.*, 2013), ultraviolet (UV) radiation (Macaya *et al.*, 2005; de la Coba *et al.*, 2009), metal

contamination (Connan and Stengel, 2011; Warneke and Long, 2015).

# 1.5.5.2 Phlorotannins content in brown algae

The total content of phlorotannins in brown algae ranges from < 1 – 14% of dry weight and has been shown to depend on environmental factors such as salinity, light and nutrient availability (Hemmi *et al.*, 2005; Jormalainen and Honkanen, 2008; Holdt and Kraan, 2011; Le Lann *et al.*, 2012). Variation of Polyphenol content seems to occur among local populations of algae (Pavia *et al.*, 1999; Hemmi and Jormalainen, 2004; Koivikko *et al.*, 2008). Such variation can be explained by genetic differentiation among populations and/or response to local changes of environmental factors (Koivikko *et al.*, 2008).

Spatial variation in levels of brown algae polyphenol compounds has been examined across very large spatial scales, such as between hemispheres and between temperate and tropical habitats (Stiger *et al.*, 2004; Le Lann *et al.*, 2012). Variation in polyphenol levels has also been examined across regional scales (tens to hundreds of km) for several species of Australasian brown algae (Steinberg, 1989), *Lobophora variegata* in Caribbean and Atlantic waters (Arnold *et al.*, 1995), *Fucus vesiculosus* in Atlantic waters (Targett *et al.*, 1992), and *Ascophyllum nodosum* in Europe (Pavia and ÅBerg, 1996). Steinberg (1989) reported that polyphenol levels of Australasian kelps and rockweeds often differed significantly among sites, but within a species the highest polyphenol concentration was generally less than twice the lowest concentration. Table 1-7 presents the seasonal variation of brown algal phlorotannins. In addition, Table 1-8 summarizes the polyphenol content with other biochemical composition from several species of brown algae.

Table 1-7 Polyphenol contents variation in different species of brown algae.

Species	<b>Polyphenols Content</b>	Type of variation	References
Cystoseira tamariscifolia	~ 2 % in winter	Seasonal	Abdala-Diaz et al., 2006
	~ 8 % in summer		
Stypopodium zonale	~ 0.5% (without herbivore)	Simulated herbivory	Ank et al., 2013
	~ 0.7 % (with herbivore)	·	
	~ 0.6 % (low epibiosis)	Epibiosis effects	
	~ 0.6% (medium epibiosis)	•	
	~ 0.4% (high epibiosis)		
Fucus serratus	4.27 % in lower intertidal zone	Interspecific	Connan et al., 2004
Ascophyllum nodosum	~ 5.8 % in lower intertidal zone	•	
Bifurcaria bifurcate	3.73 % in lower intertidal zone		
Himanthalia elongate	2.17 % in lower intertidal zone		
Laminaria digitata	0.13 % in subtidal zone		
G	3.88 % in upper intertidal zone		
F. spiralis	~ 5.8 % in upper intertidal zone		
F. vesiculosus	3.39 % in upper intertidal zone		
Pelvetia canaliculata			
L. digitata	0.09 % in summer	Seasonal	Connan et al., 2004
	0.19 % in winter		
F. vesiculosus	7.84 % in spring		
	~ 7% in summer		
H. elongate	0.49 % in winter		
-	4.84 % in summer		
P. canaliculata	3.72 % in summer		
	~ 3 % in winter		
A. nodosum	7.18 % in summer		
	~ 6% in winter		
Sargassum cristaefolium	0.08 %	Geographical location with	Le Lann et al., 2012
S. aquifolium	0.10-0.71 %	depth and reef morphology	
S. polycystum	0.09-0.74%		

Table 1-8 Polyphenol content and biochemical composition from several brown algae (expressed in % of dry material).

Species	Phenol	Carbohydrates	Protein	Ash	Reference
S. fusiforme	0.01	-	8.85	0.33	Shafay, (2014)
		17-69	11.6	19.77-26	Pereire, 2012
S . ilicifolium	1.93	38.72	38.72		Ganapathi et al., 2013
S. polycystum	0.23	34.93	7.78	21.87	Ahmad et al., 2012
	-	36.59	3.07	38.64	Triwisari, 2010
S. vulgare	0.38	93.34	5.85	0.19	Shafay, 2014
S. wightii	2.17	-	6.40	-	Seenivasan et al., 2012
Turbinaria ornate	$7-8 \times 10^{-3}$	14	15	-	Phartiban et al. 2013
Dictyota dichotoma	0.01 - 0.02	10	10		
T. conoides	0.34	41.03	7.40	21.37	Ahmad et al., 2012
Hormophy sacuneiformis	0.15	40.57	6.42	26.81	Ahmad et al., 2012
Padina gymnospora	0.25	26.86	5.93	45.04	Ahmad et al., 2012
	32.3	-	10.5	-	Rameshkumar et al., 2012
Cystoseira barbata	1.05	-	5.60	14.24	Frikha <i>et al.</i> , 2012

#### 1.5 Extraction of algal bioactive compound

Samples preparation is used to increase the efficiency of an analysis, to eliminate or reduce potential interference, to enhance the sensitivity of the analysis by increasing the concentration of the compound of interest in the assay mixture and to transform it into a more suitable form that is easily separated, detected and/or quantified. The sample produced in this step should have a high concentration of target compound free of interfering compounds from the matrix. One of the key procedures in this step is the extraction of the target compound (Palma *et al.*, 2013).

Extraction is a preliminary treatment of the analytic protocol in which a compound or group of compounds is preferentially transferred from a matrix into a different phase. The objective is making the sample available for introduction into analytic instruments either for qualitative or quantitative analysis. However, the difference between this two analyses is in qualitative the achievement of the exhaustive extractions and complete compound stability are ignored. In other word, they are not regarded as important. In the contrary, when dealing with quantitative analysis, it is highly important to achieve the complete extraction of the target compounds while preserving their original profile (Palma *et al.*, 2013).

The principal of the extraction process are related to several important properties, i.e. (1) high yield, (2) high selectivity/purity with low amount of interfering or undesirable co-extracted compounds, (3) high sensitivity allows for different quantification techniques producing a high slope in the calibration curve, (4) low limit of detection thus components in the extracts obtained can be detected at low levels (Palma *et al.*, 2013).

In general, extraction is a mass transfer process of one or more components from one phase to another one. In the most case of natural products, the samples used in the extraction process are solid materials, although in some cases liquids samples are also used. The extraction solvent is usually a liquid, but it can also be a solid or a supercritical fluid (Palma *et al.*, 2013). In understanding

the basic mechanism of extraction in any different techniques of extraction existed, both the target compounds and the extraction solvent applied must be considered. In addition, it is crucial to know the properties of the solvent medium in which the samples are to be dissolved. As a general rule, non-polar solvents dissolve non-polar solutes while polar solvents dissolve polar solutes. The interactions between solute and solvent are determined by the vapor pressure of the solute, the solubility of the solute in the solvents, the hydrophobicity, and the acid/base properties of both solute and solvents ((Palma *et al.*, 2013).

The basic mechanism of extraction as mentioned above is schematically presented in figure with following steps (Palma *et al.*, 2013):

- 1. The solvent is transferred from the fluid phase to the solid surface and pervades it,
- 2. The solvent penetrate into the solid matrix by molecular diffusion,
- 3. The soluble material is solubilized by desorption from the matrix into the extraction solvent the breakage of chemical bonds may be required for desorption of the target analytes from solid matrix,
- 4. The solution containing the solutes returns to the surface of the solid by molecular diffusion,
- 5. The solution is transferred from the solid surface to the bulk fluid by natural of forced convection.

The method of extraction applied to particular solid matrix depends on the raw material to be processed and on the final product of interest. Some types of extraction techniques have existed. The most commonly used conventional techniques are soaking extraction, Soxhlet extraction and distillation. Each one of them presents different advantages and disadvantages, thus there is no single and standard extraction technique for extracting the bioactive compounds from natural products (Palma *et al.*, 2013).

Besides the technique, the efficiency of extraction is also function of the process conditions. Certain external factors affect the concentration of the desired compounds in the extract such as solvent type, temperature, solvent to feed ratio, contact time, and particle size (Palma *et al.*, 2013). Therefore, the best extraction method with optimized conditions should be performed for preparing each particular product. Nevertheless, Cowan (1999) stated that the extraction efficiency of any conventional method mainly depends on the choice of the solvents. The polarity of the target compound is the most important factor for solvents choice. Azmir *et al.* (Azmir *et al.*, 2013) added that molecular affinity between solvents and solute, mass transfer, use of co-solvent, environmental safety, human toxicity and financial feasibility should also be considered in selection of solvent for bioactive compound extraction.

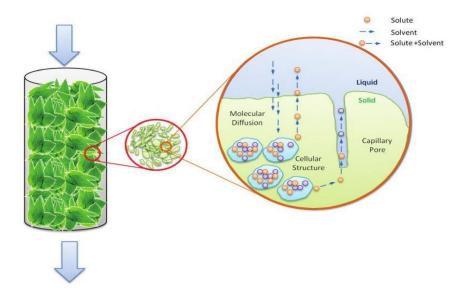


Fig 1-16 Basic mechanism of extraction based on Palma et al. (2013).

# 1.6.1 Algal model

Most of published studies concerning the extraction of marine algae natural product involved the conventional extraction method, i.e. solvents extraction consisting of liquid-liquid extraction (LLE), solid-liquid extraction (SLE) and extraction in Soxhlet apparatus (Michalak and Chojnacka, 2014). The main principal of this method is to extract the target compound by using the appropriate

solvents. Under this circumstance, the extraction yield varies from 8% to 30% of algal dry material (Robic *et al.*, 2009). This is due to the presence of various polysaccharides in large quantities in their cell wall and some compounds attached to the cell wall (cell-wall-bound compounds). Thus, it strongly reduces the efficiency of extraction. In addition, some organic solvents used for the extraction are often toxic and they result a great amount of waste after the process has finished.

Extraction of bioactive compounds is mostly performed by conventional extraction, such as solid-liquid extraction. Recently, focus is given to the production of algal extract without using the toxic organic solvents or destructive extraction conditions that could deteriorate biologically active compound from algal cells. A trend involving safer or even less solvents, more efficient, higher extraction yields compared to the conventional method and perhaps if possible with shorter time and lower cost. In order to be able to develop the application of marine natural products derived from marine algae in the future, an environmentally friendly and sustainable trend should be followed.

There are different methods of extraction that have been applied to obtain bioactive compounds of algae, like polysaccharides, polyphenols, proteins, lipids and pigments (Azmir *et al.*, 2013; Kadam *et al.*, 2015a). Figure 1-17 illustrates different methods of extraction that have been used to extract bioactive compounds in algae. Table 1-9 summarizes the application of different extraction methods to obtain bioactive compounds from algal material. The expected outputs of these new techniques of extraction are higher extraction yields, specificity and efficiency of bioactivity by preserving the original efficacy of active compounds, mild reactive conditions, shorter time, less solvents, better products and lower cost (Tanniou *et al.*, 2013a). Various type of extraction methods used to attain bioactive compounds in algae is summarized on Fig 1-17 (Jeon *et al.*, 2012; Kadam *et al.*, 2013; Michalak and Chojnacka, 2014).

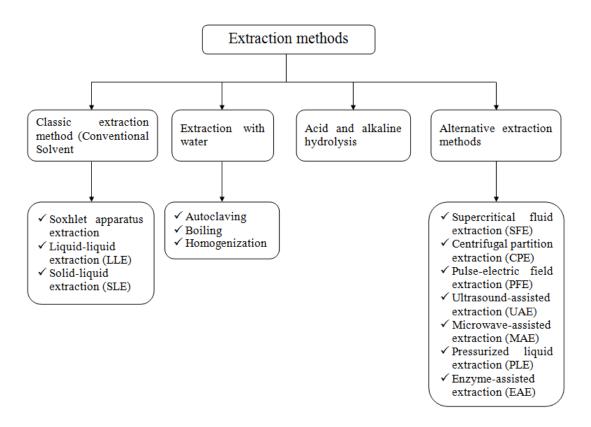


Fig 1-17 Various type of extraction methods used to attain bioactive compounds in algae (Jeon *et al.*, 2012; Kadam *et al.*, 2013; Michalak and Chojnacka, 2014).

Table 1-9 Examples of different extraction methods applied in the search of algal bioactive compounds.

Algae	Extraction method	Bioactive compound	Content	Bioactivity	References
S. muticum	<ul><li>SLE</li><li>CPE</li><li>PLE</li><li>SC-CO<sub>2</sub></li></ul>	Polyphenol	<ul> <li>18.5%</li> <li>10%</li> <li>22%</li> <li>3%</li> </ul>	Antioxidant	(Tanniou et al., 2013a)
Stypocaulon scoparium	SLE	Polyphenol	123-328 mg GAE/100g	Antioxidant	(Lopez et al., 2011)
Dictyopteris polypodioides	Acid and alkaline hydrolysis	<ul><li>Alginate</li><li>Mannuronan</li></ul>	<ul><li>0.92 and 1.25 g</li><li>0.75 and 1.15 g</li></ul>	-	(Karaki <i>et al.</i> , 2013)
S. filipendula	Alkaline hydrolysis	Sulfated polysaccharide	41-66%	Antioxidant	(Costa et al., 2011)
Gracilaria tenuistipitata	Water extraction	Polyphenols Flavonoid Ascorbic acid	98.9 μg GAE/mg 22.6 μg QE/mg 1.6 μg AA/mg	Antioxidant	(Yang et al., 2012)
Halimeda monile	Water extraction	Polyphenols Carotenoids Terpenoids	179.5 μg/g 42.2 μg/g -	Antioxidant	(Batista-Gonzalez et al., 2012)
Micro- and macroalgae	SC-CO <sub>2</sub>	Neutral lipids Phospholipids Glycolipids Liposoluble pigments	-	-	(Crampon <i>et al.</i> , 2011)
E. cava	EAE	Polyphenol	901 μg/100 ml	Antioxidant and antiproliferative	(Athukorala <i>et al.</i> , 2006)

Table 1-9 Examples of different extraction methods applied in the search of algal bioactive compounds (continue).

Algae	Extraction method	Bioactive compound	Content	Bioactivity	References
S. muticum	EAE PLE	Polyphenol	6.1 – 10 mg GAE/g 21.8-39.2 mg GAE/g	Antioxidant	(Sanchez-Camargo et al., 2015)
A. nodosum	UAE	Polyphenol Fucose Uronic acid	143.1 mg GAE/g 87.1 mg/g 128.5 mg/g	-	(Kadam et al., 2015b)
S. muticum	EAE  UAE  Hot water extraction	Polyphenol Sulfated sugar Nitrogen Polyphenol Sulfated sugar Nitrogen Polyphenol Sulfated sugar Nitrogen	200.5-300.5 μg/g 5.9-9.7 mg/g 22.1-29.6 mg/g 2.35 μg/g 10.1 mg/g 24.4 mg/g 275.8 μg/g 9.5 mg/g 24.6 mg/g	Antioxidant Prebiotic a-glucosidase inhibitory	
Osmundea pinnatifida	EAE  UAE  Hot water extraction	Polyphenol Sulfated sugar Nitrogen Polyphenol Sulfated sugar Nitrogen Polyphenol Sulfated sugar Nitrogen Nitrogen	106.0-123.1 µg/g 38.6-59.8 mg/g 11.7-19.4 mg/g 103.7 µg/g 42.8 mg/g 10.3mg/g 118.6 µg/g 41.8 mg/g 12.7 mg/g	Antioxidant Prebiotic a-glucosidase inhibitory	(Rodrigues et al., 2015)

Table 1-9 Examples of different extraction methods applied in the search of algal bioactive compounds (continued).

Algae	Extraction method	Bioactive compoun	d Content	Bioactivity	References
		Fucoidan		-	(Rodriguez-Jasso et
		• Tota	1 •	3.6-	al., 2011)
F. vesiculosus	MAE	sugar	5.6%		
		• Sulf	ates •	6.1-	
			8.3%		
Undaria pinnatifida	Sonication	Hydrocarbons	40.4-93.7%	-	(Punín Crespo et al.,
	MAE		34.6-90.5%		2006)

## 1.6.2 Solid-liquid extraction (SLE) of polyphenols

Extraction is a technique to separate compounds based on their solubility. It is a preliminary treatment of the analytic protocol in which a compound or group of compounds is preferentially transferred from a matrix into a different phase. S/L extraction is concerned with the extraction of a soluble constituent from a solid by means of solvents. If the solute is uniformly dispersed in the solid, the material near the surface will be dissolved first, leaving a porous structure in the solid residue. The solvent will then have to penetrate this outer layer before it can reach further solute, and the process might become progressively more difficult and the extraction rate will fall. This is due to the chemical structure of the solid material that some of them might have a complex bond. Thus, it makes them hard to penetrate. Generally, the process can be considered in three parts: first the change of phase solute as it dissolves in the solvent, secondly its diffusion through the solvent in the pores of the solid to the outside of the particle, and last, the transfer of the solute from the solution in contact with the particles to the main bulk of the solution (Richardson *et al.*, 2002).

Since algae become important sources of remarkable bioactive compounds, these compounds have been extracted from various genera of algae. Polyphenols from brown algae are one of bioactive compounds that have been extensively studied for their outstanding antioxidant activity. Extraction of polyphenols is usually influenced by their chemical nature, extraction method employed, sample particle size, storage time and conditions and presence of the interfering substances (Naczk and Shahidi, 2004). Polyphenols extraction from brown algal material are mostly used solid-liquid extraction and preferable solvents applied are the aqueous ethanol or methanol solutions (Stiger-Pouvreau *et al.*, 2014). As previously described (see 1.5.5), polyphenol in brown algae coexists as complexes with polysaccharides and proteins in algal cell wall. In addition, as secondary metabolite, polyphenols are not the only compounds produced by algae as defense system (Macaya et al., 2005, Jormalainen and Honkanen, 2008). Therefore, polyphenol extracts from plant material are always a mixture of different classes

Additional steps may be required to remove the unwanted polyphenols and non-polyphenols constituents, solid-phase extraction techniques and fractionation based on acidity are commonly used to eliminate the unwanted constituents (Onofrejová *et al.*, 2010). It is also important to consider the temperature applied in extracting polyphenols as they are sensitive to high temperature. In most studies, 40 °C has been proposed as the most optimal and favorable condition for polyphenol extraction (Gironi and Piemonte, 2011; Tanniou *et al.*, 2013a; Hong *et al.*, 2014; Ismail and Abdullah, 2015).

Solubility of polyphenols is defined by the polarity of solvents, degree of polymerization of polyphenol, and interaction of polyphenols with other constituents and formation of insoluble complexes (Naczk and Shahidi, 2004). Table 1-10 shows some example of solvents that have been used to obtain polyphenol from brown algae.

Polyphenol compounds in brown algae, also known as phlorotannins, are considered as a pool composed by different nature and/or polarities since they are oligomers and polymers of phloroglucinol. SLE has been selected to extract polyphenol based on the solubility preference of phlorotannins towards the organic solvents. Therefore, different solvents might extract different group of phlorotannins. If the organic solvent is favorable for polyphenols, it is highly possible that the polyphenols can be extracted in large quantities. As for productivity in SLE of phlorotannins, polar solvents extract more polyphenols. In other word, the efficiency of extraction often decreases if the solvents are not appropriate. Aside from this, other disadvantage of SLE is time-consuming, and often uses large volumes of toxic organic solvents. In addition, the complex linkage of brown algal cell wall between polysaccharides, protein and phlorotannins becomes another hindrance in SLE. Thus, the efficiency of extraction is not as optimal as expected since SLE could not break the complex linkage of constituents in algal cell wall. However, it is likely possible that even the appropriate solvent has been chosen for phlorotannins extraction, other

bioactive compounds with same preference as phlorotannins might come along. As a result, it becomes a contaminant in the extractants and creates interference in the bioactivity. For this reason, it is also crucial to consider the suitable isolation method for phlorotannins.

Table 1-10 Application of polyphenol extraction using different organic solvents and their bioactivity potential in brown algae from previous studies.

Species	Solvents	Bioactivity Potential	References
S. muticum	Ethyl-acetate; Ethanol; Acetone; Methanol; Water	Antioxidant	Tanniou <i>et al.</i> , (2013b, 2014)
S. filipendula, S. duplicatum, S. crassifolium, S. binderi, Padina sp	Ethanol, Ethyl acetate, n-Hexane	Antioxidant	Bambang <i>et al.</i> , (2013)
S. polycystum, S. duplicatum, S. echinocarpum	Hexane, Ethyl acetate, Methanol	Antibacterial and antifouling	Widowati et al., (2014)
S. longifolium	Dimethyls sulfoxide (DMSO), benzene, acetic acid, hexane, chloroform, diethyl ether		Ponnanikajamid een <i>et al.</i> , (2014)
Ecklonia cava	Hexane, diethyl ether, acetone, ethyl acetate, acetonitrile, methanol and water	Antibacterial	Choi <i>et al.</i> , (2014)
Eisenia bicyclis	Methanol, hexane, dichloromethane, ethyl acetate, n- buthanol, water	Antibacterial	Lee <i>et al.</i> , (2014)
Fucus vesiculosus, Ascophyllum nodosum, F. serratus	Ethanol (61.7 %)	Antioxidant	Keyrouz <i>et al.</i> , (2011)
Cystoseira trinodis	Dichloromethane	Antioxidant	Sathya <i>et al.</i> , (Sathya <i>et al.</i> , 2013)
F. vesiculosus	Ethanol (80%)	Antioxidant	Wang <i>et al.</i> , 2012

## 1.6.3 Enzyme-assisted extraction (EAE) of polyphenols

Marine algae are well-known by their chemically complex and heteregenous structure, i.e. sulfated and branched polysaccharides associated with proteins and various bound ions including calcium and potassium (Jeon *et al.*, 2012; Wijesinghe and Jeon, 2012a). This cell wall structure is even more complex when it comes to the family of brown algae as they have a strong interaction between protein and phlorotannin (Stern *et al.*, 1996; Koivikko, 2008). This physical barrier has limited the efficiency of solid-liquid extraction technique of biologically active compounds from the marine algae (Wijesinghe and Jeon, 2012a). Thus, it is highly recommended to degrade this complex cell wall yet in the same time it does not have the negative side effect to the target compound during the process. Enzymatic degradation of cell wall polymers as proposed in the enzyme-assisted extraction has drawn many attention and become a more attractive alternative to chemical and mechanical processes (Wijesinghe and Jeon, 2012a). Table 1-11 displays the main differences between conventional and enzyme-assisted extraction.

Table 1-11 Differences between the conventional extraction technique and enzyme-assisted extraction.

Method of extraction	Advantage	Disadvantage	References
Conventional Extraction	Easy application for industrial scale	Longer extraction time Requirement of costly and high purity solvent Low extraction selectivity Thermo labile compounds Toxic solvents are often used Low bioactive yield	Michalak and ChojnackA (2014); Azmir <i>et al.</i> (2013)
Enzyme-assisted Extraction	High bioactive yield Facilitate the solubility of bioactive compounds Food grade enzymes are applicable Nontoxic, ecofriendly Preparation simplicity Moderate conditions Applicable for a large-scale operation	Not yet applicable in the industrial- scale because of the enzyme cost	Kadam <i>et al.</i> (2013); Puri <i>et al.</i> (2012); Wijesinghe and Jeon (Wijesinghe and Jeon, 2012a); Jeon and Wijesinghe (2011); Sander <i>et al.</i> (2009); Athukorala <i>et al.</i> (2006)

Applying the cell wall degradation enzymes has received many attentions. This application is considered as a promising technique in hydrolyzing the cell wall and the bound phytochemicals to enhance the release of that high-value compound. EAE has been introduced for the past few years as one of the environmentally friendly technique of extraction. This technique is performed under moderate conditions, which protects the biologically active compounds from degradation. Carbohydrases and proteases have been proven to break down the cell wall of marine algae and release the desired bioactive compound. Some of the commonly used enzymes are Viscozymes, Celluclast, Termamyl, Ultraflo, carragenananse, agarase, xylanase, Kojizyme, Neutrase, Alcalase Umamizyme (Ahn et al., 2004; Park et al., 2004; Heo et al., 2005; Wijesinghe and Jeon, 2012a). A list of some examples of enzymes as well as their optimal conditions used in EAE of bioactive compounds from brown algae is presented in Table 1-12.

Table 1-12 Summary of the most used enzymes in enzyme-assisted extraction of algal bioactive compound, particularly phlorotannins (Kim *et al.*, 2011; Wijesinghe and Jeon, 2012b).

Enzymes	Temperature	pН	Enzyme composition
Viscozyme	(°C)	4,5	Arabanase, cellulase, ß-glucanase, hemicellulase and xylanase
Celluclast	50	4,5	Group of enzymes catalyzing the breakdown of cellulose into glucose, cellobiose, and higher glucose polymer
Termamyl	60	6,0	Heat-stable α-amylase
Ultraflo	60	7,0	Heat-stable multiactive β-glucanase
Neutrase	50	6,0	Endoprotease
Flavourzyme	50	7,0	Endoprotease and exopeptidase acitivities
Alcalase	50	8,0	$\alpha$ -endoprotease

Enzyme-assisted extracts appeared to have a higher antioxidant activity compared to those obtained using conventional extraction methods (Park *et al.*, 2004). Prior to dimethyl ether extraction, enzymatic extracts of *Undaria pinnatifida* enhanced the yields of total lipids by 15-20% and fucoxanthin by 50%

(Billakanti *et al.*, 2013). EAE is also well suited to the extraction of phlorotannins and other polyphenol compounds from marine algae (Kim *et al.*, 2006; Siriwardhana *et al.*, 2008; Wang *et al.*, 2010), as it assists in breaking the complex bonding between polyphenol and proteins. Further, this technique can be applied for large-scale operations as a promising, an environmentally friendly and a feasible alternative to traditional extraction method for algal bioactive (Michalak and Chojnacka, 2014; Kadam *et al.*, 2015c). Besides algae, the recovery of polyphenols from the terrestrial plant has been reported, they are citrus peel (Li *et al.*, 2006), grape skin (Pinelo *et al.*, 2006), grape seed (Passos *et al.*, 2009), pomegranate (Mushtaq *et al.*, 2015), unripe apples (Zheng *et al.*, 2008), black currant (Landbo and Meyer, 2004) and cauliflower (Huynh *et al.*, 2014).

Degradation of algal cell wall has been considered as an important preliminary step for bioactive compounds extraction, including polyphenols. By breaking the complex linkage of polysaccharides in algal cell wall, it allows releasing the polyphenols and makes them more accessible to extract. Regarding the yield of extraction, this method results higher yield of extraction since the EAE is a key to "open a door", in this case the complex algal cell wall, and to liberate almost everything that is "kept inside the door", in other words phlorotannins. In order to make this method more efficient, it is strongly suggested to determine the suitable enzyme for brown algal cell wall.

Table 1-13 Application of enzyme-assisted extraction in brown algae phlorotannins and their bioactivity potential.

Algae	Enzymes	Activities	References
Ecklonia cava	AMG	Antiproliferative & Antioxidant	Athukorala <i>et al.</i> (2006)
E. cava, Ishige okamure, S. fulvellum, S. horneri, S. coreanum, thunbergii, Scytosiphon lomentaria	Viscozyme, Celluclast, Termamyl, Ultraflo, Protamex, Kojizyme, Neutrase, Flavourzyme, Alcalase	Antioxidant	Heo et al. (2003)
S. muticum	Alcalase and Viscozyme	Antioxidant	Sanchez-Camargo et al. (2015)
E. cava	Viscozyme, Celluclast, Termamyl, Ultraflo, Protamex, Kojizyme, Neutrase, Flavourzyme, Alcalase	Hydrogen-peroxide scavenging	(Kim et al., 2006)
E. radiata	Viscozyme, Celluclast, Ultraflo, Alacalase, Neutrase,Flavourzyme	Antioxidant	(Charoensiddhi et al., 2014)

#### 1.6 Upgrading of *Sargassum* – bioactive compounds or extracts

Sargassum exists in large number of species with more than 350 species (Guiry and Guiry, 2015) growing mostly anchored to the bottom with its holdfast with thalli floating in water column. Its fast growth supported by its vegetative reproduction and floating ability has made Sargassum a floating jungle in the ocean causing golden tides. As it gets washed away on to the shore in substantial amounts, Sargassum forms a mini-mountain piling up along the coast. Later, Sargassum creates a putrid sulfurous odor if its massive influx of biomass is left to decompose on the beach. Inundation of Sargassum on the coast occurs naturally since it relates to Sargassum life cycle. Nevertheless, the existence of Sargassum in massive amount can be disturbing for coastal communities. As example in the Caribbean, it spent at least \$120 million to clean up the massive influx of Sargassum (Milledge and Harvey, 2016). Instead of spending the amount for merely cleaning them from the coast, it might be better to valorize

Sargassum since many studies has revealed a vast array of potential in particularly for foods, pharmacology, and cosmetics industries (Holdt and Kraan, 2011; Rebours et al., 2014). Pharmacological properties of Sargassum crude extracts or isolated pure compounds have been widely recognized including antibacterial, anticoagulant, antioxidant, antifungal, antiviral, antimelanogenic, antiinflammatory, hypoglycemic, hepatoprotective and neuroprotective activities. This suggests that Sargassum is a rich source of health and promoting agents. Prior to these exceptional findings, Sargassum has been traditionally used as medicine in the ancient times in China to treat goitre. Sargassum has been mentioned as "Hai Zao" and it is believed that Sargassum could soften hard lumps, dispel nodes, eliminate phlegm and induce urination in humans (Liu et al., 2012).

In traditional medicine of China, Sargassum has been formulated in more than 226 prescriptions to treat various diseases. S. pallidum (Turner) C. Argadh and S. fusiforme (Harvey) Setchell have been listed as treatments for arteriosclerosis, skin diseases, high blood pressure, hepatolienomegaly, neurosis, angina pectoris, chronic bronchitis and esophagitis. S. fulvellum (Turner), S. henslowianum C. Agardh, S. thunbergii (Mertens ex Roth) Kuntze, S. horneri (Turner) C. Agardh, commonly named as "Hai Qian" are another Sargassum species that are popular in traditional medicines and food ingredients in the south east region of China. Sargassum was also used exclusively for medicinal purpose at ancient times in Monte Verde, South America (Dillehay et al., 2008). Sargassum vulgare C. Agardh, and Sargassum thunbergii Yendo from Japan have been or are used as antihelmintics (Chapman, 1980; Kang et al., 2008). Beside medicines, Sargassum is also benefit in environmental application, for example U.S. has made a patent regarding the application of S. natans in sequestering gold from water (Kuyucak and Volesky, 1988). In food industry Japan, S. fusiforme also known as Hijiki is served as food, named as "vegetable for longevity". It is a side dish prepared by soaking and boiling the dried algae with water and served with soy sauce (Liu et al., 2012). S. polycystum, has been eaten as sea vegetables

in Asian countries such as Malaysia and Vietnam (Hong et al., 2007; Matanjun et al., 2009).

Records about traditional use of *Sargassum* in medicine have made this genus interesting to explore. Thus, *Sargassum* has been the most extensively as alternative natural resources with economic value. Identified bioactive meroterpenoids such as 2-methyl-6-(3-methyl-70x0-2,5-octadienyl)-1,4-benzoquinone and sargafuran (Kamei *et al.*, 2009) with significant antibacterial activity (2  $\mu$ g mL<sup>-1</sup> < MIC < 15  $\mu$ g mL<sup>-1</sup>) could provide scientific evidence for the traditional therapeutic claims of *Sargassum*. The pharmacological activity of *Sargassum* extracts and research of its bioactive constituents provide scientific evidence that underpins the traditional therapeutical claims made for *Sargassum* such as furuncle and abscesses (Liu *et al.*, 2012).

The iodine in Sargassum was believed to play an important role in therapeutical function. However, the role of iodine may have been overestimated and other bioactive metabolites may have made more significant contributions toward Sargassum's therapeutic claims (Liu et al., 2012). Research suggests that Sargassum may play a role as an immunomodulator (Yende et al., 2014). Antimalarial activity has been found Sargassum natans (Linnaeus) Gaillon al., (Orhan et2006). S. hemiphyllum lipophilic fraction of dichloromethane/methanol extract exerted potent in vitro antiplasmodial activity  $(IC_{50} = 2.8 \mu g \text{ mL}^{-1}; \text{ (Lategan et al., 2009)}). \text{ Samee et al. (2009) showed that}$ semipurified hydroethanolic extracts from Sargassum growing in Pakistan or China had an antihyaluronidase activity proportional to the phlorotannin contents of the ethyl acetate fractions. In addition, the IC<sub>50</sub> of the extract from Sargassum tennerimum was found to be similar to that of catechin. Sargassum in India is mostly exploited for its alginates production in hydrocolloid industries either for local or for export (Subba Rao and Mantri, 2006).

Table 1-14 List of important bioactive compounds from *Sargassum* species that have been extensively studied.

Bioactive compound	Reference
Polysaccharides	(Josephine et al., 2008; Sinha et al., 2010; Ji et al., 2011; Lee et al., 2011; Zhang et al., 2012; Dore et al., 2013; Yende et al., 2014)
Polyphenols  Lipids and fatty acids	(Glombitza et al., 1997a; Keusgen and Glombitza, 1997; Stiger et al., 2004; Santoso et al., 2005; Chandini et al., 2008; Ghereib, 2010; Budhiyanti et al., 2012; Tanniou et al., 2013a; Hardouin et al., 2013; Widowati et al., 2014; Perez-Lopez et al., 2014; Montero et al., 2015; Samudra et al., 2015) (Bhaskar et al., 2004; Bazes et al., 2009; Wiraswati and Warganegara, 2011; Gerasimenko et al., 2014; Balboa et al., 2016)
Pigments	(Heo and Jeon, 2009; Resita et al., 2010; Pangestuti and Kim, 2011; D'Orazio et al., 2012; Heo et al., 2012; Hu et al., 2015; Sivagnanam et al., 2015)
Proteins and amino acids Minerals	(Marín <i>et al.</i> , 2009; Peng <i>et al.</i> , 2013; Polo <i>et al.</i> , 2015) (Oyesiku and Egunyomi, 2011; Kalaivanan and Venkatesalu, 2012; Michalak and Chojnacka, 2013)

Among the bioactive compound contained in marine algae, polysaccharides and polyphenols are the two most studied especially related to their biological activity. In this study, phlorotannin, phloroglucinol derivatives from the brown algae functioning both as a primary and secondary metabolite compound will be the main subject. Brown algae are one of marine algae group well-known for their strong antioxidant property shown by this compound. However, other studies have also stated that this compound has a vast array of potential as antimicrobial, anticancer, antihypertensive, antidiabetic, antiallergic, etc (Vadlapudi, 2012; Wijesinghe and Jeon, 2012; Thomas and Kim, 2011; Wijesekara *et al.*, 2010).

The function of phlorotannins as stress compound has brought another

interesting point of view particularly related to the search of marine natural product. Allegedly, their ability to defense themselves against diverse environmental stress makes them to have number biological activities that can be further prepared as basic materials for the pharmaceutical and cosmeceutical industries. Further, these biological activities have been widely studied and one of the most outstanding activities possessed by phlorotannin is its antioxidant activity. It has been proven that brown algae phlorotannin has demonstrated as one of the strongest and important antioxidant activity (Ahn *et al.*, 2004; Shibata *et al.*, 2008; Girija *et al.*, 2013).

Strong correlation between the Polyphenol content and antioxidant activity in brown Algae has been reported by many previous studies (Wang et al., 2009; Rajauria et al., 2010; Eom et al., 2011; Namvar et al., 2013). Other studies have revealed that phlorotannins demonstrated a great potential against 1,1-diphenyl 1,2-picryhydrazyl (DPPH), hydroxyl, superoxide and peroxyl radicals (Li and Glombitza, 1991; Nakai et al., 2006; Santoso et al., 2012; Tierney et al., 2013). In general, the antioxidant properties work by inhibiting the initiation or propagation reactions by inactivating or scavenging free radicals. Therefore, it restrains or inhibits completely the lipid oxidation (Gupta and Abu-Ghannam, 2013). The phenol rings play their role as electron traps. Thus, they are responsible for the multifunctional antioxidant properties like the scavenging of hydroxyl radicals, peroxy radicals or superoxides. Further, their metal chelating potency appeared to be dependent upon their unique polyphenol structure and the number as well as location of the hydroxyl groups (Santoso et al., 2004). The chemical structure of algae phlorotannin is constituted up to eight interconnected rings (Wang et al., 2008). This factor explains why phlorotannins perform more potent antioxidant properties compared to the terrestrial polyphenols (Gupta and Abu-Ghannam, 2013) and make them as an interesting compound to further develop.

Apparently, phlorotannins do not only perform strong antioxidant but also antimicrobial property (Eom et al., 2013, 2012; Lopes et al., 2012; Choi et al.,

2010; Lee et al., 2010; Eom et al., 2008; Lee et al., 2008; Plouguerné et al., 2008; Lin et al., 2006; Ahn et al., 2004). Wang et al. (2009) reported the sensitivity of *Eschericia coli* to brown algae *Ascophyllum nodosum* phlorotannins and the terrestrial tannins. Their results demonstrated that the inhibition of E. coli growth of phlorotannins was greater than the hydrolysable and condensed tannins isolated from Rhus semialata and Schinopsis balansaei - both are terrestrial plants. The oxidation of tannins and the liberation of hydrogen peroxide become the factors that are responsible for antimicrobial properties of polyphenol compounds as mentioned by Smith et al. (2003). A different source of polyphenol produces different amounts of hydroxide peroxide under aerobic conditions. The highest production is associated with polyphenol that are rich in hydroxyl groups (Akagawa et al., 2003). In addition, the number of hydroxyl groups available to form hydrogen bonds with protein lead to the increase of tannins affinity to protein (Mueller and Harvey, 2006). Easily oxidized upon exposure to air and contain more hydroxyl groups are two characteristics shown by phlorotannins. These factors account for their greater antibacterial activity (Wang et al., 2009). Furthermore, the degree of polymerization of phloroglucinol also contributes to this outstanding antibacterial property (Nagayama et al., 2002) that does not exist in the terrestrial tannins (Wang et al., 2009). Despite the differences shown by the phlorotannins and terrestrial tannins, they seem to share similarity in the mechanisms of antimicrobial activity. Those mechanisms are inhibitions of extracellular bacterial enzymes, deprivation of substrate required for growth or direct inhibition of oxidative phosphorylation (Wang et al., 2009).

Besides those two potential, phlorotannins also perform other biological activity such as anti-proliferative and antitumor (Nwosu *et al.*, 2011; Yang *et al.*, 2010; Imbs *et al.*, 2008); antidiabetic (Nwosu *et al.*, 2011; Jung *et al.*, 2008); α-Glucosidase and α-Amylase inhibition (Lee *et al.*, 2008), angiotensin-I-converting enzyme (ACE I) inhibition (Jung *et al.*, 2006; Athukorala and Jeon, 2005), anti-inflammatory (Kim *et al.*, 2014; Kim *et al.*, 2010); matrix metalloproteinases (MMPs) inhibition (Kim *et al.*, 2006), hyaluronidase inhibition (Ferreres *et al.*,

2012; Shibata *et al.*, 2002), tyrosinase inhibition (Yoon *et al.*, 2009; Heo *et al.*, 2009; Kang *et al.*, 2004), phoroprotective (Cha *et al.*, 2012; Le Lann *et al.*, 2008; Heo *et al.*, 2009) and antiallergic (Lee *et al.*, 2008; Sugiura *et al.*, 2007).

Phlorotannin has attracted many attentions as it has a promising potential for further application in the pharmaceutical, cosmeceutical and food industries. One genus that has been well and most studied related to their phlorotannin is the *Ecklonia*, Genera of kelp belonging to the family Lessoniaceae. The phlorotannin from this genus, particulary from *E. cava*, *E. kurome*, *E. maxima*, has been successfully purified and characterized, for example phloroglucinol, eckol, dieckol, fucodiphloroethol, phlorofucofuroeckol etc. (Mwangi *et al.*, 2013; Yotsu-Yamashita *et al.*, 2013; Wijesikara *et al.*, 2010; Koivikko, 2008). Besides this Genera, the characterization of phlorotannin from other Genera of Phaeophyceae has also been successfully reported such as the *Cystoseira abies-marina* (Montero *et al.*, 2014), C. *tamariscifolia* (Hudson) Papenfuss, C. nodicaulis (Withering) M. Roberts, *C. usneoides* Linnaeus and *Fucus spiralis* Linnaeus from the Fucales (Ferreres *et al.*, 2012) and *Eisini bicyclis* (Kjellman) Setchell from the Eisenia (Fujii *et al.*, 2013).

Phlorotannins of *Sargassum* have also revealed their wide range of pharmacological properties. Many studies have revealed the promising potential of phlorotannins derived from this genus. *S. hystrix* (Harnita *et al.*, 2013; Budhiyanti *et al.*, 2011), *S. filipindeula*, *S. duplicatum*, *S. crassifolium*, *S. binderi* (Bambang et al., 2013), *S. polycystum* (Chan *et al.*, 2011), *S. myriocystum*, *S. plagiophyllum*, *S. ilicifolium* (Ganapathi *et al.*, 2013), *S. muticum* (Namvar *et al.*, 2013; Tanniou *et al.*, 2012), *S. boveanum* (Zahra *et al.*, 2007) are some examples of *Sargassum* species that have been reported for their potential antioxidant properties of their polyphenol compounds. However, only few studies have reported the characterization of phlorotannin from this genus, i.e fuhalols, deshydroxyfuhalols and fucophloroethols from *S. spinuligerum* (Glombitza *et al.*, 1997; Glombitza and Keusgen, 1995), bifuhalol from *S. ringgoldianum* (Nakai *et* 

*al.*, 2006), 2-(4-(3,5(dihydroxyphenoxy)-3,5-dihydroxyphenoxy) benzene-1,3,5-triol (DDBT) from *S. patens* (Kawamura-Konishi *et al.*, 2012).

# 1.7 Production and potential application of brown algae

Most of algal cultivation activities come from the red algae, Rhodophyceae. Major red algae species that are cultivated include *Kappaphycus* and *Euchema*, (primary raw materials for carrageenan), followed by *Gracilaria* (raw materials for agar) and *Poryphyra* (nori, for direct human consumption). Figure 1-19 presents the division of algal industrial application and most of algal dry materials are used for direct human consumption known as sea vegetables. Second place of algal application is for hydrocolloid and last is for other applications, like medical applications, fertilizer, animal feeds, and other biotechnological applications. In 2011, 83% algal production was designated for human consumption (Craigie, 2011). Afterwards, processed food industry, the primary market for algal hydrocolloids continue to grow rapidly. Hydrocolloids products from algae, i.e. agar, alginate and carrageenan, serve as texturing agents and stabilizers (Bixler and Porse, 2011).

The brown algae processing industry in Europe is divided into two main categories, (1) alginic acid production and (2) products for agriculture such as fertilizer and animal feeds (Frangoudes *et al.*, 2012). Colloids have been applied to multiple industrial applications, such as food processing, cosmetics and pharmaceuticals. Alginate, carrageenan and agar are widely known for their gelling, stabilizing and thickening properties. These products have been commercialized as additives with an average 1 million tons of wet algaes are used every year in order to extract these three compounds. In France itself, there are nearly 90 companies that process or sell algae-based products (Mesnildrey *et al.*, 2012). Hydrocolloid extraction and commercialization have contributed 75% of algae-based products. Meanwhile, for direct consummation of fresh or dried algae, algae production only contributes 1% with 19 species authorized for general consumption. As for the focus of French stakeholders interested in algae-

based products, the majority of cultivated algae is used for material purposes, for example in the specialty of chemical (cosmeceutical, nutraceuticals and pharmaceuticals) (Brinker and Sternberg, 2014).

In European level, France is in the second position after Norway based on their algae production in 2012 with 57 % and 14 % respectively. Approximately, 95 000 tons of fresh algae were collected. Every year, France produces between 40.000 and 70.000 tons of fresh algae (Kerlero de Rosbo and Bernard, 2014). Brown algaes contributed 95 % of the production mainly from the genus *Laminaria* – with total 75% consisted of *L. digitata* (60%) and *L. hyperborea* (15%) (BDI, 2012; Mesnildrey *et al.*, 2012).

The distinguished annual production of algae in France is due to the fact that their production still depends on the natural stock. Furthermore, as a four season country, natural biomass of algae is extremely seasonal-dependent. Aquaculture production only contributes 50 tons of algae with two principal species cultivated, i.e. *Undaria pinnatifida* (Wakamé) and *S. latisima* (Kombu) (Kerlero de Rosbo and Bernard, 2014). They are mainly harvested from the coast of Brittany, particularly in the Finistère and Côtes d'Armor (Kerlero de Rosbo and Bernard, 2014). Figure 1-17 describes the percentage of algae applications in France dominating by the food-processing industry, chemistry and microbiology as the main markets.

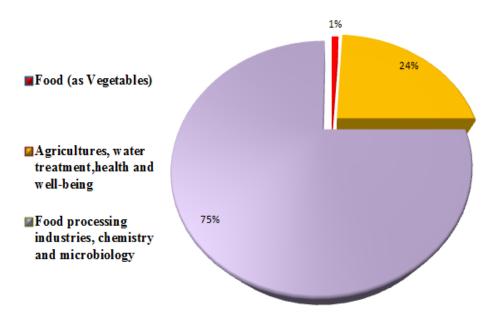


Fig 1-18 Classification of algae based on their utilization in France (French production and imports) according to CEVA (2009(Mesnildrey *et al.*, 2012)

There are 24 algae authorized for human consumption in France, in which 6 of them are particularly present in the littoral zone; kombu, wakame, nori, dulse, sea lettuce and bean (Figure 1-18). Table 1-12 displays the lists of marine algae that are allowed for human consumption (CEVA, 2014).

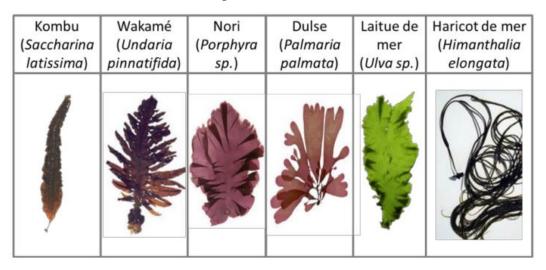


Fig 1-19 Macroalgae mainly consumed in France (CEVA, 2014).

Table 1-15 Macroalgae that are allowed for human consumption (CEVA, 2014).

Species	Common name
Brown algae	
A. nodosum	-
F. vesiculosus, F. serratus	-
Himanthalia elongate	"Haricot de mer" (bean)
U. pinnatifida	Wakame
L. digitata	Kombu
L. saccharina	Kombu royal
L. japonica	Kombu
Alaria esculenta	Atlantic wakame
Red algae	
Palmaria palmate	Dulce
Porphyra umbilicalis	Nori
P. tenera	Nori
P. dioica	Nori
P. purpurea	Nori
P. laciniata	Nori
P. leucostica	Nori
Chondrus crispus	Pioca
G. verrucosa	Ogonori
Lithothamnium calcareum	Maerl
Green algae	
Ulva sp.	"Laitue de mer" (sea lettuce)
Enteromorpha sp.	Aonori

In 2009, Indonesia was the second world producer contributing 16% of algae production, second consumer (13%) and first exporter of macro-algae by 32% of flow rate. 87% of its market was concentrated in the production of *Euchema* spp. In contrary with China and Japan, and similar to France, the Indonesian production of algae contributes the market with their colloids, particularly carrageenan used as stabilizer, emulsifier and gelling agents for food-processing industries (Kerlero de Rosbo and Bernard, 2014).

Algae production is mostly concentrated on Sulawesi, producing almost 65% of the total production volume. Most of the productions come from the exports of dry materials and the rest from the hydrocolloid production, especially carrageenan and agar-agar from the red algae. Neish (2012) stated that red algal galactan algae (RAGS) from the genus *Kappahycus/ Euchema* and *Gracilaria* are the most cultivated algae in Indonesia. In 2010, the yields of carragenophytes

reached 9.000 tons with different quality levels; 9% was refined and 21% was semi-refined for non-food purposes (Neish, 2012).

Basically, there seven main species that are commercially harvested, namely *K. striatus*, *E. spinosum*, *K. alvarezzi*, *Gracilaria sp.*, *Sargassum sp.*, *Gellidium sp.*, *and Caulerpa sp.* Predominant Indonesian algae crops are *K. alvarezzi* – commercial name "cottonii", and *K. striatus* – commercial name "sacol". About 50% of *Kappaphycus* crops are exported to China; and 20% are processed by domestic processors and the rest is exported to Philippines and other countries (Neish *et al.*, 2015). *K. alvarezzi*, *E. spinosum* and *Gracilaria sp.* are known to produce kappa- and iota- carrageenan, and agar. Accordingly, algal production in Indonesia is most likely designated for hydrocolloid production. For carrageenan, there are two types of products according to the production process, *i.e.* refined and semi-refined carrageenans, known as RC and SRC, respectively.

Almost similar to France's case, Sargassum species in Indonesia is also in the state of being underexploited. For local anglers, the application of Sargassum is still for animal's food (fodder) (pers. comm.) despite of the fact that alginates from brown algae are important substances for food, beverages, medical/pharmaceuticals, and non-food industries (Wouthuyzen et al., 2016). Previous studies have revealed that Sargassum species extracts or compounds from Indonesia have shown some pharmacological potential. As an example, alginate of Sargassum sp. in Indonesia has been acknowledged to enhance immune system in catfish *Clarias* sp (Isnansetyo *et al.*, 2014).

French and Indonesian *Sargassum* have been long considered as a potential natural resource with economic value. *Sargassum* is normally proposed as alginates sources for hydrocolloids industry as texturing agents. There have been many studies carried out regarding the prospect of *Sargassum* extracts in food, pharmacological and cosmetics industries beside the hydrocolloid industry. Results reveal the hidden yet remarkable pharmacological properties of *Sargassum* as it contains extremely active compounds that are possible for further

formulation. Moreover, this species might also be suggested as proteins, vitamins and minerals source for human consumptions. As can be seen from Table 1-12, *Sargassum* has not been included as one of macroalgae authorized for human consumption in France, though *Sargassum* has been consumed in certain countries such as in Japan. In order to authorize this species as a 'safely-consumed' product, it is highly essential to perform a series analysis required according to the regulations existed in each designated countries. Thus, *Sargassum* can be included as one of the authorized marine resources for commercial interest.

Others have started considering exploiting the massive biomass of Sargassum. For example, Japan has recognized Sargassum as source of sustainable bioenergy. Lipids and starch from Sargassum have been proven to be exploitable for bioenergy. Exploitation of Sargassum biomass is considered one of the best options for increasing renewable energy due to its massive influx along the coast. Bioethanol production from seaweed in Japan, entitled the "Ocean Sunrise Project," aims to produce seaweed-based bioethanol by farming and harvesting S. horneri, utilizing 4.47 million km<sup>2</sup> (the sixth largest in the world) of unused areas of the Exclusive Economic Zone (EEZ) and maritime belts of Japan (Aizawa et al., 2007). As Sargassum contains important proteins, fiber, macroand microminerals (especially in potassium, nitrates, calcium, iron and PUFAs), this species has been targeted as alternative source of animal feeds, fertilizer and biostimulants. Dry material of Sargassum has been integrated as supplements in animal feeds. It is believed that this species could improve the growth rate, survival rate and immune system of cultivated animals as reported in shrimp Litopenaeus vannamei, Fenneropenaeus chinensis, and rainbow Oncorhynchus mykiss (Huynh et al., 2011; Hafezieh et al., 2014; Sudaryono et al., 2015; Zamannejad et al., 2016). As fertilizer, algae have been applied to soils as manure and as conditioning agents (Hong et al., 2007). Extract of Sargassum also stimulates the germination, improve the growth and biochemical constituents and enhance the immunity of certain plants, like lentil (Vigna mungo), mung bean (Vigna radiata), mustard (Brassica juncea L.), garden cress (Lepidium sativium),

and tomato (*Solanum lycopersicum* L.) (Kavipriya *et al.*, 2011; Kalaivanan and Venkatesalu, 2012; Hernández-Herrera *et al.*, 2014; Siregar *et al.*, 2014; Godlewska *et al.*, 2016). Sargasso Sea Alliance has revealed that there are about 90 patents referencing *Sargassum*. Various industrial, medical and nutritional uses are proposed including applications focused on inhibiting HIV infection, as an antibiotic, antifungal and antifouling substance, biofuel, fertilizer and animal feeds (Laffoley *et al.*, 2011).

Table 1-16 Several studies on Sargassum bioactive compounds with pharmacological potential.

Species	Location	Solvent	Result	Reference
S. fillipendulla	Pulau Talango, Kab. Sumenep Madura	methanol, aceton, hexane, ethil acetate	Antioxidant activity and preservation for fish	(Khotimah <i>et al.</i> , 2013)
S. duplicatum	-	Ethanol, kloroform, ethil acetate	alternative therapy for IBD treatment	(Aulanni'am et al., 2011)
S. echinocarpum	Talango Island, Madura	Ethanol, methanol, acetone	Antioxidant activity	(Firdaus et al., 2012)
S. duplicatum	Pantai Raca, Nusa Kambangan	Ethanol, methanol, hexane, ethil acetate	Inhibition of manoldehide and peroxidose	(Septiana and Asnani, 2013)
S. crassifolium	Pulau Sumba	Methanol, hexane, dichlorometana, water	Antioxidant activity	(Supriyono, 2007)
S. hystrix	South Coast of Yogyakarta	Methanol, chloroform, deionized water, ethyl acetate	Candidate for photoprotection of oxidative damage to the DNA	(Harnita et al., 2013)

Table 1-16 Several studies on *Sargassum* bioactive compounds with pharmacological potential (continue).

Species		Location	Solvent	Bioactivity	Reference
S. crassifolium, S. polycystum, S. echinocarpum	S. S.	Craggy Coast of Gunung Kidul, Yogyakarta	alginate extraction	Potential to produce alginate	(Mushollaeni and S, 2012)
S. echinocarpum		Pari Island	alginate extraction	Potential to produce alginate as the viscosity reach the SIGMA's standar	(Rasyid, 2010)
r	S. S.	Teluk Awur, Jepara	Hexane, Ethyl acetate, Methanol	Antibacterial activity	(Widowati et al., 2014)
S. muticum		Grand Dellec, Plouzané	chloroform	Antibacterial, antifungal and anti-microalgal	(Plouguerne <i>et al.</i> , 2010)
		Cote-d'armor, Brittany	Alginate extraction and biomethane	Methane production	(Jard et al., 2013)
		Grand Dellec, Plouzané	Enzymatic hydrolysis	Anti-herpetic activity	(Hardouin et al., 2013)
		Northern and southwestern of Brittany	Methanol:water (50%; v/v)	Antioxidant activity	(Connan et al., 2006)
		Locmariaquer	Ethanol:water and ethanol:dichloromethane	Antifouling activity	(Bazes et al., 2009)

## **Chapter 2 Material and Methods**

## 2.1 Sampling Sites and abiotic factors

In this work, two locations, France and Indonesia, were chosen in order to compare the biochemical composition of *Sargassum* sp. extracts along with their biological activities. Illustration of sampling location is described in Fig 2-1.

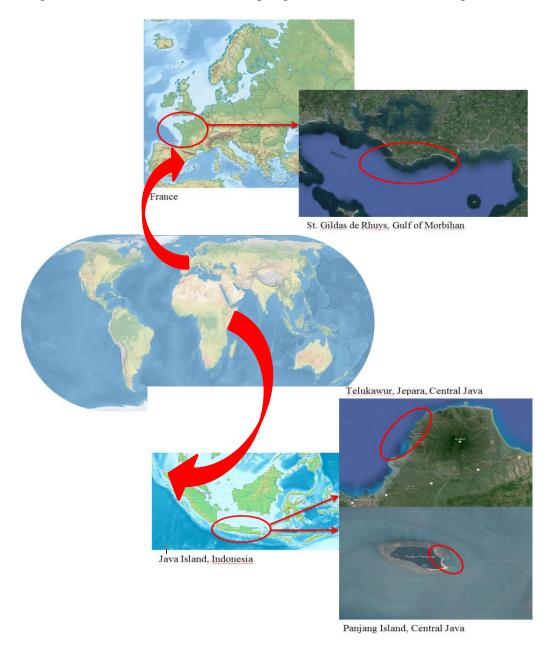


Fig 2-1 Overall view of sampling sites (Above) Saint Gildas de Rhuys, Morbihan, France; (Below) Java Island, Indonesia.

#### 2.1.1 Overview of studied location in France

The region of Brittany is located in the temperate climatic zone of the Northern hemisphere bordered with the English Channel – known as La Manche (in French) – on the North, south and west of Atlantic Ocean. This region is between two marine eco-regions; they are the Celtic Sea and Atlantic Continental Shelf of South Europe (Spalding *et al.*, 2007). As a consequence, this difference leads to a distinguished environmental conditions and fauna indicated between the coastal of North and South Brittany (Ehrhold *et al.*, 2006).

The coastal border of south Brittany is located on the northern continental shelf of Gascogne Gulf. It extends 150 km from east and west, between Guérande peninsula and the point of Raz bordered with a series of islands on the south (Houat, Hoëdic, Belle-Île, Groix and the archipelago of Glénans). The size of this area represents around 400.000 ha. The eastern side of South Brittany coast is bordered with the estuarine of Vilaine and Loire. From east to west, the zone includes four big bays, which are Vilaine, Quiberon, Concarneau and Audierne (Dutertre *et al.*, 2014).

Morbihan is one of the French departments within south Brittany, situated on the Northwest of France with 6.823 km<sup>2</sup> of total area. This department is surrounded by others departments namely, Finistère, Côtes-d'Armor, Ille-et-Vilaine and Loire-Atlantique with Atlantic Ocean on the southwest. Morbihan Gulf, an inland sea with 11.500 ha area widespread with a number of small islands, spreads to the North of Quiberon Bay. This area has a total size of 117.000 ha corresponding to "Mor-Braz" (Dutertre *et al.*, 2014).

Sargassum muticum (Yendo) Fensholt was collected monthly from February 2014 – February 2016 from the St. Gildas de Rhuys (47° 30′ 03″ N 2° 50′ 12″ W), one of the commune within the Morbihan department of South Brittany, France. Based on their position, the hydrology characteristic of this area is influenced by the Biscay Bay characteristic. Biscay Bay region ranges from the West Brittany (48°N) to the Strait of Gibraltar (36 °N). The continental shelf located in this Bay is about 140 km wide and it becomes narrower to the south

with about 50 km off the southern France. Sediments covering the continental margin mainly consist of thick turbidity sheet-fan deposits. In the south-western Brittany, the substrate is known as Grande Vasière – large muddy area.

S. muticum was hand-sampled by cutting the basal part of the thalli near the holdfast. They were rinsed with seawater after sampling and transported into the laboratory in closed boxes with no excess seawater. The collected samples were then rinsed thoroughly using tap water to remove the epiphytes and residues attached to samples. S. muticum was then grinded with Jupiter Universal (Wernau, Germany) using a 3 mm perforated grill. The grinded seaweeds were then lyophilized to obtain a dry material for further analysis and experiments.

In the north east of Biscay Bay, mainly in summer, upwelling events occur off Southern Brittany and the Landes coastline and may induce low-salinity (Koutsikopoulos and Le Cann, 1996; Puillat *et al.*, 2006). Temperature in this region varies between 7 and 15 °C in the sea surface and between 5.5 and 7 °C in the deep waters. The average salinity is about 35 % or higher in both surface and deep water.

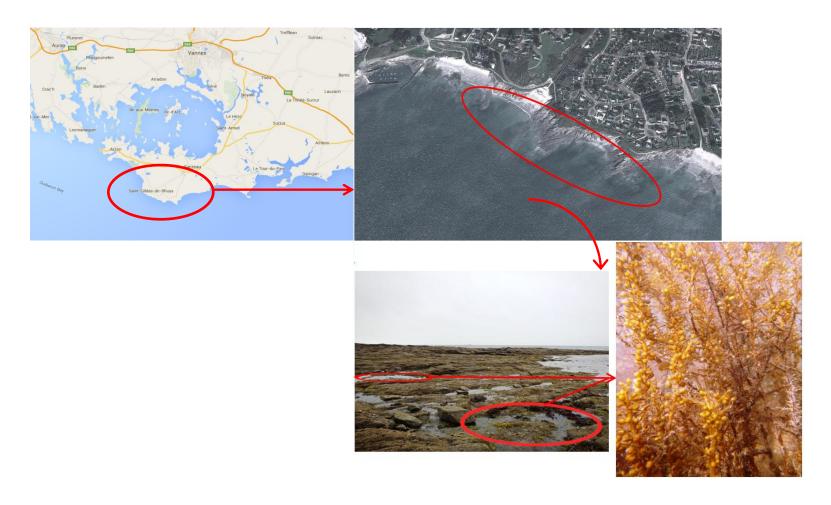


Fig 2-2 Overall illustration of sampling sites at St. Gildas de Rhuys, Vannes, France.

#### 2.1.2 Overview of studied location in Indonesia

The second location represented the tropical region is located in one of the main island in Indonesia, which is at Jepara, Central Java. Jepara is one of a city in the North of Central Java with 82.73 km<sup>2</sup> of coastal zone with total area covers 100.413,189 ha, located between 5°43'20,67" - 6°47'25,83" South and 110°9'48,02" – 110°58'37,40" East. This region extends up to the North where the Karimunjawa Islands is bordered by the Java Sea on the north and west side.

Jepara has a tropical climate with average air temperature between 21 and 32 °C with maximum precipitation (600 mm) occurring during the rainy season between January and March usually followed by the highest high tide. During this season, strong waves might reach 5 m high with wind speed 12-16 knots. The driest season happens on August with only 20 mm precipitation. Most of the substrate found in this region is fringing reef with white sands.

There were two different sampling sites, *i.e.* Panjang Island (06°34"30' South and 110°37"30' East) and Teluk Awur (6,1268° South 110,4° East) with 30 min away between these two sites. Both of them are located at Jepara, Java Central. *S. polycystum* was collected from Panjang Island, Jepara in October 2013 then *S. aquifolium* and *S.ilicifollium* were collected from Teluk Awur, Jepara in April 2014. All seaweeds were hand-sampled by cutting the basal part of the thalli near the holdfast. They were rinsed with seawater after sampling and transported into the laboratory in closed boxes with no excess seawater. The collected samples were then rinsed thoroughly using tap water to remove the epiphytes and residues attached to samples. *S. aquifolium*, *S. ilicifolium* and *S. polycystum* were air-dried and chopped to pieces and grinded using multiuse blender.

The water quality of Teluk Awur where *S. aquifolium* and *S. ilicifolium* were collected, and Panjang Island in Jepara, Indonesia where *S. polycystum* was sampled, was measured *in-situ* for its salinity, temperature, pH, and dissolved oxygen, and *ex-situ* for its nitrate and phosphate content. Details on water quality are presented in the table 2-1.

Table 2-1 Water quality of sampling sites based on *in-situ* measurements.

Water Quality	Teluk Awur	Panjang Island
Salinity (‰)	27	30.3
Temperature (°C)	28.2	31.6
рН	7.7	8.13
Dissolved oxygen (DO/ mg/l)	6.3	8.09
Phosphate (mg/l)	0.1	n.d
Nitrate (mg/l)	1.1	n.d

n.d: not determined



Fig 2-3 Overall view of sampling sites at Jepara, Central Java, Indonesia, (Left) Panjang Island, (Right) Teluk Awur.

## 2.2 Preparation of algal material

## 2.2.1 Species, sites and period of collection

Algal species used in this study was *S. aquifolium*, *S. ilicifolium*, *S. polycystum*, and *S. muticum*. Table 2.2 presents the detail of *Sargassum* species used in this study collected from different sites in different period.

Table 2-2 List of *Sargassum* species, location and period of collection from this study.

Species	Location	Region	Period of collection		
S. muticum	Saint Gildas de Rhuys, Morbihan	Temperate (France)	February 2014 – February 2016		
S. aquifolium S. ilicifolium	Teluk Awur, Jepara	Tropical (Indonesia)	April 2014		
S. polycystum	Panjang Island, Jepara	Tropical (Indonesia)	October 2013		

#### 2.2.2 Preparation of the samples, drying and grinding

Thallus of French *Sargassum*, *S. muticum*, were cut by scissor, approximately 5 cm from the holdfast. Algal material was quickly cleaned with seawater and was kept in closed box to keep them away from direct contact with sunlight. Further, algal material was washed with tap water from the remaining sand and epiphytes and was grinded with a Jupiter's Universal grinder (Wernau, Germany) using a 3-mm perforated grill. Grinded algal material was lyophilized, and the dry algal materials were stored in the dark for further analysis.

Indonesian *Sargassum*, *i.e. S. aquifolium*, *S. ilicifolium* and *S. polycystum*, was collected by cutting the thallus as well, approximately 5 cm from the holdfast. Algal material was quickly cleaned with seawater and was kept in closed box to keep them away from direct contact with sunlight. Further, algal material was washed with tap water from the remaining sand and epiphytes. Indonesian *Sargassum*, was naturally dried for five to seven days. To maintain the quality of the target compound, it is crucial to prevent any direct contact with sunlight. Thus, algal material was dried under the shade. After the drying, they were cut to small pieces with scissor and were grinded by using multi-use home blender.

#### 2.2.3 Determination of the dried algal material

Fresh algal material of 3 g was weighed and placed in glass tube and dried in 100 °C until constant weight of the sample was attained (Gupta *et al.*, 2011). The dry algal material was determined by following formula:

$$DW (\%) = \frac{W_f}{W_i} \times 100$$

D = dry weight of algae express in %

 $W_f$  = final weight of algae (g)

 $W_i$  = initial weight of algae (g)

#### 2.3 Biochemical composition of algal material

#### 2.3.1 Acid hydrolysis

Ten mg of dry material were introduced to 5 mL of HCl 1 M in different vials. All prepared samples were then put in the heating block of 100 °C for 2 h with agitation every 30 min to optimize the process. Samples were left in the box of ice for 5 min. In order to complete the samples concentration to 1 mg mL<sup>-1</sup>, 5 mL NaOH 1 M were added to each different vials. All tubes were kept in 4 °C before further analysis and the designated analysis were protein, sugar, sulfates groups, uronic acid, nitrogen, ash, and polyphenol content analysis.

#### 2.3.2 Water hydrolysis

Ten mg of dry material were introduced to 5 mL of H<sub>2</sub>O in different vials. All prepared samples were then put in the heating block of 100 °C for 2 h with agitation every 30 min to optimize the process. Samples were left in the box of ice for 5 min. In order to complete the samples concentration to 1 mg mL<sup>-1</sup>, 5 mL H<sub>2</sub>O were added to each different vial. All tubes were kept in 4 °C before further analysis and the designated analysis were protein, sugar, sulfates groups, uronic acid, nitrogen, ash, and polyphenol content analysis.

#### 2.3.3 Protein content analysis

Dosage of total protein is using the BSA (Bovine Serum Albumin) method based on the reaction of Biuret reagent which results in the reduction of Cu<sup>2+</sup> ion to Cu<sup>+</sup> by the protein in the alkaline medium. Twenty-five μL of standards (Bovine Serum Albumin) or samples was introduced into the micro plate well then added with 200 μL of reactive A + B (Pierce BCA protein assay kit, Thermo Scientific, 23227). Bovine Serum Albumin was prepared in different concentration ranged from 0-100 μg mL<sup>-1</sup>. The micro plate was then incubated for 1 h in 37 °C. Optical density was then measured at 562 nm. All samples were made in triplicate. Content of protein was calculated using the following formula:

$$\% = \left[ \frac{(OD - b)}{a} \right] \times D \times \frac{1}{1000} \times 10$$

With OD is the optical density of the designated analysis b is the intercept of the calibration curve, a is the slope of the calibration curve, D is 1/factor of dilution, 1/1000 is the conversion of mg from µg, 10 is from 1 mg samples and conversion to % (Smith *et al.*, 1985; Kessler and Fanestil, 1986; Wiechelman *et al.*, 1988; Brown and Jeffrey, 1992).

#### 2.3.4 Sugar content analysis

The principle of this analysis is based on the formation of degradation products of sugars by the action of a strong acid, derivatives of furfural. These chromogenic products are then condensed with Polyphenol to obtain a chromophore. Anhydrous glucose served as standard for the calibration curve prepared in different concentration ranged from 0-100 µg mL<sup>-1</sup>.

One mL of samples or standard was put in the tubes then added with 50  $\mu$ L of phenol 75% and followed by agitation by using the vortex. 2.5 mL of sulfuric acid was added and put them in the cold water of 20 °C for 10 min. After 10 min, all samples were agitated and placed in the water bath of 30 °C for 10 min. Optical density is read against the blank at 490 nm. All samples were made in triplicate (Dubois *et al.*, 1956).

$$\% = \left[ \frac{(OD - b)}{a} \right] \times D \times \frac{1}{1000} \times 10$$

With OD is the optical density of the designated analysis b is the intercept of the calibration curve, a is the slope of the calibration curve, D is 1/factor of dilution, 1/1000 is the conversion of mg from µg, 10 is from 1 mg samples and conversion to % (Smith *et al.*, 1985; Kessler and Fanestil, 1986; Wiechelman *et al.*, 1988; Brown and Jeffrey, 1992).

## 2.3.5 Sulfates groups content analysis

This assay is based on the colorimetric method by Jacques *et al.* (1986) (Guillaume, 2010) which reacts specifically with sulfate groups linked to the polysaccharide backbone. This reaction is based on the formation of a complex between sulphate groups and the 3- amino-7-(dimethylamino) phenothizin-5-ium (Azure A, Sigma-Aldrich A6270) to give a purple. A solution of 17% dextran sulfate in concentration ranging from 0-100 µg mL<sup>-1</sup> served as standard. 200 µl of samples or standard was added with 2 mL of Azure A (10 mg L<sup>-1</sup>) and the optical density against the blank was read at 535 nm. All samples were made in triplicate.

$$\% = \left[ (OD - b)/a \right] \times D \times \frac{1}{1000} \times 10$$

With OD is the optical density of the designated analysis b is the intercept of the calibration curve, a is the slope of the calibration curve, D is 1/factor of dilution, 1/1000 is the conversion of mg from µg, 10 is from 1 mg samples and conversion to % (Smith *et al.*, 1985; Kessler and Fanestil, 1986; Wiechelman *et al.*, 1988; Brown and Jeffrey, 1992).

#### 2.3.6 Uronic acid content analysis

Uronic acid could react to the presence of meta-hydroxydiphenyl (MHDP) and this reaction creates a pink coloration. To enhance the coloration, tetraborate is added and the addition of sulfamic acid minimizes the interferences of neutral sugars detection in the presence of sulfuric acid (Blumenkrantz and Asboe-

Hansen, 1973).

Diluted samples solution or standard of 400  $\mu$ L was added with 40  $\mu$ L of sulfamic acid 4 M. Prior to the introduction of 2.4 mL of sodium tetraborate 75 mM, the solution was agitated and re-agitated afterward. All samples were incubated for 20 min at 80 °C then put in the ice for 5 min. Then, 80  $\mu$ L of MHDP 0.15% was then added and the optical density was read at 525 nm. Glucuronic acid with concentrations ranged from 0-400  $\mu$ g mL<sup>-1</sup> served as standards for the calibration curve (Guillaume, 2010).

$$\% = \left[ \frac{(OD - b)}{a} \right] \times D \times \frac{1}{1000} \times 10$$

With OD is the optical density of the designated analysis b is the intercept of the calibration curve, a is the slope of the calibration curve, D is 1/factor of dilution, 1/1000 is the conversion of mg from µg, 10 is from 1 mg samples and conversion to % (Smith *et al.*, 1985; Kessler and Fanestil, 1986; Wiechelman *et al.*, 1988; Brown and Jeffrey, 1992).

#### 2.3.7 Nitrogen content analysis

By using the Kjeldahl method, there are three main steps to follow. First is the digestion, i.e. the decomposition of nitrogen in organic samples by utilizing a concentrated acid solution. This is accomplished by boiling the samples in concentrated acid in order to obtain ammonium sulfate solution as a final product. Second step is distillation – adding excess base to the acid digestion mixture to convert NH<sub>4</sub><sup>+</sup> to NH<sub>3</sub> followed by boiling and condensation of the NH<sub>3</sub> gas in a receiving solution. Last step is the titration, i.e. to quantify the amount of ammonia in the receiving solution (Bradstreet, 1965).

One hundred mg of dry material and 2.5 g of catalyst were introduced in to the glass tube. The catalyst is composed of 100 g potassium sulfate, 20 g of copper sulfate pentahydrated and 2 g of selenium. Later, 15 mL of sulfuric acid (96–97%) was added along with several glass beads. All the glass tubes specialized for the nitrogen analysis were put in the mineralizer support at 350 °C

for 30 min. BSA (Bovine Serum Albumin) was used as control. All samples and controls were analyzed in triplicate.

After 30 min of mineralizing process, the samples were then distilled by introducing 50 mL of boric acid, concentration 40 g L<sup>-1</sup>, in to the distiller. The distiller was set up before running the analysis with following conditions: 60 mL of concentrated sodium hydroxide (30%) in 5 min (time of analysis). 0.5 mol L<sup>-1</sup> of sulfuric acid was prepared for the titration. The acid was introduced slowly while in to the samples that have been distilled (the distillate) added with a few drops of Taschiro indicator while mixing them on the magnetic stirrer. During the addition of acid sulfuric onto the distillate, the changing color of distillate was observed and the volume of acid needed to cause the color to change was noted. As the molarity of acid sulfuric is changeable depends on the samples.

#### 2.3.8 Ash content analysis

Ash is the general term used to describe the inorganic matter in a fuel. In biomass fuels, the ash content may originate from the biomass itself, e.g. materials that the plant absorbed from the water during its growth. In food analysis, ash content represents the total mineral content. Ashing is the first step in preparing a food sample for specific elemental analysis. Since certain foods are high in particular minerals, ash becomes important (Marshall, 2010).

The principle is to incinerate materials at high temperature (525 °C or higher). 100 mg of dry material was put in the oven of 100 °C for 30 min then burnt on the Bunsen flame for 10 min. Later, the samples were kept in the incinerator of 585 °C for 3 h. Prior to measure their final weight, all samples were put in the desiccators for 1 h.

$$\% = \frac{M_f}{M_i} \times 100$$

 $M_f$  = final mass of algae (g)

 $M_i$  = initial mass of algae (g)

#### 2.3.9 Polyphenol content analysis

Folin-Ciocalteu is a commonly used method for measuring the polyphenol content. This assay is based on the reduction-oxidation (redox) reactions, which are usually considered to be relatively stoichiometric and on the redox potential of the polyphenol hydroxyl group (Hagerman and Butler, 1989).

Solution of 20  $\mu$ l samples (0.25 mg mL<sup>-1</sup>) or standard (Phloroglucinol) was added with 130  $\mu$ l H<sub>2</sub>O was then introduced to the samples or standard solution followed by 40  $\mu$ l of Folin-Ciocalteu. In the end, 40  $\mu$ l mL<sup>-1</sup> of Na<sub>2</sub>CO<sub>3</sub> 200 g L<sup>-1</sup> was added. The reaction was then incubated in 70 °C for 10 min and put in the box of ice to stop the reaction for 5 min. Then the optical density was read against the blank prepared at 620 nm. Phloroglucinol was used as a standard for the calibration curve prepared in different concentrations ranged from 0-100  $\mu$ g mL<sup>-1</sup> (Le Lann *et al.*, 2008).

$$\% = \left[ (OD - b)/_a \right] \times D \times \frac{1}{1000} \times 10$$

With OD is the optical density of the designated analysis b is the intercept of the calibration curve, a is the slope of the calibration curve, D is 1/factor of dilution, 1/1000 is the conversion of mg from µg, 10 is from 1 mg samples and conversion to % (Smith *et al.*, 1985; Kessler and Fanestil, 1986; Wiechelman *et al.*, 1988; Brown and Jeffrey, 1992).

#### 2.4 Extraction of polyphenol compounds

#### 2.4.1 Solid-liquid extraction (SLE)

There were four different solvents chosen for the polyphenol extraction, there were water, methanol 50% (v/v), pure methanol and ethanol 75% (v/v). Later, 40 °C and 3 h were chosen as temperature and time of extraction, it was based on the study performed by Tanniou (2014).

Twenty-five g of dry algal material was diluted in 300 mL of solvents. The filtered samples were then evaporated until dryness, then, the dried extract

obtained was then added by 20 mL of  $H_2O$  in order to get the crude extract. Finally, the raw extracts were frozen and lyophilized for further analysis, *i.e.* polyphenol content, bioactivity analysis and chromatographic analysis.

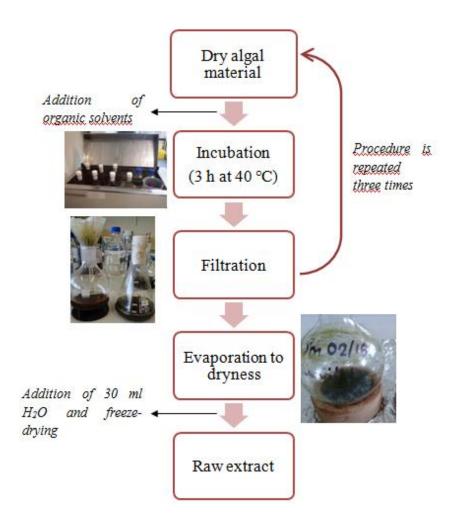


Fig 2-4 Solid-liquid extraction of *Sargassum* polyphenol.

#### 2.4.2 Enzyme-assisted extraction (EAE)

In this study, 19.5 g of dry algal material was diluted in 300 mL of water, then 2.5 % (of dry algal material) of commercial enzyme (two protease and six carbohydrase) was added. Sample solutions were incubated for 3 h in 50 °C. In advance of filtration, all samples were denatured for 20 min in 90 °C. The soluble samples were then frozen and lyophilized for further analysis. As for the insoluble

samples (residues), they were stored in –20 °C. The eight enzymes used were Neutrase, Ultraflo, Alcalase, AMG, Shearzyme, Termamyl, Viscozyme and Celluclast. Method was based on Heo *et al.* (2003) and Hardouin *et al.* (2014). Table 2-3 shows details of the commercial enzymes used in the EAE screening. Aqueous extraction served as control, *i.e.* extraction without enzyme. Figure 2-4 shows the screening of *Sargassum* enzyme-assisted extraction of polyphenol compounds.

Table 2-3 Commercial enzymes applied for the EAE and their optimal conditions based on Kadam *et al.* (2013) and Wijesinghe and Jeon (2012).

Enzymes	Type of enzymes	pH (optimal)	T °C (optimal)
Neutrase	Protease	5.5-7.5	50
Ultraflo	Carbohydrase	7.0	60
Alcalase	Protease	6.5-8.5	50
AMG	Carbohydrase	4.5	60
Shearzyme	Carbohydrase	6.8-7.2	50
Termamyl	Carbohydrase (endoamylase)	6.0	60
Viscozyme	Carbohydrase	3.3-5.5	50
Celluclast	Carbohydrase	4.5	50

For optimizing the conditions of phlorotannin extraction, several different parameters were chosen, i.e. temperature and concentration of enzyme applied. Based on the first screening of EAE by using eight commercials enzymes, enzyme that extracted the highest polyphenol was chosen for optimization. In addition, another enzyme was introduced in this part of work, which is the Protamex as a protease. Protamex is known as enzyme that works optimal at 40 °C, which is also favorable for polyphenol extraction (Park *et al.*, 2005).

The extraction procedure was almost the same as previously described (Heo *et al.*, 2003; Hardouin *et al.*, 2013). The difference lied in the temperature and the concentration of enzyme. Instead of 50°C, 40°C was applied. Then, as for the enzymes concentration, 5 and 10% were determined in order to comprehend whether or not the increase of enzyme concentration would also increase the polyphenol extraction. In order to enhance the recovery of polyphenol extraction, denaturation of enzyme by heating in 90 °C for 20 min was eliminated. Therefore, after the filtration, soluble enzymatic extracts were directly stored in the freezer of

-20 °C in order to stop the reaction. 40 °C has been proven to be the most optimal temperature for the polyphenol compound extraction. An elevated temperature above 40 °C might degrade the compound. All steps of extraction were conducted in the dark in order to prevent the oxidation of the compound.

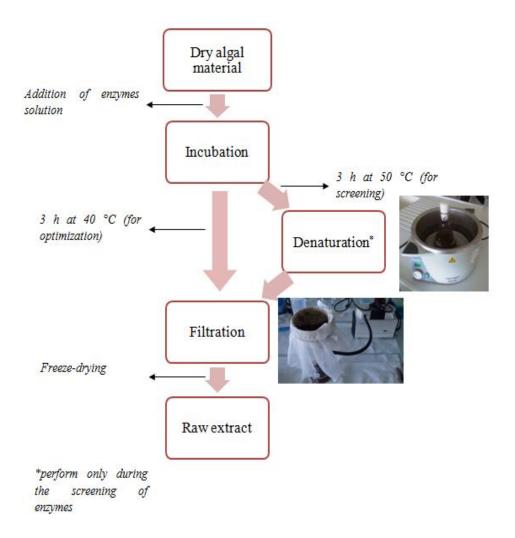


Fig 2-5 Enzyme-assisted extraction of *Sargassum* sp. polyphenol (screening of enzymes).

## 2.4.2.1 Solid-phase extraction of Sargassum muticum enzymatic extract

Solid-phase extraction (SPE) is extraction method using a solid and a liquid phase to isolate and to concentrate a target compound, also called as analyte, from a solution. Main principle of this method is to introduce a sample into compatible sorbent, then to allow it to be absorbed by the sorbent and then to elute it with suitable solvents.

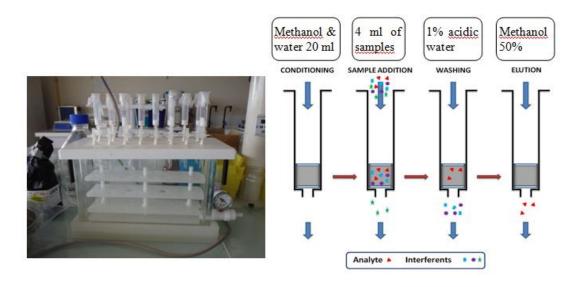


Fig 2-6 Solid-phase extraction of *S. muticum* polyphenol.

SPE was performed by using reversed phase C-18 cartridges based on method performed by Tanniou (2013), with slight modification (Fig 2-6). Prior to the fractionation process, the cartridges were conditioned with the following elution; 20 mL methanol and 20 mL ultrapure H<sub>2</sub>O successively. Later, 4 mL of raw extracts (5 mg/mL) was loaded into the cartridge HyperSep<sup>TM</sup> C18 from Thermo Scientific<sup>TM</sup> (40-60 µm, 1 g, 60 Å, 6 mL) and left to absorb the column (time). Prior to elution, the cartridge was washed by 1% acidic water to remove sugars, organic acids, and other water-soluble constituents (Dai and Mumper, 2010). Phlorotannins were then eluted with 40 mL methanol 50% (v/v) (flow rate ~3 mL min<sup>-1</sup> at positive pressure). All processes were repeated four times in order to accumulate sufficient amount of fractions. Same fractions were pooled and evaporated under reduced pressure at 40 °C by a rotary evaporator. Each fraction was redissolved using acetonitrile-methanol (2:1; v/v). Polyphenol content of methanol fraction was determined with the Folin-Ciocalteu method and its antioxidant activity was tested via DPPH and FRAP.

#### 2.5 Characterization of enzymatic extracts

## 2.5.1 Fourier Transform Infra Red (FT-IR) spectroscopy

FT-IR is a vibrational spectroscopic technique and its principle is based on the facts that most molecules absorb light in the infra-red wavelength region of the electromagnetic spectrum. This absorption corresponds specifically to the chemical bonds present in molecules. FT-IR can facilitate to confirm the identity of a pure compound by using the unique collection of absorption bands.

Spectra of *S. muticum* enzymatic extracts was determined using a Nicolet TM iSTM 5 FT-IR spectrometer (Thermo ScientificTM) with diamond crystals plates. The spectra were collected with subsequent scanning of samples in triplicate ranging from 4000-400 cm<sup>-1</sup>. Two mg of dry enzymatic extracts were introduced onto the crystal plate. The spectra were then acquired as the percentage of absorbance. The detected peaks were then compared with a standard of phloroglucinol.

#### 2.5.2 High Performance Liquid Chromatography (HPLC) analysis

Chromatography is a technique to separate mixtures of substances into their components on the basis of their molecular structure and composition. This involves a stationary phase (a solid, or a liquid supported on a solid) and a mobile phase (a liquid or a gas). The mobile phase flows through the stationary phase and carries the components of the mixture with it. Reversed phase HPLC (RP-HPLC) used in this study has a non-polar stationary phase and an aqueous, moderately polar mobile phase. One common stationary phase is silica, which has been surface-modified with RMe<sub>2</sub>SiCl, where R is a straight chain alkyl group such as C<sub>18</sub>H<sub>37</sub> or C<sub>8</sub>H<sub>17</sub>. With such stationary phases, retention time is longer for molecules which are less polar, while polar molecules elute more readily (early in the analysis).

The analytical chromatographic analysis was performed by UltiMate 3000 Rapid Separation LC (RSLC) System consisted of pump, auto-samplers and diode array detector (DAD-3000RS) with UV-vis detection (Thermo Scientific<sup>TM</sup>). The

column was a Nucleodur® RP  $C_{18}$  (250 x 4.6 mm i.d; 5 µm) of Macherey-Nagey. The reversed-phase HPLC analysis was performed based on the modified method of Koivikko *et al.* (2007) and Sanchez-Camargo *et al.* (2015). *S. muticum* extracts were eluted by a multi-step gradient of solvents A (1% acid formic in water) and B (acetonitrile). The elution was as following; 0–20 min: 5% B, 20–35 min: 20 % B, 35–45 min: 80% B, and 45–55 min: 0% B. Polyphenol compounds were detected at 280 nm, while the wavelength was set ranging from 240–300 nm. The injection volume was 100 µL with 1 mL min<sup>-1</sup> flow-rate at 150 bar. Internal standard range of phloroglucinol was prepared with concentration ranging from  $100-800 \ \mu \text{g mL}^{-1}$ .

#### 2.6 Antioxidant activity

#### 2.6.1 DPPH radical scavenging assay

2,2-diphenyl-1-picrylhydrazyl or DPPH is a stable free radical. DPPH assay is based on the theory that a hydrogen donor is an antioxidant. It measures compounds that are radical scavengers. DPPH shows a strong absorption maximum at 517 nm.

A series of butylated hydroxyl-anisole (BHA), butylated hydroxyl-toluene (BHT) (respectively 2(3)-*t*-Butyl-4-hydroxyanisole, and 2,6-Di-*tert*-butyl-4-methylphenol, Sigma, Saint Quentin Fallavier, France) solution – serves as positive standard – is prepared in different concentrations (2-50 μg mL<sup>-1</sup>). DPPH solution of 0.25 mM is introduced to 100 μL samples solution. Samples solution of *S. muticum* hydrolyzates was made in different concentration by diluting the solution stock in methanol (0–10%). Before reading the optical density at 517 nm, all samples are incubated in 40 °C for 30 min. Then, the percentage of inhibition is calculated by suing following formula:

$$I(\%) = \left[ \frac{\left(A_{control} - A_{sample}\right)}{A_{control}} \times 100 \right]$$

With  $A_{control}$  = absorbance of control and  $A_{sample}$  = absorbance of samples

 $IC_{50}$  of samples is determined based on the regression obtained from dose response curve. IC is defined as the concentration sufficient to obtain 50 % of a maximum scavenging capacity.

### 2.6.2 Ferric Reducing Antioxidant Power (FRAP)

This method is based on the reduction of a ferroin analog, the complex of tripyridyltriazine  $Fe(TPTZ)^{3+}$ , to the intensely blue colored complex  $Fe(TPTZ)^{2+}$  by the presence of antioxidants in acidic medium. Then, the results are obtained from the absorbance at 593 nm and expressed as  $\mu m Fe^{2+}$  equivalents (Benzi and Strain, 1996; Benzie and Strain, 1999).

The reagent of TPTZ consisted of 300 mM of acetate buffer with pH 3.6, 10 mM TPTZ (2, 4, 6-tripyridyl-s-triazine) and 20 mM FeCl<sub>3</sub> <sub>6</sub>H<sub>2</sub>O solution. Fifty µL of *S. muticum* enzymatic extracts was introduced in microplate mixed with 150 µL reagent. After 15 min of incubation, the absorbance was read at 593 nm. The results were expressed in µM equivalent to FeSO<sub>4.7</sub>H<sub>2</sub>O calculated from the calibration curve (Shahwar *et al.*, 2012).

#### 2.7 Tyrosinase inhibition activity

Tyrosinase or polyphenol oxidase is an oxidoreductase that participates in the biosynthesis of melanin, a ubiquitous biological pigment found in human's skin, hair, etc. Presence of L-DOPA (L-3,4-Dihydroxyphenylalanine) as substrate results the formation of dopaquinone leading to melanin production. Under distressed condition, melanine production could cause hypo or hyper pigmentation on human's skin. Potential tyrosinase inhibitors are examined in the presence of L-tyrosine or L-DOPA as enzyme substrate, and tyrosinase activity is assayed in terms of dopachrome formation (Chang, 2012).

Solution of 80  $\mu$ L phosphate buffer is introduced to 20  $\mu$ L of sample solution added with 20  $\mu$ L of DMSO (dimethyl sulfoxide) 50%. Later, 40  $\mu$ L of L-DOPA is added to the solution followed with 40  $\mu$ L tyrosinase to complete the reaction. Before reading the absorbance at 475 nm, all microplates were incubated

at 37 °C for 30 min. Kojic acid and Quercetin serving as control were prepared at different range of concentration (0.02-0.4%). Percentage of inhibition is calculated by following formula:

$$I(\%)_{at \times min} = \left[ \frac{\left( A_{control} - A_{sample} \right)}{A_{control}} \right] \times 100$$

Where:

$$\begin{split} A_{control} &= \left(DO_{control\,(x\,min)} - DO_{control\,(x\,min)}\right) \\ &- \left(DO_{control\,(0\,min)} - DO_{control\,(0\,min)}\right) \\ A_{sample} &= \left(DO_{sample\,(x\,min)} - DO_{sample\,(x\,min)}\right) \end{split}$$

$$- \left(DO_{sample\ (0\ min)} - \ DO_{sample\ (0\ min)}\right)$$

 $A_{control}$  = absorbance (optical density/DO) of control

A<sub>sample</sub> = absorbance (optical density/DO) of samples

## 2.8 Elastase inhibition activity

As inhibitor, trypsin inhibitor type II-O was used prepared in phosphate buffer in final concentration 1 g L<sup>-1</sup>. All tested samples were dissolved in DMSO 10% with final concentration same as standard. The activity of porcine pancreatic elastase (Type I; Sigma Chem. Co) was examined using substrate (N-methoxysuccinyl-ala-ala-pro-val-p-nitroanilide/N-Suc-(Ala)<sub>3</sub>-nitroanilide), and the release of ρ-nitroaniline at 405 nm was measured.

The reaction of 20  $\mu$ L samples was carried out in 80  $\mu$ L of 100 mM Tris and 500 mM NaCL pH 8, 50  $\mu$ L of 0.4 U/mL elastase solution, and 50  $\mu$ L of 0.64 mM substrate. The elastase inhibition was assessed every 10 min at 25°C during 40 min of reading. Same reactions applied for standard, but the volume of standard introduced in the well ranged between 0-100  $\mu$ L ( $c_f = 0$ -0.5 mg mL<sup>-1</sup>) in order to obtain the calibration curve. All reactions were made in triplicate (Sallenave *et al.*, 1998). The percent inhibition of elastase was calculated as follows:

$$I(\%) = \left[ \frac{(A-B)}{A} \right] \times 100$$

Where A = absorbency at 405 nm without the samples/standards, and B = the change in absorbency with samples/standards.

#### 2.9 Antimicrobial activity

#### 2.9.1 Antibacterial activity

Agar diffusion method (Bauer *et al.*, 1996) was chosen as method for evaluating the antibacterial activity of *S. muticum* hydrolyzates. Two gram positive bacteria – *Staphylococcus hominis* and *Bacillus subtilis* – as well as two gram negative bacteria – *Pseudomonas aeruginosa* and *Escherichia coli* – were chosen for this test. These bacteria were prepared in liquid nutrient broth media with bacterial density 1.5 x 10<sup>8</sup> cfu mL<sup>-1</sup>. Bacterial suspension was poured on to solidified agar media and incubated for 1 h in 37 °C.

Sargassum sp. enzymatic extracts of final concentration 5 mg mL<sup>-1</sup> were prepared in sterile physiological water. Twenty μL of tested samples solution – enzymatic extracts solution or positive control (Phosphomycin, Ampicil and Streptomycin) – was introduced to sterile paper disc. All samples were made in triplicate. Later, the impregnated sterile paper discs were placed on the agar media in accordance with samples coding made in advance. The petri dishes were then incubated at 37 °C for 48 h. Zone of inhibition established on the agar media was then measured and expressed in cm. Antibacterial activity was represented by following formula:

**Antibacterial activity** = Diameter of inhibition zone (cm) -0.6 cm (diameter of paper disc)

#### 2.9.2 Biofilm Inhibition Activity

This test is based on the Crystal Violet Assay (Adetunji and Isola, 2011). *E. coli, P. aeruginosa* and *B. subtilis* were selected as biofilm forming bacteria. In this assay, the dye that is bounded to adherent cells is resolubilized and measured in

optical density (OD). Prior to analysis, bacterial solution of 1.5 x 10<sup>8</sup> cells in physiologic sterile water 10% was made by incubating the bacteria culture in Lisogeny Broth (LB) sterile medium at 37 °C for 24 h. Later, the bacteria culture was centrifuged for 5 min at 5000 rpm and the supernatant was discarded. The remaining pellet was dissolved in physiologic sterile water 10%.

Aliquot of 200  $\mu$ L bacterial solution was then introduced into a sterile 96-well microplate and left for 24 h at 37 °C in order to allow the bacteria to form a film layer in the microplate. After 24 h of incubation, the plate was washed two times with the physiologic sterile water 10%. Then, 200  $\mu$ L of *S. muticum* enzymatic extracts (1 mg mL<sup>-1</sup>) dissolved in LB were added into the microplate and incubated for 24 h in 37 °C. In the next 24 h, the microplate was rinsed two times with physiologic sterile water 10% and introduced with 200  $\mu$ L of 10% crystal violet – prepared in ultrapure water. Fifteen minutes after the crystal violet was added, subsequently, the microplate was rinsed and the biofilm formed in the microplate was then suspended in 200  $\mu$ L of ethanol. The microplate was later read in 600 nm. Negative and positive control consisted of physiologic sterile water 10% and bacteria were applied. Both the control and the samples were made in triplicates.

#### 2.9.3 Antiviral and cytotoxicity activity

Antiviral activity of *S. muticum* enzymatic extracts and their cytotoxicity were analyzed. For cytotoxicity, the kidney cells of African green monkey (Vero cells) were used. Herpes-simplex virus type-1 (HSV-1) model was used to determine the antiviral effect.

Aliquots of 100  $\mu$ L cell suspension (3.5×10<sup>5</sup> cells mL<sup>-1</sup>) in Eagle's MEM (Minimum Essential Medium) containing 8% FCS (Fetal Calf Serum) was incubated with 50  $\mu$ L of diluted enzymatic extract (concentration from 10 to 500  $\mu$ g/mL) in 96-well plates (72 h, 37 °C, 5% CO<sub>2</sub>). Three replicates were infected using 50  $\mu$ L of medium and a virus suspension at a Multiplicity of Infection (MOI) of 0.001 ID<sub>50</sub> cells<sup>-1</sup>. After incubation, antiviral activity was evaluated by the neutral red dye method. Zovirax, a commercial antiherpetic acyclovir (9-2(2-

hydroxyethoxymethyl) guanine), was used as reference inhibitor (positive control).

For antiviral, 50% effective antiviral concentration (EC $_{50}$ ) was expressed as the concentration that achieved 50% protection of virus-infected cells from virus-induced destruction. The optical density (OD) was related directly to the percentage of viable cells, which was inversely related to the cytopathic effect (CPE). The linear regression was determined for each assay on the basis of cell controls (0% CPE) and virus controls (100% CPE). Data were expressed as a percentage of protection (%P):

$$\left| \frac{((OD_t)_{virus} - (OD_c)_{virus})}{((OD_c)_{MOCK} - (OD_c)_{virus})} \right| \times 100$$

where  $(OD_t)_{virus}$  is the OD of the test sample,  $(OD_c)_{virus}$  is the OD of the virus control and  $(OD_c)_{MOCK}$  is the OD of the mock-infected control (McLaren *et al.*, 1983; Langlois *et al.*, 1986).

For cytotoxicity, 50% cytotoxic concentration ( $CC_{50}$ ) was defined as the concentration that reduced the OD of treated cells to 50% of that of untreated cells. Data were expressed as a percentage of protection (%D):

$$\left| \frac{((OD_c)_C - (OD_c)_{MOCK})}{(OD_c)_C} \right| \times 100$$

where  $(OD_c)_C$  and  $(OD_c)_{MOCK}$  are the OD values of untreated cells and treated cells, respectively (Langlois *et al.*, 1986).

## 2.10 Statistical Analysis

Data obtained in this study was based on the average value from triplicate and the

data was presented as mean value  $\pm$  standard of deviation, with n = 3. Analysis of variance was performed to analyze the significant difference caused by the independent variables (p<0.05) by using IBM SPSS Statistics 23. If there was any significant difference observed than Tukey's post-hoc test was used to evaluate which variables showed a significant difference to the other. A significant difference from single independent variable was assigned by different superscript letters. As for analysis with multiple independent variables, the data was presented together with its post-hoc test for better description of which independent variables contributed to the significant difference.

Spectrophotometer data obtained in this study were presented as the mean value with its standard deviation (mean  $\pm$  standard deviation), n = 3. Later, for the seasonal variation data of *S. muticum*, they were statistically analyzed using analysis of variance in order to determine whether there was a significant difference according to the seasons of collection (p<0.05). Tukey's post-hoc test was conducted to evaluate the significant difference. Pearson's correlation was performed to view the relationship between the environmental condition and the biochemical composition of *S. muticum*. Principal Coordinate Analysis was applied to view the tendency of *S. muticum* biochemical composition content in different seasons.

For the spectroscopy data of *S. muticum* from FT-IR analysis, the Principal Component Analysis (PCA) was chosen to group the biochemical composition based on seasons. This grouping was facilitated by selecting certain wavenumbers that corresponded to the analyzed *S. muticum* biochemical composition. Later, for the chromatogram data of *S. muticum* from HPLC analysis, the Principal Component Analysis (PCA) was chosen to group the polyphenol content based on seasons. This grouping was facilitated by selecting certain retention time of *S. muticum* polyphenol that corresponded to phloroglucinols.

## **Chapter 3 Results and Discussions**

#### 3.1 Biochemical composition of *Sargassum*

#### 3.1.1 Indonesian Sargassum sp.

Table 3-1 presents the biochemical composition of *S. aquifolium*, *S. ilicifolium*, and *S. polycystum*. *S. aquifolium* and *S. ilicifolium* were species collected on April 2014 at Telukawur, Jepara and *S. polycystum* was collected on October 2013 at Panjang Island, Jepara.

The biochemical compositions of *Sargassum* analyzed in this study were protein, sulfates groups, neutral sugar, uronic acid, nitrogen, polyphenol and ash. Prior to analysis, all dry materials were hydrolyzed using hydrochloric acid (HCl) and ultrapure water (H<sub>2</sub>O Milli-Q), except for the nitrogen and ash content that used directly the dry material.

Uronic acid, protein, sugar and polyphenol contents of *S. ilicifolium* were the highest with  $7.8 \pm 3.3\%$ ,  $30.5 \pm 1.2\%$ ,  $10.8 \pm 0.2\%$ , and  $2.1 \pm 0.4\%$  of dry algal material, respectively. In *S. aquifolium*, there were only  $4.5 \pm 2.0\%$ ,  $20.9 \pm 1.1\%$ ,  $8.4 \pm 0.1\%$ ,  $10.7 \pm 0.6\%$ , and  $1.7 \pm 0.3\%$  of uronic acid, protein, sugar, sulfates groups, and polyphenol contents, respectively. *S. polycystum* contained the highest sulfates groups ( $13.7 \pm 0.7\%$ ) However, the nitrogen content of these two species was the same, with 1.6% of dry algal material (Table 3-1).

Ash content of *S. polycystum*  $(33.6 \pm 1.9\%)$  was higher compared to *S. aquifolium*  $(32.5 \pm 0.3\%)$  and *S. ilicifolium*  $(29.0 \pm 2.6\%)$ . Different case showed with the protein and nitrogen content of *S. polycystum*, apparently, it had the lowest content with  $19.8 \pm 0.6$  and  $1.0 \pm 0.2\%$  of dry algal material, respectively. Nevertheless, *S. polycystum* contained more uronic acid, sugar, and sulfates groups contents than the *S. aquifolium*. In the other hand, they both had 1.7% of polyphenol.

Table 3-1 Biochemical composition of Indonesian Sargassum species.

Algon	Content (% dry algal material)								
Algae	Uronic acid	Protein	Sugar	Sulfates groups	Polyphenol	Ash	Nitrogen	Total	Other
S. aquifolium	$4.5 \pm 2.0$	$20.9 \pm 1.1$	$8.4 \pm 0.1$	$10.7 \pm 0.6$	$1.7 \pm 0.3$	$32.5 \pm 0.3$	$1.6 \pm 0.1$	69.2	30.8
S. ilicifolium	$7.8 \pm 3.3$	$30.5 \pm 1.2$	$10.8 \pm 0.2$	$11.4 \pm 0.5$	$2.1 \pm 0.4$	$29.0 \pm 2.6$	$1.6 \pm 0.1$	77.1	22.9
S. polycystum	$6.1 \pm 2.6$	$19.6 \pm 0.6$	$10.3 \pm 0.1$	$13.7 \pm 0.7$	$1.7 \pm 0.3$	$33.6 \pm 1.9$	$1.0 \pm 0.2$	72.0	28.0

Table 3-2 Average of *S. muticum* biochemical composition based on year of collection.

	Content (% dry algal material)								
Year	Uronic Acid	Protein	Sugar	Sulfates groups	Polyphenol	Ash	Nitrogen	Total	Other
2014	$2.0 \pm 0.4$	$24.5 \pm 6.4$	$13.5 \pm 2.6$	$12.2 \pm 1.7$	$1.9 \pm 0.7$	$25.1 \pm 4.1$	$2.0 \pm 0.7$	81.1	18.9
2015	$2.6 \pm 1.0$	$27.2 \pm 7.1$	$12.2 \pm 4.3$	$8.2 \pm 1.0$	$3.1 \pm 1.1$	$23.9 \pm 2.8$	$1.9 \pm 0.6$	79.1	20.9

#### Conclusion

Indonesian *Sargassum* contained quiet high of ash or total mineral. We acknowledge that there was intra-species variation regarding the biochemical composition, unless for the nitrogen content showing almost the same value.

#### 3.1.2 French S. *muticum* – seasonal variation

Seasonal variation of French *Sargassum* was studied, *i.e. S. muticum*. In order to acknowledge the seasonal pattern of their biochemical composition, the analysis was conducted for two complete years, *i.e.* from February 2014 to February 2016 (Table 3-2 and 3-3).

In 2014, the highest intensity of sunlight occurred in June with 270.7 h/month<sup>1</sup> with 46.2 mm/month of precipitation. The water surface temperature recorded in this month was 18 °C and the maximal air temperature ( $T_{max}$ ) was 25 °C<sup>1</sup>. Meanwhile in 2015, the seawater surface temperature (SST) in June was 16 °C<sup>1</sup>, 2° lower than in 2014 with sunlight intensity 20 h more (290.7 h month<sup>-1</sup> (1)). During this period, the precipitation rate was the lowest with only 4.2 mm month<sup>-1</sup> (1).

In 2014, the average content of *S. muticum* uronic acid, protein, sugar, sulfates groups, ash, and nitrogen contents were 2.0%, 24.5%, 13.5%, 12.2%, 1.9%, 25.0% and 2.0%, respectively. Meanwhile, the average content of *S. muticum* uronic acid, protein, sugar, sulfates groups, ash, and nitrogen in 2015 were 2.6%, 27.2%, 12.2%, 8.2%, 3.1%, 23.9% and 1.9%, respectively. It seems like that there was a decrease in sugar, sulfate groups, ash and nitrogen content of *S. muticum* in 2015. Results of *S. muticum* biochemical composition are presented on Table 3-2.

Moisture and dry material of *S. muticum* showed a seasonal variation. *S. muticum* moisture content ranged from 74.4-84.4% of total weight, while its dry

<sup>(1)</sup> The meteorological data was based on the real time observation provided the <u>www.meteociel.fr</u>

material was at 12.9-25.6% of total weight. In 2014, the minimum moisture content was from June as the maximum was on September. Then, in 2015, the minimum moisture content of *S. muticum* was obtained on March and the maximum was on September. Appendix 2 shows the figure of *S. muticum* moisture content and dry material from two years of sampling.

Fig 3-1 illustrates the dry material of *S. muticum* in different period of collection. As can be seen from the picture, the dry material from samples collected from February to July tended to increase followed by a decline from August to December in 2014. Unfortunately, this tendency was not followed in the next year of sampling.

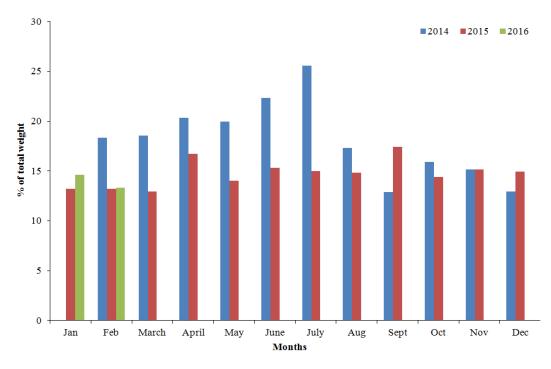


Fig 3-1 Dry material of *S. muticum* in different period of collection (February 2014 – February 2016).

# 3.1.2.1 Analysis of variance of seasonal effects towards S. muticum biochemical composition

Analysis of variance of seasonal effects towards S. muticum biochemical composition was analyzed and it revealed that there was a significant difference of biochemical composition caused by season (p<0.05). Dry material of S. muticum

was significantly influenced by the seasons. As presented in Table 3-3, the significant difference occurred between the spring-summer and the autumn-winter (p<0.05).

Table 3-3 Seasonal variation of *S. muticum* dry material.

Seasons	Dry material
	(% of total weight)
Spring 2014-2015	$17.1 \pm 3.0^{a}$
Summer 2014 and 2015	$18.4 \pm 4.5^{a}$
Autumn 2014 and 2015	$15.1 \pm 1.6^{b}$
Winter 2014, 2015, and 2016	$14.3 \pm 1.9^{b}$

The protein content of *S. muticum* in summer was significantly lower than spring, autumn and winter (p<0.05). In winter and spring, the sugar content was also significantly lower from summer and autumn (p<0.05). Polyphenol content of *S. muticum* in autumn appeared to be significantly higher than spring and summer (p<0.05). In the contrary with its ash content, *S. muticum* contained significantly higher ash content in autumn and winter than in spring and summer (p<0.05). As for its nitrogen content, a significant difference was remarked between summer and spring (p<0.05). Apparently, there was no significant seasonal effect towards *S. muticum* uronic acid and sulfates groups content (p>0.05). Table 3-4 presents the average value of *S. muticum* biochemical composition in each season of collection from February 2014 – February 2016.

Based on the Pearson's correlation, the biochemical composition of *S. muticum* appeared to be related to some of the collected meteorological data. An inversely weak correlation yet significant was shown by the protein content of *S. muticum* towards  $T_{max}$  (r = -,257; p < 0.05) and  $T_{min}$  (r = -,250; p < 0.05). Same case with ash content which had an inverse yet moderate correlation with  $T_{max}$  (r = -,454; p < 0.01), an inversely significant weak correlation with  $T_{min}$  (r = -,377; p < 0.01) and sea surface temperature (r = -,331; p < 0.01). Sunlight gave a significantly inverse but strong correlation with *S. muticum* ash content (r = -,509; p < 0.01). Meanwhile, the correlation of ash content with precipitation was a positive weak correlation (r = ,237; p < 0.05).

Regarding the sugar content of *S. muticum*, the effect of SST and precipitation were positively and significantly moderate (r = ,271; p < 0.05; r = ,350; p < 0.01). The sugar content also showed a significantly positive yet weak correlation with the  $T_{max}$  (r = ,267; p < 0.05) and  $T_{min}$  (r = ,336; p < 0.01). Sunlight had a very weak correlation with sugar content (r = ,099; p > 0.05 & 0.01). Sulfates groups significantly and moderately correlated to sunlight,  $T_{max}$  and SST with r = ,378; p < 0.01; r = ,298; p < 0.01 and r = ,298; p < 0.01, respectively. In addition, the sulfates groups of *S. muticum* showed an insignificantly and inversely very weak correlation with precipitation (r = -,31; p > 0.05 & 0.01).

Uronic acid, polyphenol and nitrogen content of *S. muticum*, based on the result obtained in this study, were insignificantly affected with sunlight,  $T_{max}$ ,  $T_{min}$ , precipitation and SST (p>0.05 & 0.01). Furthermore, they mostly had a very weak correlation with r=.00-.19. Further detail concerning the Pearson's correlation of *S. muticum* biochemical composition and meteorological data is presented on Appendix 5.

Component loadings and biplot showed the distribution tendency of *S. muticum* biochemical composition in different seasons as presented on Fig 3-2. Interestingly, the component loading also described the correlation between the biochemical compositions and the meteorological as previously described. In the winter, *S. muticum* mostly contained ash and uronic acid in the spring. During winter and spring, *S. muticum* was dominated by the protein content. In the mean time, the polyphenol content was quite abundant in summer and autumn season. As for the sulfates groups, the axe was covered with spring and summer. It indicated that this compound was mostly found during these seasons.

#### Conclusion

Season, indeed, contributes to the biochemical composition of *S. muticum*, especially in protein, sugar, polyphenol, nitrogen and ash content. Protein and nitrogen content demonstrate same pattern regarding their changing value. Seasonal effects towards *S. muticum* biochemical composition are usually due to

the fluctuating environmental conditions, for example the temperature, salinity, precipitation, and duration of sunlight. Protein and ash content showed an inverse correlation with temperature, however, the ash content of *S. muticum* has a positive correlation with sunlight. Sugar and sulfate groups content in this study had positive correlation with precipitation and sunlight, respectively. In addition, both of these compounds demonstrated a positive correlation with air temperature.

Table 3-4 Seasonal variation of *S. muticum* biochemical composition during February 2014-2016 (content expressed as % of dry algal material). Superscript letters indicate significant difference (p<0.05).

Season	Uronic Acid	Protein	Sugar	Sulfates	Polyphenol	Ash	Nitrogen	Total	Other
				groups					
Spring 2014-	$2.4 \pm 1.3^{a}$	$27.1 \pm 3.4^{a}$	$10.9 \pm 3.2^{a}$	$10.2 \pm 2.0^{a}$	$2.2 \pm 0.4^{a}$	$23.5 \pm 2.8^{a}$	$2.3 \pm 0.5^{a}$	78.6	21.4
2015		1	ī				1		
Summer 2014	$2.3 \pm 0.8^{a}$	$21.8 \pm 6.4^{b}$	$14.8 \pm 2.4^{b}$	$10.7 \pm 3.4^{a}$	$2.4 \pm 1.0^{a}$	$22.4 \pm 3.4^{a}$	$1.7 \pm 0.7^{\rm b}$	76.1	23.9
and 2015									
Autumn 2014	$2.2 \pm 0.4^{a}$	$27.2 \pm 10.4^{a}$	$14.3 \pm 3.4^{b}$	$9.6 \pm 2.4^{a}$	$3.2 \pm 1.7^{b}$	$25.4 \pm 3.9^{b}$	$2.0 \pm 0.6^{a}$	83.9	16.1
and 2015									
Winter 2014,	$2.2 \pm 0.5^{a}$	$27.4 \pm 2.6^{a}$	$11.6 \pm 3.7^{a}$	$9.3 \pm 2.0^{a}$	$2.5 \pm 0.6^{a}$	$25.8 \pm 2.3^{b}$	$2.0 \pm 0.4^{a}$	80.8	19.2
2015, and									
2016									

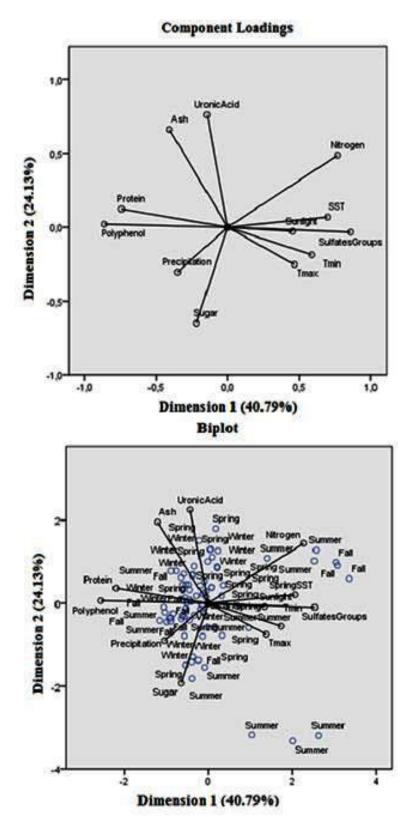


Fig 3-2 Component loadings and biplot of Principal Coordinate Analysis of *S. muticum* biochemical composition

#### 3.2 Extraction of *Sargassum* polyphenol

#### 3.2.1 Seasonal variation of S. muticum extracted with different solvents

Extraction yield of *S. muticum* ranged from 8.0% to 24.1% of dry algal material as presented in Table 3-5. Results indicated that there was a seasonal effect towards the extraction yield of *S. muticum* (p<0.05). From four solvents applied for the extraction of *S. muticum*, water resulted higher yield than the other solvents. Aqueous methanol placed in second after water for the extraction yield of *S. muticum*, followed by the aqueous ethanol. Methanol yielded the lowest extraction yield.

Table 3-5 Extraction yield of *S. muticum* extracted with different solvents (expressed in % of dry algal material). Superscript letters indicate significant difference at p<0.05.

Season	Water	Methanol	Aqueous methanol 50%	Aqueous ethanol 75%
Spring	$24.0 \pm 5.0^{a}$	$11.6 \pm 2.6^{a}$	$14.8 \pm 3.5^{a}$	$8.7 \pm 3.6^{a}$
Summer	$21.5 \pm 5.8^{ab}$	$9.8 \pm 2.4^{a}$	$10.8 \pm 3.9^{b}$	$12.3 \pm 3.1^{a}$
Autumn	$17.7 \pm 0.8^{b}$	$11.3 \pm 1.2^{a}$	$13.9 \pm 2.8^{a}$	$14.9 \pm 2.2^{b}$
Winter	$24.1 \pm 4.5^{a}$	$10.0 \pm 1.9^{a}$	$8.6 \pm 2.4^{b}$	$8.0 \pm 1.4^{a}$

## 3.2.1.1 Polyphenol content of *S. muticum* extracted with different solvents based on season

Seasonal variation of *S. muticum* polyphenol extracted in different solvents, *i.e.* water, methanol, aqueous methanol 50% and aqueous ethanol 75% was evaluated. Fig 3-3, 3-4, 3-5 and 3-6 present the seasonal variation of polyphenol content from water, methanol, aqueous methanol 50% and aqueous ethanol 75% extracts of *S. muticum*. Different letters that is assigned in each bars represented a significant difference between variables (p<0.05).

Results from analysis of variance exhibited that seasonal collection of S. muticum gave significant effect to its polyphenol content (p<0.05). The results from Tukey's post-hoc test informed that in water extract of S. muticum, its polyphenol showed significant difference in its content between winter-spring and autumn-summer (p<0.05). Then, the content of S. muticum polyphenol, extracted

with aqueous methanol 50% and collected in summer, was significantly higher than that of from spring and winter (p<0.05). There was no significant difference in polyphenol content of S. muticum from winter and spring-autumn for the same extract (p>0.05). In aqueous ethanol 75% extracts, the significant content of polyphenol was shown by S. muticum collected in summer with those from spring and winter (p<0.05). It seems that there was no significant variation of polyphenol content in the methanolic extracts of S. muticum (p>0.05).

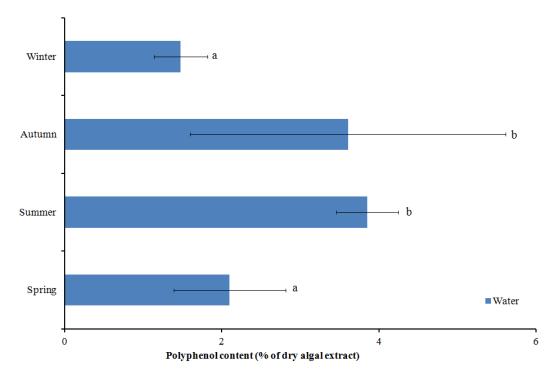


Fig 3-3 Polyphenol content of *S. muticum* water extract. Different alphabets on the bar indicate that there was significant difference (p>0.05).

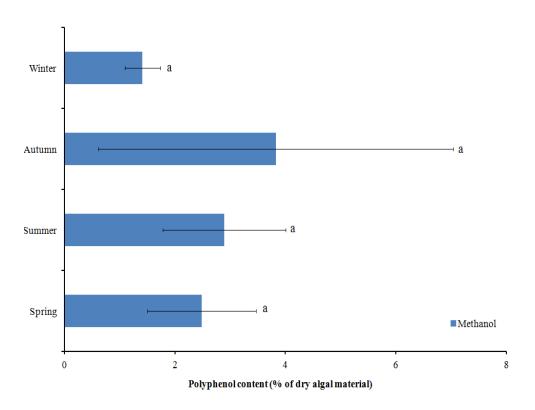


Fig 3-4 Polyphenol content of *S. muticum* methanol extract. Different alphabets on the bar indicate that there was significant difference (p>0.05).

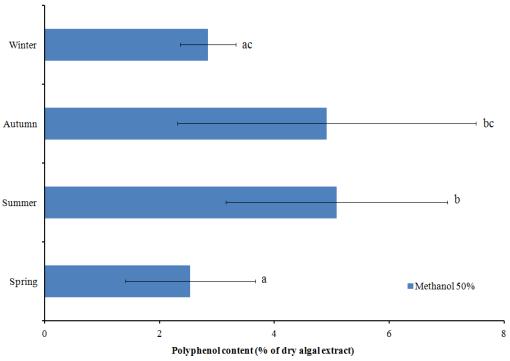


Fig 3-5 Polyphenol content of *S. muticum* aqueous methanol 50% extract. Different alphabets on the bar indicate that there was significant difference (p>0.05).

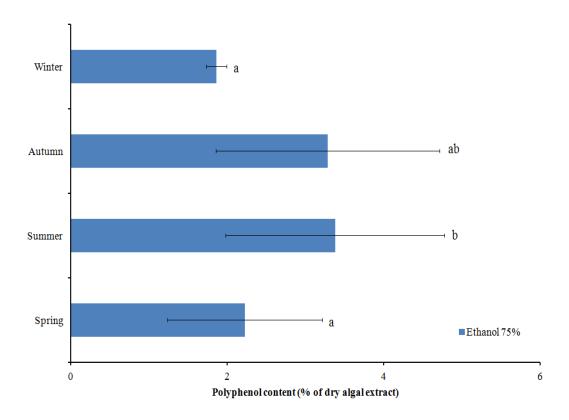


Fig 3-6 Polyphenol content of *S. muticum* aqueous ethanol 75% extract. Different alphabets on the bar indicate that there was significant difference (p>0.05).

Pearson correlation analysis from these data was performed. It helped to inform the correlation between polyphenol content of *S. muticum* extracted with different solvents and the meteorological data. It seems that in all extracts – water, methanol, aqueous methanol 50% and aqueous ethanol 75%, they were inversely correlated with precipitation. However, the correlation was weak since the value of r ranging at -.336 to -.110. Moreover, a significant correlation was only revealed with the water extract at r = -.336, p < 0.01 while the other had no significant correlation (p>0.05 and 0.01, 2-tailed) In the contrary, these extracts had significantly and positively weak to strong correlation with the sunlight, the maximum and the minimum air temperature ( $T_{max}$  and  $T_{min}$ ) and the sea surface temperature (SST).

Polyphenol content in water extracts showed a significantly strong correlation with sunlight (r = .700, p<0.01),  $T_{max}$  (r = .777, p<0.01),  $T_{min}$  (r = .757,

p<0.01) and SST (r=.763, p<0.01). Methanol extracts of *S. muticum* had significantly weak correlation sunlight (r=.370, p<0.01),  $T_{max}$  (r=.324, p<0.05),  $T_{min}$  (r=.318, p<0.05) and SST (r=.272, p<0.05). Meanwhile, significantly moderate correlation was exhibited between the aqueous methanol 50% extracts with sunlight (r=.451, p<0.01),  $T_{max}$  (r=.492, p<0.01),  $T_{min}$  (r=.516, p<0.01) and SST (r=.469, p<0.01). As for aqueous ethanol 75% extracts, a significantly weak correlation was presented with sunlight (r=.288, p<0.05). Further, the correlation of this extract with  $T_{max}$ ,  $T_{min}$  and SST was significantly moderate with r=.432, p<0.01, r=.483, p<0.01 and r=.409, p<0.01, respectively. Details to the statistical results for this study are presented on Appendix 7.

## 3.2.1.2 Polyphenol content of S. muticum in different solvents

Aside from the seasonal effect, effect of different solvents in *S. muticum* polyphenol was evaluated as well, as presented in Fig 3-7, 3-8, 3-9, and 3-10. Solvents gave significant effect towards polyphenol content of *S. muticum* collected in summer and winter. In summer, the aqueous methanol 50% extract of *S. muticum* polyphenol was significantly high than water, methanol, and aqueous ethanol 75% (p<0.05). Later, aqueous methanol 50% of *S. muticum* in winter showed a significant difference of polyphenols towards the methanol and water extracts (p<0.05). Aqueous methanol 50% extract of *S. muticum* had the highest polyphenol content in all seasons ranging from 2.5±1.1 to 4.9±2.6% of dry algal extract.

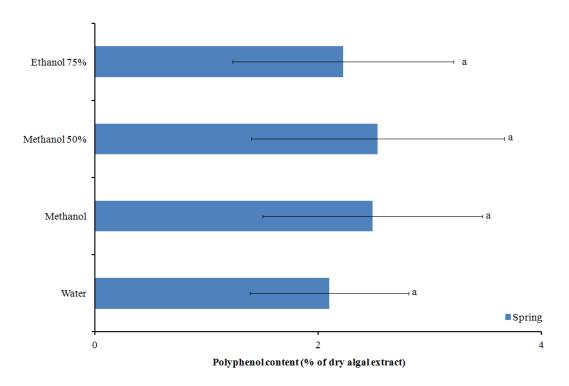


Fig 3-7 Polyphenol content of *S. muticum* collected during spring in different solvents. Alphabets on the bar indicate that there was no significant difference (p>0.05).

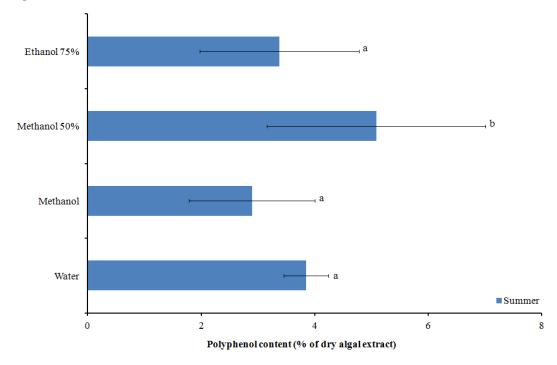


Fig 3-8 Polyphenol content of *S. muticum* collected during summer in different solvents. Different alphabets on the bar indicate that there was significant difference (p<0.05).

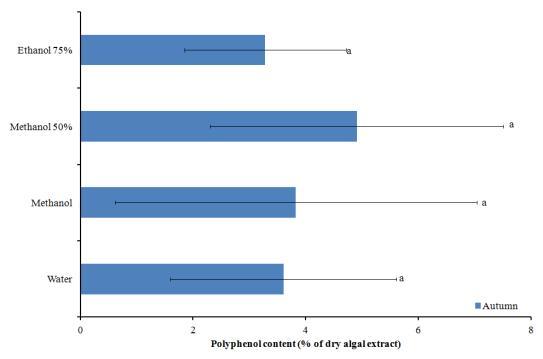


Fig 3-9 Polyphenol content of *S. muticum* collected during autumn in different solvents. Different alphabets on the bar indicate that there was significant difference (p<0.05).

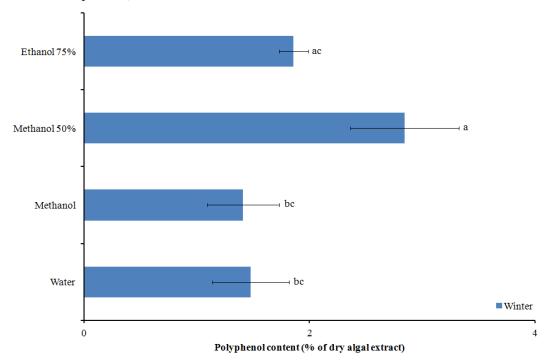


Fig 3-10 Polyphenol content of *S. muticum* collected during winter in different solvents. Different alphabets on the bar indicate that there was significant difference (p<0.05).

#### Conclusion

From four different solvents used, water had the highest extraction yield with 24% of dry algal material. Seasons and solvents gave significant effects in *S. muticum* polyphenol content. In seasonal variation point of view, *S. muticum* collected in summer contained higher polyphenol when it was extracted with water, aqueous methanol 50% and aqueous ethanol 75%. As for methanol extract, it contained more polyphenol content in autumn sample than the other season. If we look the polyphenol content based on the variation of solvent then the polyphenol from all seasons was more favorable with aqueous methanol 50%.

### 3.2.2 Solid-liquid extraction of Sargassum polyphenol

In solid-liquid extraction of *Sargassum* polyphenol, there were two main treatment of statistical analysis performed. The first analysis concerned about the effect of different solvents in extracting polyphenol in tropical *Sargassum* from Indonesia, *i.e. S. aquifolium*, *S. ilicifolium* and *S. polycystum*. Additionally, effect of different species was also statistically evaluated. Then, the second comparison analysis was performed merely on *S. muticum*. These two different analyses were performed because of the difference in dry material due to different drying treatment. *S. aquifolium*, *S. ilicifolium* and *S. polycystum* were naturally dried, Meanwhile, *S. muticum* was lyophilized. As a result, the characteristics of final products obtained were not the same. The first three species of *Sargassum* were in medium pieces of thalli and *S. muticum* dry form was powder-like material.

Table 3-6 and 3-7 presents the extraction yield and polyphenol content of Indonesian Sargassum in different solvents. Water extraction resulted better yielded than the extraction with methanol, aqueous methanol 50% and aqueous ethanol 75% and S. polycystum had the highest yield of extracts with  $22.8 \pm 1.1$ % of dry algal material. Methanol extraction showed the lowest yield ranging from  $3.1 \pm 0.7$  to  $8.3 \pm 3.2$ % of dry algal material. Polyphenol of S. aquifolium, S. ilicifolium and S. polycystum was best extracted with methanol (6.4  $\pm$  1.5% of dry algal material), water (7.0  $\pm$  1.3% of dry algal material) and aqueous methanol

50% (8.1  $\pm$  2.2% of dry algal material), respectively. In order to evaluate whether there was a significant effect shown by different solvents or not towards polyphenol content of Indonesian *Sargassum*, statistical analysis of variance was performed to analyze whether there was a significant difference or not.

Table 3-6 Extraction yield of Indonesian *Sargassum* with the resume of significant difference caused by species and type of solvents.

Algae	Solvents	Yield (% of dry algal material)	Sig. ( <i>p</i> <0.05) of Species	Sig. ( <i>p</i> <0.05) of Solvents
	Water	$17.3 \pm 2.2$		
	MeOH	$5.7 \pm 1.8$		
S. aquifolium	MeOH 50%	$15.8 \pm 1.5$		
	EtOH 75%	$12.3 \pm 0.4$		
	Water	$14.7 \pm 0.8$		
	MeOH	$3.1 \pm 0.7$	.005	.000
S. ilicifolium	MeOH 50%	$7.3 \pm 1.0$	$(\mathbf{df} = 2 ; \mathbf{F} =$	$(\mathbf{df} = 3; \mathbf{F} =$
	EtOH75	$7.5 \pm 1.3$	3.399)	18.024)
S. polycystum	Water	$22.8 \pm 1.1$		
	MeOH	$8.3 \pm 3.2$		
	MeOH 50%	$13.4 \pm 1.9$		
	EtOH 75%	$12.5 \pm 0.5$		

Statistical analysis showed that there was a significant effect caused by different species and solvents used (p<0.05). This significant effect was related to the extraction yield of Indonesian *Sargassum*. As presented on Appendix 10, extraction yield of *S. aquifolium* and *S. polycystum* was significantly different from *S. ilicifolium* (p<0.05). In addition, water extracts of Indonesian *Sargassum* had a significant difference to methanolic (methanol and aqueous methanol 50%) and ethanolic (aqueous ethanol 75%) extracts (p<0.05). Then, methanol extracts with aqueous methanol 50% and aqueous ethanol 75% extracts showed a significant difference in their extraction yield with methanol (p<0.05).

Table 3-7 Polyphenol content of Indonesian *Sargassum* with the resume of significant difference caused by species and type of solvents.

Algae	Solvents	Polyphenol (% of dry algal extract)	Sig. ( <i>p</i> <0.05) of Species	Sig. ( <i>p</i> <0.05) of Solvents
	Water	$1.8 \pm 0.1$		
	MeOH	$6.4 \pm 1.5$		
S. aquifolium	MeOH 50%	$3.6 \pm 0.2$		
	EtOH 75%	$6.0 \pm 0.4$		
	Water	$5.3 \pm 0.3$		
	MeOH	$6.0 \pm 0.4$	.000	.000
S. ilicifolium	MeOH 50%	$1.4 \pm 0.0$	$(\mathbf{df} = 2 ; \mathbf{F} =$	$(\mathbf{df} = 3; \mathbf{F} =$
	EtOH75%	$4.8 \pm 0.3$	50.569)	98.976)
	Water	$7.0 \pm 1.3$		
G 1	MeOH	$3.9 \pm 0.1$		
S. polycystum	MeOH 50%	$8.1 \pm 2.2$		
	EtOH 75%	$3.5 \pm 0.0$		

In polyphenol content, there was significant difference caused by different species of Sargassum (p<0.05), which was shown by the three species of Sargassum. In addition, a significant difference was also observed in different type of solvents that were used to extract polyphenols from Indonesian Sargassum. Table 3-7 presents the polyphenol content of Indonesian Sargassum using different solvents. As can bee seen, type of species and solvents gave significant difference towards polyphenol content. Appendix 8 shows the significant difference in Sargassum polyphenol content due to different solvents (p<0.05). Aqueous methanol 50% and aqueous ethanol 75% had a significant difference with water and methanol (p<0.05). It can also be said that in this study water extraction yielded better of dry algal material while aqueous methanol 50% and aqueous ethanol 75% extracted higher content of polyphenol in Indonesian Sargassum.

Table 3-8 Extraction yield and polyphenol content of French S. muticum June 2014

Solvents	Yield	Polyphenol
	(% of dry algal material)	(% of dry algal extract)
Water	$24.8 \pm 0.3^{a}$	$3.8 \pm 0.1^{a}$
MeOH	$13.3 \pm 2.9^{c}$	$2.8 \pm 0.4^{\rm b}$
MeOH 50%	$15.7 \pm 1.5^{\rm b}$	$6.6 \pm 0.2^{a}$
EtOH 75%	$19.0 \pm 1.5^{\rm b}$	$4.0 \pm 0.6^{\rm b}$

In *S. muticum*, aqueous methanol 50% was more effective in extracting polyphenol containing  $6.6 \pm 0.2\%$  of dry algal extract as shown in Table 3-8. Nevertheless, water yielded higher dry algal material (24.8  $\pm$  0.3% of dry algal material) than other solvents. Statistical analysis, presented in Appendix 9, revealed that there was a significant difference in extraction yield and polyphenol content of *S. muticum* caused by different solvents (p<0.05). Extraction yield of *S. muticum* water extract was significantly different from methanol, aqueous methanol 50% and aqueous ethanol 75% extracts (p<0.05). A significant difference was also shown by methanol and aqueous ethanol 75% extracts (p<0.05). However, there was no significant difference between aqueous methanol 50% and aqueous ethanol 75% in their extraction yields (p<0.05). Type of solvents also gave a significant effect in *S. muticum* polyphenol (p<0.05). Water and aqueous methanol 50% extracts was significantly different from methanol and aqueous ethanol 75% extracts was significantly different from methanol and aqueous ethanol 75% extracts (p<0.05).

#### Conclusion

Solid-liquid extraction has been widely used to extract bioactive compound in marine algae, particularly polyphenol. Efficacy of this technique relies in the appropriate selection of extraction conditions, such as type of solvents, temperature, particle sizes of samples, etc. By applying different type of solvents, our study demonstrated that aqueous organic solvents extracted more polyphenol than water only. On the opposite, water is the best media to obtain high extraction yield in comparison to organic solvents since it co-extracted the polysaccharides and salts as well. Regardless, intra-species variation of polyphenol also needs to

be considered. As presented in this study, each *Sargassum* species showed different preference of solvents. *S. aquifolium* and *S. ilicifolium* contained high polyphenol in methanol extract. Meanwhile, 50% aqueous methanol extract of *S. muticum* and *S. polycystum* showed more polyphenol content than the other extracts; water, methanol, 75% aqueous ethanol extracts.

## 3.2.3 Enzyme-assisted extraction of Sargassum polyphenol

# 3.2.3.1 Screening of enzymes

In enzyme-assisted extraction, the insoluble sludge was separated from the soluble aqueous phase. Only the soluble aqueous phase was studied. Yields and polyphenol content were quantified and compared. Table 3-9 presents the extraction yield and polyphenol content of *S. muticum* enzymatic extracts.

There was no significant difference in the yield of enzyme-assisted extraction (p>0.05). However, compared to the control ( $26.5 \pm 4.7\%$  of dry algal material) – an extraction without added enzyme – the enzymatic extracts produced higher yield of dry material; except for Shearzyme extract ( $26.2 \pm 1.7\%$  of dry algal material). As a protease enzyme, Neutrase extract yielded  $32.6 \pm 4.9\%$  of dry algal material while other protease, Alcalase, only yielded  $28.7 \pm 3.5\%$  of dry material. Second highest yield was obtained from Ultraflo extract (carbohydrase),  $31.6 \pm 4.5\%$  of dry algal material. The yield of enzyme-assisted extraction from other carbohydrase, such as AMG, Termamyl, Viscozyme and Celluclast ranged between  $27.3 \pm 0.2$  and  $30.2 \pm 5.2\%$  of dry algal material.

Insoluble and soluble polyphenol content of *S. muticum* enzymatic extract were determined and results are expressed as % of dry hydrolysate. For the insoluble polyphenol, all *S. muticum* enzymatic extracts showed the same content of polyphenol, *i.e.* 0.9% of dry hydrolysate. As for the control, there was  $1.0 \pm 0.0\%$  of dry hydrolysate. Despite the low value, this result indicated that there were still some traces of polyphenol in the residues of *S. muticum* enzymatic extracts.

Neutrase extract had the highest yield of extraction with  $32.6 \pm 4.6\%$  of dry hydrolysate; however, the highest polyphenol was shown by Viscozyme extract,  $6.4 \pm 0.5\%$  of dry algal material followed by Termamyl  $5.0 \pm 0.1\%$  of dry hydrolysate. The protease i.e. Neutrase and Alcalase only extracted polyphenol at  $3.8 \pm 0.0\%$  of dry hydrolysate. Nevertheless, in comparison with the control ( $2.6 \pm 0.0\%$  of dry hydrolysate); all eight enzymes enhanced the extraction of polyphenol compound in *S. muticum*.

Table 3-9 Yield of extract and polyphenol content of S. muticum enzymatic extracts (different letter superscript letters indicates a significant difference (p< 0.05)).

Samples	Yield of extracts	Polyphenol
	(% of dry algal material)	(% of dry hydrolysate)
Control	$26.5 \pm 4.8^{a}$	$2.6 \pm 0.0^{a}$
Neutrase	$32.6 \pm 4.9^{a}$	$3.8 \pm 0.0^{ac}$
Ultraflo	$31.6 \pm 4.5^{a}$	$4.2 \pm 0.0^{ac}$
Alcalase	$28.7 \pm 3.5^{a}$	$3.8 \pm 0.0^{ac}$
Celluclast	$29.9 \pm 0.8^{a}$	$4.5 \pm 0.0^{ m bc}$
AMG	$30.2 \pm 5.2^{a}$	$4.8 \pm 0.0^{\rm bc}$
Shearzyme	$26.2 \pm 1.7^{a}$	$4.7 \pm 0.0^{\rm bc}$
Viscozyme	$28.1 \pm 3.0^{a}$	$6.4 \pm 0.5^{ m bc}$
Termamyl	$27.3 \pm 0.3^{a}$	$5.0 \pm 0.1^{bc}$

Beside polyphenol, other biochemical compositions of *S. muticum* enzymatic extracts such as uronic acid, protein, sugar and sulfate groups were also determined. Table 3-10 presents the biochemical composition of *S. muticum* enzymatic extracts. This result informs the presence of polysaccharides in the enzymatic extracts of *S. muticum*.

Table 3-10 Biochemical composition of *S. muticum* enzymatic extracts.

Comples	Content (% of dry hydrolysate)			
Samples -	Uronic acid	Protein	Sugar	Sulfates groups
Control	$0.5 \pm 0.0$	$33.0 \pm 2.3$	$8.2 \pm 2.1$	$7.0 \pm 0.1$
Neutrase	$0.6 \pm 0.1$	$31.8 \pm 0.8$	$7.7 \pm 1.5$	$8.8 \pm 0.0$
Ultraflo	$0.6 \pm 0.0$	$40.3 \pm 0.3$	$6.8 \pm 0.7$	$6.8 \pm 0.1$
Alcalase	$0.7 \pm 0.2$	$47.4 \pm 5.4$	$12.9 \pm 3.6$	$7.4 \pm 0.1$
Celluclast	$0.6 \pm 0.1$	$52.1 \pm 0.6$	$7.4 \pm 0.8$	$7.6 \pm 0.1$
AMG	$0.5 \pm 0.0$	$53.1 \pm 0.0$	$7.9 \pm 0.2$	$7.5 \pm 0.4$
Shearzyme	$0.6 \pm 0.1$	$54.3 \pm 0.7$	$8.5 \pm 0.9$	$7.5 \pm 0.2$
Viscozyme	$0.5 \pm 0.1$	$27.1 \pm 0.7$	$10.6 \pm 1.6$	$7.4 \pm 0.1$
Termamyl	$0.5 \pm 0.1$	$41.3 \pm 0.8$	$9.9 \pm 0.4$	$7.6 \pm 0.2$

### 3.2.3.2 Optimization of S. muticum polyphenol enzyme-assisted extraction

According to the screening, Viscozyme and Termamyl extracted more polyphenol than the other enzymes and control – the aqueous phase, as previously presented. Based on the previous result of this study, these two enzymes were further selected to evaluate the efficiency of these enzymes in extracting polyphenol with adjusted conditions. Such conditions referred to the temperature and concentration of enzymes. As mentioned by many previous studies, 40 °C has been the most favorable temperature to extract polyphenol. Additionally, the concentration of enzymes was augmented to 5 and 10% assuming that it might increase the extracted polyphenol content as well. Result is presented on Table 3-10.

This study was meant to acknowledge whether or not by applying same enzyme in samples collected from different month would give same results. As previously described, Viscozyme managed to extract polyphenol until 6% of dry algal material along with Termamyl. Both of these enzymes are carbohydrase. Then, in this part, we decided to attempt an enzymatic extraction using Protamex – a protease.

Based on the observation of soluble enzymatic extracts, there was a distinguish color produced between *S. muticum* from June and November 2014. In screening, we used *S. muticum* from June 2014 while in this optimization; *S.* 

muticum from November 2014 was selected. Soluble enzymatic extracts of *S. muticum* from June 2014 had dark brown color. As for that in November, they showed dark green color. As can be seen on Table 3-11, extraction yield of *S. muticum* extracted with 10% of Viscozyme and Termamyl slightly increased from 33 to 35% and from 39 to 42%. However, for *S. muticum* that was extracted with 10% of Protamex did not show any increase in polyphenol content. Viscozyme extracted more polyphenol than Protamex and Termamyl in 5% of concentration. In comparison with 10%, 5% of enzymes had more content of polyphenol.

Table 3-11 Extraction yield and total polyphenol content of *S. muticum* from November 2014 extracted with enzymes in different concentration at 40°C without denaturation.

Enzymes	Concentration (%)	Yield (% of dry algal material)	Polyphenol (% of dry hydrolysate)
Viscozyme		$33.8 \pm 0.6$	$3.6 \pm 1.5$
Protamex	5	$42.8 \pm 0.2$	$2.0 \pm 0.7$
Termamyl	3	$39.8 \pm 2.6$	$1.5 \pm 0.0$
Viscozyme		$35.4 \pm 1.2$	$1.8 \pm 0.0$
Protamex	10	$42.6 \pm 0.6$	$1.7 \pm 0.0$
Termamyl	10	$42.1 \pm 0.4$	$2.7\ \pm0.0$
Water		$33.4 \pm 1.1$	$3.4 \pm 0.0$

According to Table 3-11, the next extraction procedure would use Viscozyme and Protamex. As presented in above table, Viscozyme and Protamex extracts of *S. muticum*, using 5% of enzyme, contained higher polyphenol than Termamyl. Table 3-12 presents results of *S. muticum* extraction yield and polyphenol content from June 2014. The extraction was performed at 40 °C for 3 h and without denaturation. The condition of extraction was selected in the assumption that this condition was favorable for polyphenol extraction. Nevertheless, results showed that the selected parameters did not extract more polyphenol in *S. muticum*. Interestingly, the extraction yield of S. muticum Viscozyme extract was higher than the yield from the screening. As a reminder, parameters that were applied in polyphenol extraction of *S. muticum* were 50°C, 3

h and with denaturation at 90 °C. By extracting *S. muticum* with Viscozyme and Protamex at 40°C for 3 h and without denaturation, extraction yield attained 45.5  $\pm$  1.4 and  $46.4 \pm 0.5\%$  of dry hydrolysate.

Table 3-12 Extraction yield and total polyphenol content of *S. muticum* from June 2014 extracted with 5% enzymes at 40°C without denaturation.

Enzymes	Yield (% of dry algal material)	Polyphenol (% of dry hydrolysate)	
Viscozyme	$45.5 \pm 1.4$	$2.1 \pm 0.2$	
Protamex	$46.4 \pm 0.5$	$3.0 \pm 0.0$	

## 3.2.3.3 Enzyme-assisted extraction of Indonesian Sargassum

Same enzymes and condition of extraction was also applied to Indonesian *Sargassum*, *i.e.* Viscozyme and Protamex at  $40^{\circ}$ C. Results are presented in Table 3-13. *S. polycystum* was best extracted with Protamex,  $38.2 \pm 6.8\%$  of dry algal material. On the other hand, *S. aquifolium* and *S. ilicifolium* had higher extraction yield with Viscozyme than Protamex.

Table 3-13 Extraction yield of Indonesian *Sargassum* extracted with enzymes at 40°C without denaturation.

Algae	Enzymes	Yield (% of dry algal material)	Sig. ( <i>p</i> <0.05) of Species	Sig. ( <i>p</i> <0.05) of Enzymes
S. aquifolium	Viscozyme Protamex	$29.5 \pm 5.4$ $26.2 \pm 5.9$		
S. ilicifolium	Viscozyme Protamex	$27.9 \pm 1.5$ $21.4 \pm 2.1$	.001 (df = 2; F = 11.964)	.276 (df = 1; F = 1.300)
S. polycystum	Viscozyme Protamex	$35.8 \pm 2.7$ $38.2 \pm 6.8$		

Table 3-14 Polyphenol content of Indonesian *Sargassum* extracted with enzymes at 40°C without denaturation.

Algae	Enzymes	Polyphenol (% of dry hydrolysate)	Sig. ( <i>p</i> <0.05) of Species	Sig. ( <i>p</i> <0.05) of Enzymes
S. aquifolium	Viscozyme Protamex	$4.7 \pm 0.2$ $4.2 \pm 0.0$		
S. ilicifolium	Viscozyme Protamex	$3.6 \pm 0.1$ $5.0 \pm 0.3$	.000 (df = 2; F = 18.995)	.000 (df = 1; F = 77.395)
S. polycystum	Viscozyme Protamex	$4.0 \pm 0.4$ $6.1 \pm 0.1$		

Table 3-14 shows the polyphenol content of Indonesian Sargassum extracts. Analysis of variance exhibited that there was a significant difference in extraction yield and polyphenol content between different species of Indonesian Sargassum (p<0.05). Extraction yield and polyphenol content of S. polycystum was significantly different from that of S. aquifolium and S. ilicifolium (p<0.05). In addition, Viscozyme and Protamex gave a significant difference towards polyphenol content in Indonesian Sargassum (p<0.05) but not in the extraction yield (p>0.05).

### **Conclusion**

Extraction of polyphenol from *S. muticum* using enzyme-assisted extraction was successfully demonstrated at laboratory-scale. Application of carbohydrase and protease managed to enhance polyphenol content, likewise, extraction yield of *S. muticum* was also improved compared to water extraction. In this study, introduction of Viscozyme increased more polyphenol content yet the extraction yield was improved by Neutrase.

Enzyme-assisted extraction, in certain extent, improves extraction yield and facilitated release of target bioactive compound to the medium. Moreover, this technique is performed under moderate conditions that maintain the bioactivity of target compounds. To do so, it is important to apply the appropriate condition of extraction according to type of enzymes used. Most of the enzymes work optimally at 50 °C, except for Protamex that works optimally at 40 °C.

Interestingly, polyphenol compounds from *S. ilicifolium* and *S. polycystum* extracted with Protamex at 40 °C showed slightly higher polyphenol content than Viscozyme extract. It can be concluded that in optimizing the enzyme-assisted extraction for polyphenol, it is important to consider the optimal condition for each enzymes, inter-species and seasonal variation.

## 3.3 Characterization of Sargassum biochemical composition and extracts

### 3.3.1 Fourier transform infrared spectroscopy (FT-IR)

In evaluating the seasonal variation of *S. muticum* biochemical composition, dry algal materials were analyzed using the FT-IR spectroscopy. Results from this analysis are presented in sub-chapter 3.3.1.1.

Sub-chapter 3.3.1.2 and 3.3.1.3 present the FT-IR analysis from *S. muticum* and Indonesian *Sargassum* enzymatic extracts, respectively. Presence of polysaccharides (sugars, uronic acid and sulfates groups), proteins, and polyphenols were determined by selecting the corresponding wavenumbers.

### 3.3.1.1 Biochemical composition of *S. muticum*-seasonal variation

Fig 3-11 presents an example of *S. muticum* biochemical composition spectra from April 2014. The FT-IR spectra managed to detect interesting peaks at 3270-3250, 2940-2925, 2300-1900, 1660-1600, 1500-1200 and 1100-1000 cm<sup>-1</sup>. Table 3-15 presents details on these peaks. The peaks detection was analyzed based on the single analysis run in each samples.

Certain wavenumbers corresponded to particular biochemical composition could be selected. The interest was to see the seasonal pattern of specific biochemical composition. For that reason, the presence of polysaccharides, protein, uronic acid, sulfates groups and polyphenol were detected by selecting

the corresponded wavenumbers. Later, Principal Component Analysis (PCA) was performed to further describe the seasonal pattern in each compound.

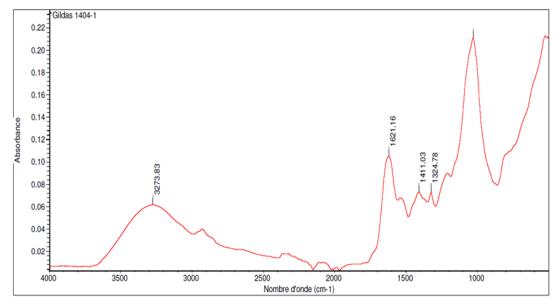


Fig 3-11 FT-IR spectra of *S. muticum* biochemical composition (sample from April 2014).

Wavenumbers chosen for polysaccharides were 1320, 1085-1070, 1030-1025, 895-875, and 825-815 cm<sup>-1</sup>. As for the protein, 3600-3450 and 1660-1440 cm<sup>-1</sup> were selected. 1300-1015 cm<sup>-1</sup> was the range of wavenumbers chosen to detect the presence of sulfates groups. 1705-1595, 1415-1395, 1090-1035, and 940-930 cm<sup>-1</sup> were wavenumbers for determining the uronic acid compound in *S. muticum*. The detection of polyphenol by FT-IR analysis was indicated by the following wavenumbers 3300-3200, 1620-1605, 1460-1445, 1090-1075, 935-920, 880-865, and 675- 660 cm<sup>-1</sup>.

Table 3-15 FT-IR characterization of algal biochemical compositions (Bi *et al.*, 2007; Prati *et al.*, 2010; Gomez-Ordonez and Ruperez, 2011; Marimuthu *et al.*, 2012; Girija *et al.*, 2013; Kannan, 2014; Sugiono *et al.*, 2014; Rodrigues *et al.*, 2015; Kusumaningsih *et al.*, 2016).

Compositions	Wavenumbers (cm <sup>-1</sup> )	Liaisons	
	3423.87-1633	OH, C=O, and COOH; hydroxyl, carbonyl, carboxyl (alginic aicd)	
	1730	C=O; carboxylic acid ester	
	1675 and 1600	COO stretch; carboxyclate anion	
	1320	C=C stretch	
Polysaccharides	1200-970	C-C, C-O pyranoid ring and C-O-C streetching of glycosidic	
	1139.21-1036.60	Acidic polysaccharides	
	1100	Mannuronic units	
	1027 (1030-1025)	Guluronic units	
	817 (825-815)	Mannuronic units	
	1705-1595	C-O stretch	
	1609 and 1420	-COO- asymmetric and symmetric stretch of uronic acid	
TT ' '1	1611	Aromatics nucleus	
Uronic acid	1415-1395	C-OH stretch	
	1090-1035	C-O vibration	
	940-930	C-O-C stretch	
	1280-1210	(S=O) Stretch of sulfate groups	
	1250-1040	S=O stretching	
Sulfates groups	1030	S=O stretching	
	836.75-836.20	Sulfate groups	
	670-600	C-S and C=S stretching	
	3600-3450	N-H stretch	
D	2347	N-H bond	
Proteins	1660-1440	C=O, N-H, $\delta$ CH <sub>2</sub> -CH <sub>3</sub>	
	939	N-O; nitrogen grouping	
Polyphenol	3300-3200	O-H and C-H stretch; phenol hydroxyl carboxylic acid	
	2945-2930	C-H stretch	
	1620-1605	Aromatic nucleus	
	1384	-OH phenolic groups	
	1086	C-H stretch; aromatic nucleus	

Principal component analysis, also known as PCA, is most commonly used to condense the information contained in a large number or original variables into a smaller set of dimensions. This analysis facilitates the explanation of variance-covariance structure of a set of variable. This set of variable is converted into

principal components. Fig 3-12, 3-13, 3-14, 3-15 and 3-16 show PCA scores plot of S. muticum biochemical composition (polysaccharides, protein, uronic acid and sulfates groups) based on FT-IR spectra. Score plot is a projection of data set onto subspace and used for interpreting relations among observations, in this case, a relation between biochemical composition and seasonal variation. In general, the PCA score plot presented on Fig 3-12, 3-13, 3-14, 3-15 and 3-16 distinguished between the spring-summer and autumn-winter samples. In polysaccharides, protein, uronic acid, sulfates groups and polyphenol of S. muticum, the summer and winter samples were grouped in the same region in the PCA score plot. However, the pattern of the grouping from these biochemical compositions was different between one to another. Polysaccharides and sulfates groups showed that the autumn-winter samples had grouped in the left region of the score plot towards the PC-2. Meanwhile, protein, uronic acid and polyphenol content of S. muticum tended to form a group towards the right region of the plot. It could be suggested that the seasonal variation of polysaccharides and sulfates groups were on the same pattern and the other way around with protein, uronic acid and polyphenol.

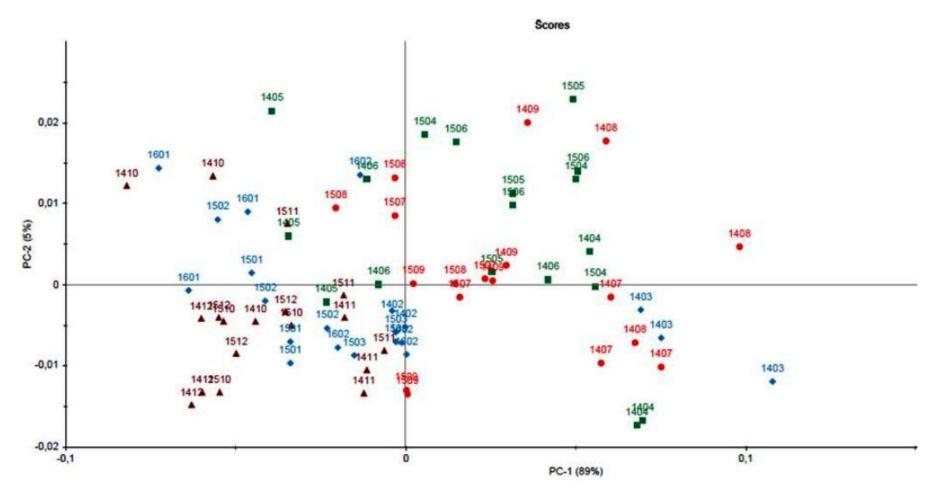


Fig 3-12 Principal component analysis of *S. muticum* polysaccharides from selected wavenumbers of 1320, 1085-1070, 1030-1025, 895-875, and 825-815 cm<sup>-1</sup>. Green box: Spring, Red circle: Summer, Purple triangle: Autumn, and Blue diamond: Winter.

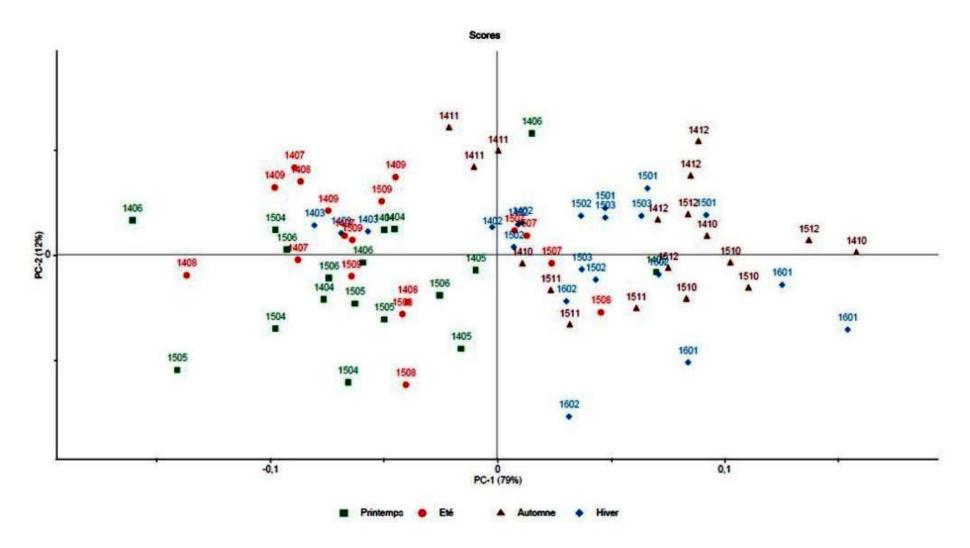


Fig 3-13 Principal component analysis of *S. muticum* proteins from selected wavenumbers of 3600-3450 and 1660-1440 cm-1. Green box: Spring, Red circle: Summer, Purple triangle: Autumn, and Blue diamond: Winter.

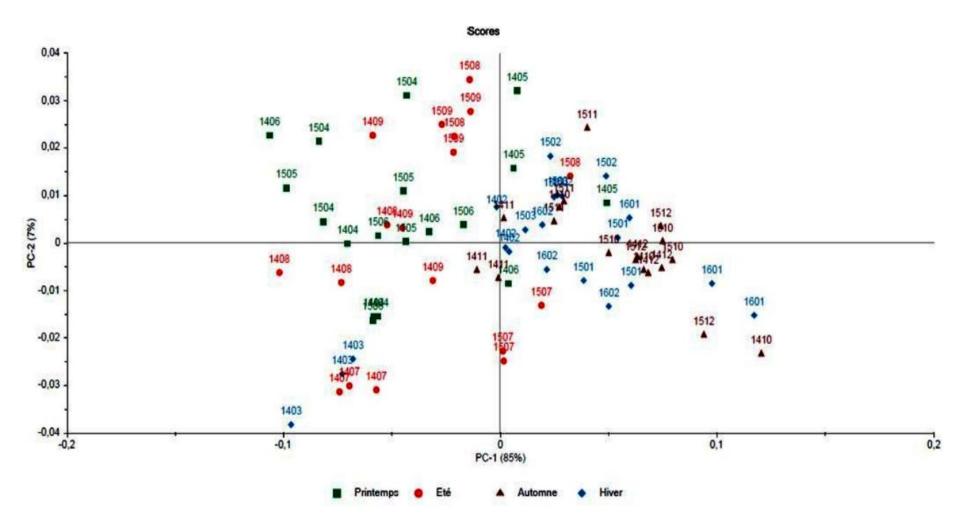


Fig 3-14 Principal component analysis of *S. muticum* uronic acid from selected wavenumbers of 1705-1595, 1415-1395, 1090-1035, and 940-930 cm-1. Green box: Spring, Red circle: Summer, Purple triangle: Autumn, and Blue diamond: Winter.

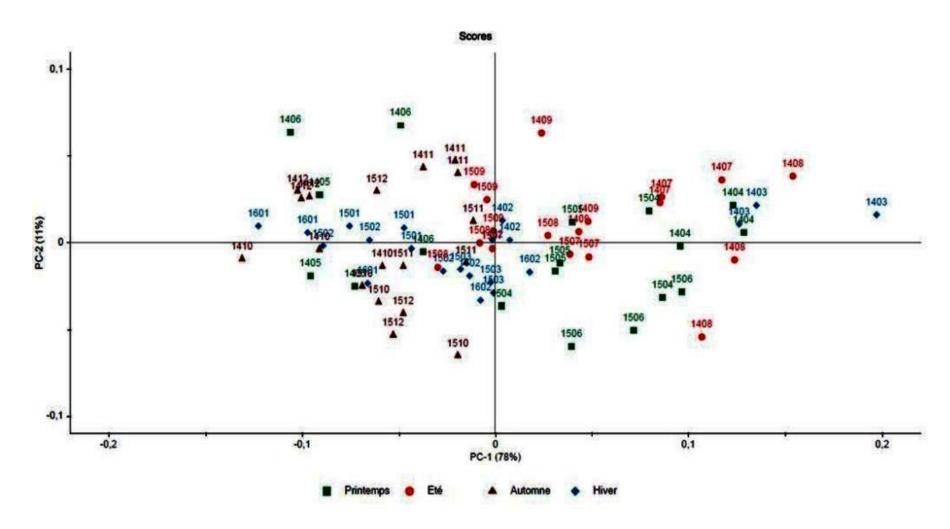


Fig 3-15 Principal component analysis of *S. muticum* sulfates group from selected wavenumbers of 1300-1015 cm-1. Green box: Spring, Red circle: Summer, Purple triangle: Autumn, and Blue diamond: Winter.

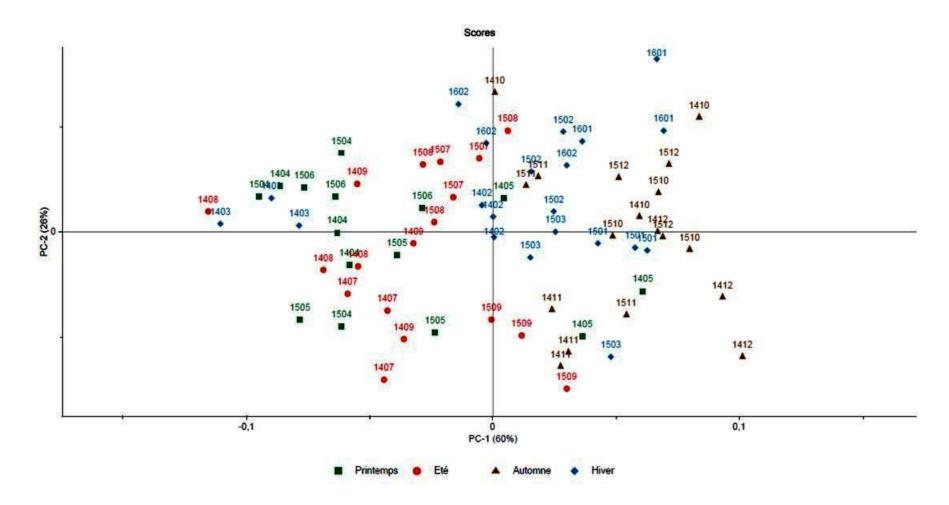


Fig 3-16 Principal component analysis of *S. muticum* polyphenol from selected wavenumbers of 3300-3200, 1620-1605, 1460-1445, 1090-1075, 935-920, 880-865, and 675-660 cm-1. Green box: Spring, Red circle: Summer, Purple triangle: Autumn, and Blue diamond: Winter.

#### Conclusion

FT-IR analysis is a helpful tool for preliminary identification of certain active compound in algal dry material. This method offers a rapid yet precise analysis. Application of FT-IR in this study is to evaluate the biochemical composition, especially the seasonal variation pattern in dry algal material, and dry enzymatic extract. PCA analysis in the seasonal variation of *S. muticum* biochemical composition reveals that season gives substantial effects towards the contents. Regarding the polyphenol in *Sargassum* enzymatic extracts, FT-IR has detected the presence of several chemical structures resembling those in phloroglucinol (standar).

In this study, the application of FT-IR is used to polyphenol, particularly phlorotannins in *Sargassum*. In addition, this method allows characterizing the seasonal effects towards *S. muticum* polyphenol. Based on the statistical analysis, there are two main group of season influencing the polyphenol production in *S. muticum*, namely the warm/hot season (spring-summer) and cold season (autumn-winter).

## 3.3.1.2 Enzymatic extract of S. muticum polyphenol compounds

Fig 3-17 presents the PCA from selected FT-IR spectra (1800-600 cm<sup>-1</sup>) of *S. muticum* enzymatic extract. Aqueous and Alcalase extract of *S. muticum* were spotted away from the other samples. Other samples that formed a distinctive group were Neutrase and Ultraflo extracts.

Viscozyme extract was chosen among the other eight as this enzyme had the highest phenolic content based on the Folin-Ciocalteu method. FT-IR managed to detect the presence of phenolic compound based on the identified wavenumbers. The liaison of polyphenol compounds detected from the raw and purified fraction of Viscozyme extract was at 3239.54, 2934.75, 1450.96, 1081.72, 930.47, and 873.72 cm<sup>-1</sup>. These wavenumbers were in accordance to those detected in phloroglucinol (standard) and from previous studies. Additional bonds which were not present in the purified fraction were detected at 1325 and

899.15 cm<sup>-1</sup> in raw Viscozyme extract, related to the C=CH<sub>2</sub> and C-H deformation bonds, respectively. In addition, the PCA analysis of *S. muticum* polyphenol from the FT-IR spectra revealed that there were two groups of variance. The first group consisted of aqueous (control), Alcalase, Neutrase, Ultraflo extracts, and the second group were AMG, Viscozyme, Shearzyme, Termamyl and Celluclast, as shown by Fig 3-18.

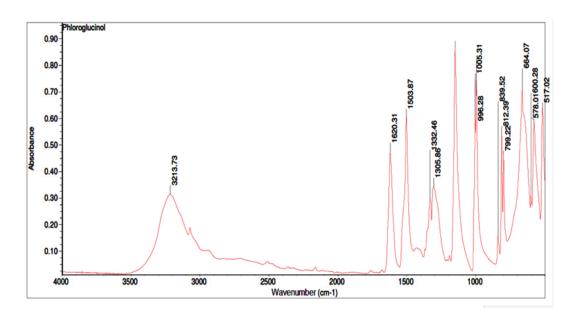


Fig 3-17. FT-IR spectra of phloroglucinol (standard).

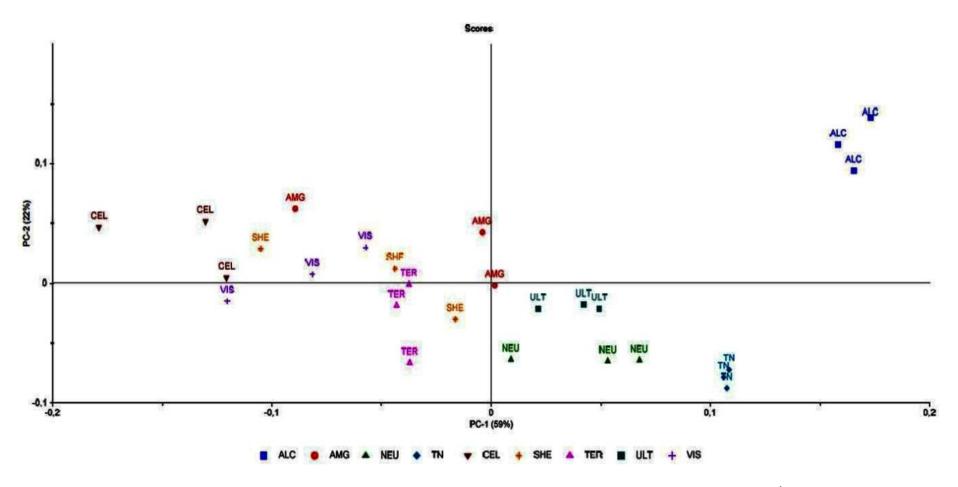


Fig 3-18 Principal component analysis of *S. muticum* enzymatic extract from selected wavenumbers of 1800-600 cm<sup>-1</sup>. ALC: Alcalase, AMG: Amyloglucosidase, NEU: Neutrase, TN: Aqueous extraction (control), CEL: Celluclast, SHE: Shearzyme, TER: Termamyl, ULT: Ultraflo, VIS: Viscozyme

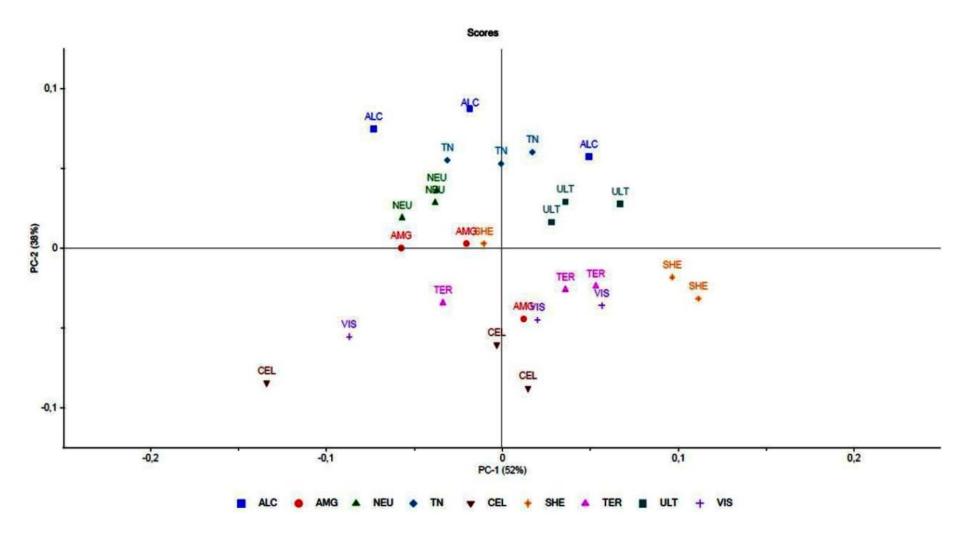


Fig 3-19 Principal component analysis of *S. muticum* enzymatic extract for polyphenol compounds from selected wavenumbers of 3300-3200, 1620-1605, 1460-1445, 1090-1075, 935-920, 880-865, and 675- 660 cm<sup>-1</sup>. ALC: Alcalase, AMG: Amyloglucosidase, NEU: Neutrase, TN: Aqueous extraction (control), CEL: Celluclast, SHE: Shearzyme, TER: Termamyl, ULT: Ultraflo, VIS: Viscozyme.

### 3.3.1.3 Enzymatic extract Indonesian *Sargassum* polyphenol compounds

Enzymatic extracts of Indonesian *Sargassum* were characterized using the Fourier-Transform Infrared (FT-IR) spectroscopy as presented on Fig 3-20 and 3-21. In general, the spectra of *S. aquifolium* and *S. ilicifolium* had the same pattern.

The spectra of *S. aquifolium, S. ilicifolium* and *S. polycystum* Viscozyme extracts ranged from 3290.89-615.76-cm<sup>-1</sup>, 3284.48-615.47-cm<sup>-1</sup>, and 3854.41-1285.15-cm<sup>-1</sup>, respectively. Then, the detected spectra of *S. aquifolium, S. ilicifolium* and *S. polycystum* Protamex extracts ranged from 3304.13-614.73 cm<sup>-1</sup>, 3277.25-615.51cm<sup>-1</sup>, and 3274.49-1259.91. Meanwhile, the phloroglucinols spectrum was detected at 3217.73, 1620.31, 1503.87, 1332.46, 1305.86, 1150.82, 1005.31, 996.28, 839.52, 812.39, 517.02, 799.22, 664.07, 600.28, and 578.01 cm<sup>-1</sup>.

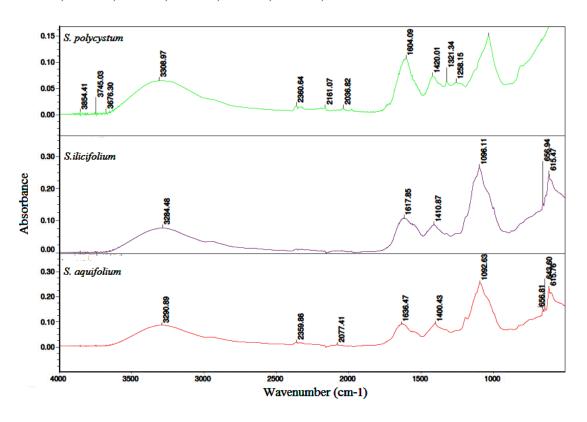


Fig 3-20 FT-IR spectra of *S. polycystum*, *S. ilicifolium*, and *S. aquifolium* extracted with Viscozyme.

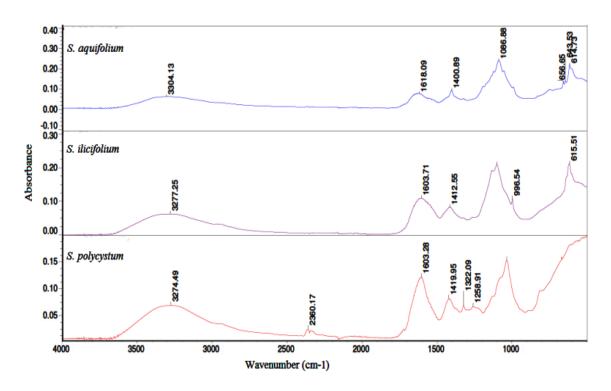


Fig 3-21 FT-IR spectra of S. aquifolium, S. ilicifolium and S. polycystum extracted with Protamex.

The presence of phloroglucinol-like (polyphenol) compounds was detected in *S. aquifolium, S. ilicifolium* and *S. polycystum*. The spectra that were assumed to correspond with this compound was 1636.47 and 3290.89 cm<sup>-1</sup> from *S. aquifolium,* 1617.85 and 3284.48 cm<sup>-1</sup> from *S. ilicifolium* and 3308.97 cm<sup>-1</sup> from *S. polycystum*. All of these spectra were from the Viscozyme extracts. As for the Protamex extracts of *S. aquifolium, S. ilicifolium* and *S. polycystum,* the spectra that corresponded to the polyphenol were 3304.13 and 1618.09 cm<sup>-1</sup>; 3277.25, 1603.71 and 996.54 cm<sup>-1</sup>; and 3274.49, 1603.28, and 1322.09 cm<sup>-1</sup>, respectively.

### Conclusion

Preliminary characterization using FT-IR showed the presence of phloroglucinol-like compounds indicating the detection of phlorotannins in enzymatic extracts. Such detection was based on the corresponding peaks obtained from literature and peak of phloroglucinol as standard. However, there was also the presence of other compounds based on their functional structure, such as carbohydrates and proteins, especially in the raw extracts.

## 3.3.2 High performance liquid chromatography (HPLC)

### 3.3.2.1 Polyphenol of S. muticum – seasonal variation

For further understanding the seasonal variation of polyphenol, the extract of *S.muticum* in different solvents were then analyzed by using HPLC as presented on Appendix 8 Apparently, HPLC chromatograms merely revealed qualitative variation of *S. muticum* polyphenol based on the peaks intensity. The intensity of these peaks varied in function of months and they were detected at 7-8 minutes and 9.5-11 minutes of retention time. Polyphenol compounds referred in this analysis were phlorotannins. In comparison to our samples, the phloroglucinols – commercial standard used to evaluate the presence of phlorotannins – had retention time at 10 minutes. Peaks of *S. muticum* extracts were detected at wavenumbers between 260 and 280 nm; 280 nm was the same wavenumbers in which phloroglucinols peak was detected.

Fig 3-22 and 3-23 are HPLC chromatogram in 3D of *S. muticum* polyphenol extracted in water from the solid-liquid extraction representing autumn, winter, spring and summer in 2014. From these four images, noticeable difference is observed between 6 to 8 min and 10 to 11 min of retention time (sphere-like forms in blue enveloped in red line). 3D image helps to illustrate the detected compound that was elucidated by the HPLC system. *S. muticum* water extracts collected in autumn, winter and spring might contained same character of compound since the HPLC managed to record the detected compound that was elucidated at 6 to 8 min. In summer, the same pattern was not observed (Fig 3-23). In addition, there was a compound detected at 10-11 min; however, it was assumed that the quantity was this compound was fluctuated. In autumn and spring, the compound was clearly distinguished in color yet it was not appeared in *S. muticum* extract collected in summer and winter. We hypothesized that the compound detected in the HPLC system might be a phloroglucinol-like compound, in other words, phlorotannins.

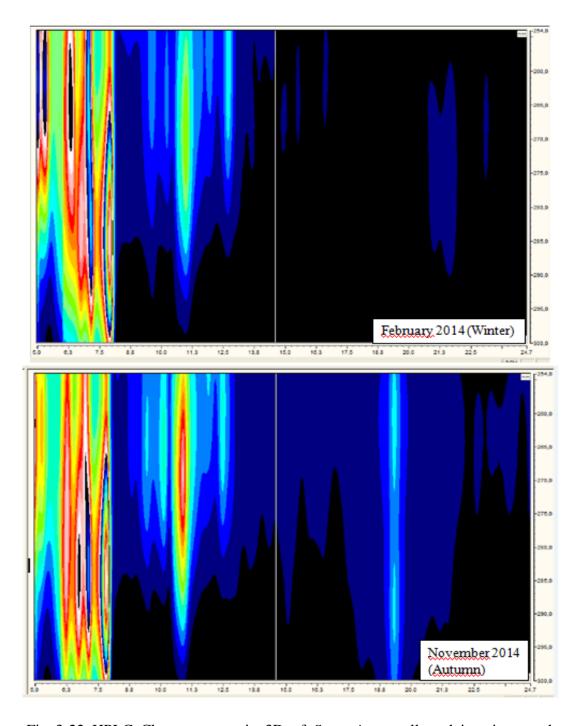


Fig 3-22 HPLC Chromatogram in 3D of *S. muticum* collected in winter and autumn. Axis represents the retention time (vertical) and wavenumbers (horizontal).

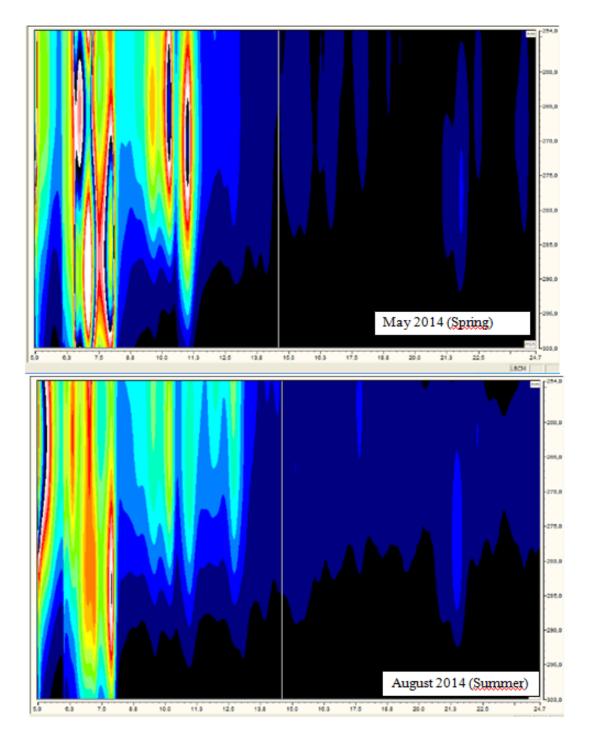


Fig 3-23 HPLC Chromatogram in 3D of *S. muticum* collected in spring and summer. Axis represents the retention time (vertical) and wavenumbers (horizontal).

### 3.3.2.2 Purified fraction of S. muticum Viscozyme extract

Chromatogram of HPLC from the raw extract and the purified fraction of *S. muticum* Viscozyme extract are presented on Fig 3-24. Based on the HPLC analysis, Viscozyme extract of *S. muticum* contained a compound with same characteristic as phloroglucinol (Fig 3-21 (Above)). This was due to the peak that was detected at the same retention time it had as phloroglucinols, *i.e.* at 10.483 min.

Purified fraction of *S. muticum* Viscozyme extract showed a well separated fraction assumed to have similar chemical structure as phloroglucinol (Fig 3-24 (Below)). We attempted to fractionate and collect the fraction from the selected peak. Unfortunately, it was not a successful attempt as the collected fractions was too diluted, thus, the quantity obtained was not sufficient for further analysis. Though, it would be interesting to further evaluate this active fraction.

#### Conclusion

HPLC analysis performed to the seasonal data of *S. muticum* polyphenol showed that the production of polyphenols was mostly affected by the sunlight, air temperature and sea surface temperature. However, the strength of this correlation varied depending on the solvents used. Water extracts had strong correlation, and aqueous methanol 50% showed a moderate correlation with sunlight, air temperature and sea surface temperature. As for methanol and aqueous ethanol 75% extracts, they demonstrated a weak correlation with those three conditions.

Regarding the pure fraction of *S. muticum* polyphenol extracted with Viscozyme, HPLC managed to separate the peak corresponding to phloroglucinol, even though the intensity was relatively low. It indicated that the eluted compound was not too diluted. As a consequence, this fraction was hardly collected after the fractionation using HPLC.

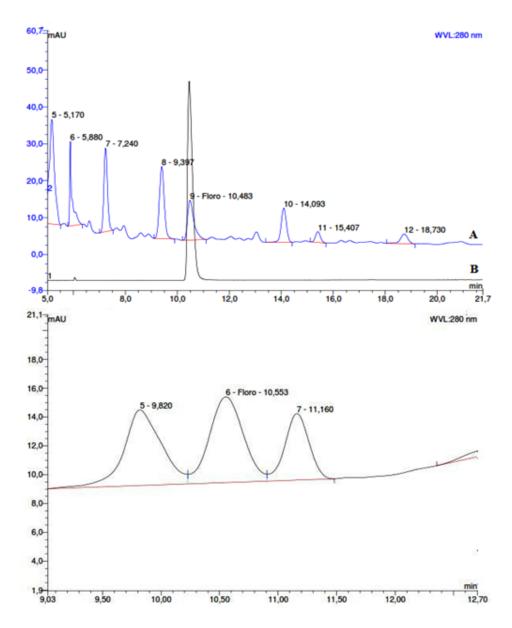


Fig 3-24 Chromatogram of *S. muticum* enzymatic extract (Above) Raw extract (A) with phloroglucinol (B) and (Below) purified fraction.

### 3.4 Antioxidant activity

### 3.4.1 DPPH radical scavenging and FRAP

## 3.4.1.1 Screening of enzymes and optimization

The antioxidant activities of *S. muticum* enzymatic extracts were determined through the free radical scavenging-ability and the reducing power by means of DPPH and FRAP analysis. Table 3-16 exhibits the antioxidant of *S. muticum* from June 2014 extracted with different enzymes at 50°C for 3 h without denaturation. All samples

showed a significant difference (p<0.05) of DPPH IC<sub>50</sub> compared to control (aqueous extract). Having low value of IC<sub>50</sub> indicates that the sample has a strong antiradical activity. In this case, Viscozyme and Celluclast extracts ( $0.6 \pm 0.1 \text{ mg mL}^{-1}$ ) would be mostly effective against the free radical compared to the other. Ultraflo ( $2.2 \pm 0.9 \text{ mg mL}^{-1}$ ) and Alcalase extracts ( $1.9 \pm 0.3 \text{ mg mL}^{-1}$ ) was less effective against the free radical as shown by their high value of DPPH IC<sub>50</sub>. Nevertheless, the free radical scavenging activity of *S. muticum* enzymatic extracts was inferior to the commercial antioxidant BHA and BHT (IC<sub>50</sub>:  $0.008 \pm 0.5 \text{ and } 0.012 \pm 4.8 \text{ mg mL}^{-1}$ ).

All samples had a significantly different reducing capacity from the aqueous extract (65.7  $\pm$  1.0  $\mu$ M of Fe<sup>2+</sup>). The most effective reducing capacity was shown by AMG extract in 60.8  $\pm$  1.8  $\mu$ M of Fe<sup>2+</sup>. In the contrary, *S. muticum* Viscozyme extract did not show a strong reducing capacity with only 48.7  $\pm$  0.9  $\mu$ M of Fe<sup>2+</sup>. The reducing power activity of *S. muticum* enzymatic extracts was still inferior to commercial Vitamin C, 58.1  $\pm$  0.7  $\mu$ M of Fe<sup>2+</sup>. Results are presented on Table 3-16. DPPH IC<sub>50</sub> of BHA and BHT and reducing power of Vitamin C as standards were 8 x 10<sup>-3</sup>  $\pm$  0.5, 12 x 10<sup>-3</sup>  $\pm$  0.5 mg mL<sup>-1</sup> and 58.1  $\pm$  0.7  $\mu$ M of Fe<sup>2+</sup>.

Table 3-16 Antioxidant activity of *S. muticum* enzymatic extracts from June 2014 (different letter superscript letters indicates a significant difference (p<0.05).

Samples	DPPH IC <sub>50</sub> (mg mL <sup>-1</sup> )	FRAP (µM of Fe <sup>2+</sup> )
BHA	$8 \times 10^{-3} \pm 0.5$	-
BHT	$12 \times 10^{-3} \pm 0.5$	-
Vitamin C	-	$58.1 \pm 0.7$
Control	$0.7 \pm 0.2^{a}$	$65.7 \pm 1.0^{a}$
Neutrase	$0.9 \pm 0.1^{a}$	$58.8 \pm 0.6^{bd}$
Ultraflo	$2.2 \pm 0.9^{c}$	$58.4 \pm 0.8^{b}$
Alcalase	$1.9 \pm 0.3^{bc}$	$56.1 \pm 3.0^{\circ}$
Celluclast	$0.6 \pm 0.1^{a}$	$59.3 \pm 1.7^{bd}$
AMG	$0.7 \pm 0.3^{a}$	$60.8 \pm 1.8^{ad}$
Shearzyme	$1.0 \pm 0.3^{ab}$	$58.4 \pm 0.7^{bd}$
Viscozyme	$0.6 \pm 0.1^{a}$	$48.7 \pm 0.9^{c}$
Termamyl	$0.8 \pm 0.2^{a}$	$57.4 \pm 1.0^{b}$

Antioxidant activity of *S. muticum* from November 2014 that was extracted with Viscozyme, Termamyl and Protamex was evaluated and results are presented on

Table 3-17. This time, algal material was extracted at 40°C for 3 h without denaturation and the enzyme concentration was increased to 5%. As for samples that were extracted with 10% enzymes, antioxidant activity was not determined.

Table 3-17 Antioxidant activity of *S. muticum* enzymatic extracts from November 2014 after optimization.

Samples	Concentration (%)	DPPH IC <sub>50</sub> (mg mL <sup>-1</sup> )	FRAP (µM of Fe <sup>2+</sup> )
Viscozyme		$7.7 \pm 1.7$	$15.5 \pm 0.4$
Protamex	5	$3.3 \pm 0.8$	$16.0 \pm 0.3$
Termamyl	3	$0.6 \pm 0.0$	$17.5 \pm 0.5$
Viscozyme		n.d	n.d
Protamex	10	n.d	n.d
Termamyl		n.d	n.d
Water		$0.7 \pm 0.0$	$31.0 \pm 2.1$

n.d: not determined

# 3.4.1.2 Indonesian *Sargassum* extracted with solid-liquid and enzyme-assisted extraction

A comparison related to the bioactivities between solid-liquid and enzyme-assisted extraction was performed. In order to compare these two methods, 40 °C was chosen as extraction condition. The fact that 40 °C is the most optimal condition for polyphenol extraction in solid-liquid becomes the main fundamental in choosing this condition. Table 3-18 present the results of antioxidant activity from Indonesian *Sargassum* extracted in two different methods.

Based on the result, Protamex extract of *S. aquifolium* had better antiradical activity by its IC<sub>50</sub>  $1.4 \pm 0.7$  mg mL<sup>-1</sup>. As for the reducing power, water extract of *S. polycystum* showed promising activity with  $46.5 \pm 5.3$  µM of Fe<sup>2+</sup>. Statistical analysis was performed to know whether there was a significant difference or not in Indonesian *Sargassum* antioxidant activity. Analysis of variance (see Appendix 11), showed that there was a significant difference (p>0.05) in antioxidant activity of Indonesian *Sargassum* due to species and method of extraction. Table 3-18 exhibits the antiradical scavenging activity of Indonesian *Sargassum* represented by IC<sub>50</sub>.

S. aquifolium and S. ilicifolium showed a significant difference in DPPH IC<sub>50</sub> (p<0.05). Later, the DPPH IC<sub>50</sub> of methanol extracts was significantly different to the aqueous methanol 50% (p<0.05). For the reducing activity, all Sargassum species showed significant difference (p<0.05). As for the solvents, water, aqueous methanol 50% and Viscozyme extracts had significant difference to each other and with the other methods (p<0.05). Water, aqueous methanol 50% and Viscozyme extracts extract were significantly different from aqueous ethanol 75%, methanol and Protamex extracts (p<0.05). Such results implied that different methods gave significant effect towards the reducing activity. Protamex showed no significant difference with methanol and aqueous ethanol 75%.

Table 3-18 Antiradical scavenging activity of Indonesian *Sargassum* in solid-liquid and enzyme-assisted extraction.

Samples	Methods	DPPH IC <sub>50</sub> (mg mL <sup>-1</sup> )	Sig. (p<0.05) of Species	Sig. ( <i>p</i> <0.05) of Solvents
BHA		7.5±0.4	_	
BHT		$11.7 \pm 4.5$		
	Water	$8.9 \pm 2.1$		
	MeOH	n.a		
	MeOH 50%	$3.2 \pm 1.7$		
S. aquifolium	EtOH 75%	n.a		
	Viscozyme	n.a		
	Protamex	$1.4 \pm 0.7$		
	Water	n.a		
	MeOH	$3.4 \pm 0.8$	04.4	0.00
	MeOH 50%	$9.0 \pm 0.7$	.014	.008
S. ilicifolium	EtOH 75%	$9.9 \pm 1.8$	(df = 2; F = 4.802)	(df = 5; F = 3.601)
	Viscozyme	$9.9 \pm 6.1$	4.802)	3.691)
	Protamex	$4.4 \pm 2.4$		
	Water	n.a		
	MeOH	n.a		
S. polycystum	MeOH 50%	$20.3 \pm 2.5$		
	EtOH 75%	$5.5 \pm 0.3$		
	Viscozyme	n.a		
	Protamex	$9.5 \pm 2.4$		

It might be possible that in certain extracts as in methanol extract of *S*. *aquifolium* with no antiradical activity, they were not active in small concentrations since the concentration used in this study ranged from 0 to 1 mg mL<sup>-1</sup>.

Table 3-19 shows the reducing antioxidant power (FRAP) activity of Indonesian *Sargassum* in solid-liquid and enzyme-assisted extraction. *S. aquifolium*, *S. ilicifolium* and *S. polycystum* showed a significant difference in FRAP activity (p<0.05). Water extract had a significant difference with the all other extracts (p<0.05). Viscozyme and aqueous methanol extracts showed significant difference with methanol, aqueous ethanol 75%, and Protamex extracts (p<0.05). Interestingly, Protamex extracts did exhibit a significant different with methanol and aqueous ethanol 75% extracts (p>0.05).

Due to different drying treatment between French and Indonesian *Sargassum*, we could not combine the analysis. As a consequence, the statistical analysis was separated. Table 3-20 presents the antioxidant activity of *S. muticum* in different methods of extraction. *S. muticum* extracted in aqueous methanol 50% and Viscozyme showed stronger antiradical activity at  $0.5 \pm 0.0$  mg mL<sup>-1</sup>. Meanwhile, water extract of *S. muticum* had more efficient reducing power with  $51.4 \pm 0.5 \mu M$  of Fe<sup>2+</sup>. DPPH IC<sub>50</sub> of *S. muticum* showed that there was no significant effect generated by different methods of extraction (p>0.05).

A significant difference in consequence of different methods was observed in the reducing activity of *S. muticum*. As shown by Table 3-20, the reducing power activity of water, methanol, aqueous methanol 50% and aqueous ethanol 75% extracts were significantly different from Protamex and Viscozyme extracts. However, there was no significant effect in the reducing power activity of Protamex and Viscozyme extract (p>0.05).

Table 3-19 Reducing antioxidant power (FRAP) activity of Indonesian *Sargassum* in solid-liquid and enzyme-assisted extraction.

Samples	Methods	FRAP (µM of Fe <sup>2+</sup> )	Sig. (p<0.05) of Species	Sig. (p<0.05) of Solvents
Vitamin C		58.1±0.7		
	Water	$21.8 \pm 1.7$		
	MeOH	$12.7 \pm 2.5$		
	MeOH 50%	$22.4 \pm 4.9$		
S. aquifolium	EtOH 75%	$14.4 \pm 0.3$		
s. aquijoiiini	Viscozyme	$33.2 \pm 2.5$		
	Protamex	$16.2 \pm 0.2$		
	Water	$24.3 \pm 1.6$		
	MeOH	$11.5 \pm 0.8$		
	MeOH 50%	$16.0 \pm 0.8$	.000	.000
S. ilicifolium	<b>EtOH 75%</b>	$10.1 \pm 1.2$	(df = 2; F =	(df = 5; F =
S. titely ettilin	Viscozyme	$15.3 \pm 1.0$	46.436)	80.014)
	Protamex	$8.6 \pm 0.6$		
	Water	$46.5 \pm 5.3$		
	MeOH	$12.7 \pm 3.1$		
	MeOH 50%	$16.9 \pm 2.6$		
S. polycystum	EtOH 75%	$13.7 \pm 2.0$		
	Viscozyme	$27.7 \pm 4.9$		
	Protamex	$16.3 \pm 0.1$		

Table 3-20 Antioxidant activity of French *S. muticum* from June 2014 polyphenol extracted in solid-liquid and enzyme-assisted extraction.

Samples	DPPH IC <sub>50</sub> (mg mL <sup>-1</sup> )	FRAP (µM of Fe <sup>2+</sup> )
BHA	$7.5 \pm 0.4$	-
BHT	$11.7 \pm 4.5$	-
Vitamin C	-	$58.1 \pm 0.7$
Water	$1.8 \pm 0.0^{a}$	$51.4 \pm 0.8^{a}$
MeOH	$1.9 \pm 0.0^{a}$	$32.1 \pm 5.0^{b}$
MeOH 50%	$0.5 \pm 0.0^{a}$	$44.9 \pm 5.4^{ab}$
EtOH 75%	$0.7 \pm 0.0^{a}$	$38.9 \pm 6.7^{ab}$
Viscozyme	$0.5 \pm 0.0^{a}$	$15.0 \pm 1.3^{c}$
Protamex	$1.6 \pm 1.3^{a}$	$11.8 \pm 1.6^{c}$

# Conclusion

Based on the screening, if a sample had strong antiradical activity, it showed an opposite reducing power capacity in which it had an inverse relationship. As an

example, Viscozyme extract of S. muticum polyphenol demonstrated a promising antiradical activity but not reducing power. From the eight commercial enzymes used, Celluclast and Viscozyme extracts exhibited similar activity based on their IC<sub>50</sub> value. On the contrary, the best reducing power activity was shown by aqueous extracts.

Concerning the antiradical activity of Indonesian *Sargassum* extracted in different methods, in certain case, enzymatic extracts showed stronger activity than the solvent extracts. However, the reducing power activity of enzymatic extracts was less effective than the organic solvents.

## 3.5 Tyrosinase and elastase inhibition activity

# 3.5.1 Screening of enzymes and optimization

First result presented in this part is the tyrosinase and elastase inhibition activities of *S. muticum* enzymatically extracted at 50 °C and denaturated at 90 °C with 2.5% of enzymes. Results are presented on Table 3-21. For the tyrosinase inhibition activity of *S. muticum* enzymatic extracts, their activity showed no significant difference (p>0.05) and their effects on tyrosinase enzyme were still low compared to kojic acid  $(67.6 \pm 6.5\%)$  of inhibition) and quercetin  $(42.7 \pm 9.4\%)$  of inhibition) – serving as positive controls. The inhibition activity of *S. muticum* enzymatic extracts ranged from  $6.6 \pm 1.5$  (Neutrase extract) to  $24.5 \pm 2.6\%$  of inhibition (Ultraflo extract).

In elastase inhibition analysis, a significant difference was only showed between the AMG ( $21.6 \pm 0.8\%$  of inhibition) and Shearzyme extracts ( $32.8 \pm 0.9\%$  of inhibition). It seems that AMG extract has the lowest activity while Shearzyme extract has the highest one. All *S. muticum* enzymatic extracts showed no significant difference with the control ( $23.1 \pm 4.8\%$  of inhibition). Furthermore, the elastase inhibition activity of *S. muticum* enzymatic extracts was weaker than that of the trypsin inhibitor ( $50.0 \pm 0.0\%$  of inhibition). Results are presented on Table 3-21. Kojic acid and Quecertin served as standard and they exhibited  $67.6 \pm 6.5\%$  and  $42.7 \pm 9.4\%$  of tyrosinase inhibition activity.

Table 3-21 Tyrosinase and elastase inhibition activity of *S. muticum* enzymatic extracts, significant difference is assigned with different superscript letter (p<0.05).

Samples	Tyrosinase Inhibition Activity (% of inhibition)	Elastase Inhibition Activity (% of inhibition)
Kojic Acid	$67.6 \pm 6.5$	-
Quecertin	42.7±9.4	-
Trypsin inhibitor	-	$50.0\pm0.0$
Control	$14.7\pm3.1^{a}$	$23.1 \pm 4.8^{ac}$
Neutrase	$6.6 \pm 1.5^{\mathrm{b}}$	$23.4\pm3.3^{ac}$
Ultraflo	$24.5 \pm 2.6^{c}$	$24.0\pm4.2^{ac}$
Alcalase	$10.7\pm3.5^{a}$	$30.9 \pm 5.9^{ac}$
Celluclast	19.9±5.1 <sup>a</sup>	$29.9 \pm 0.5^{ac}$
AMG	$15.4\pm2.3^{a}$	$21.6\pm0.8^{a}$
Shearzyme	$13.0\pm0.7^{a}$	$32.8 \pm 0.9^{bc}$
Viscozyme	$23.5 \pm 2.6^{\circ}$	$29.7 \pm 1.9^{ac}$
Termamyl	14.2±2.5 <sup>a</sup>	$28.7 \pm 3.8^{ac}$

Tyrosinase inhibition activity of *S. muticum* (extraction at 40°C with 5% of enzyme) was analyzed and results are presented on Table 3-22. Unfortunately, the activity was less appealing than the previous samples compared to the results from Table 3-21.

Table 3-21 Tyrosinase inhibition activity of *S. muticum* enzymatic extracts.

Samples	Concentration (%)	Tyrosinase inhibition activity (% of inhibition)
Kojic Acid		$83.3 \pm 5.5$
Quecertin		$46.7 \pm 3.9$
Viscozyme		$9.4 \pm 0.3$
Protamex	5	n.a
Termamyl	3	$14.8 \pm 0.8$
Viscozyme		n.d
Protamex	10	n.d
Termamyl	10	n.d
Water		n.d

n.d: not detemined

# 3.5.2 Enzymatic extract of Indonesian Sargassum

Tyrosinase inhibition activity of *Sargassum* enzymatic extracts was then analyzed and the results are presented on Table 3-23. *S. ilicifolium* extracted with Viscozyme showed no inhibition. Meanwhile, *S. aquifolium and S. polycystum* Viscozyme extracts showed an inhibition of  $11.5 \pm 9.7\%$  and  $10.2 \pm 4.1\%$ . For Protamex extracts, the tyrosinase inhibition activity of *S. aquifolium*, *S. ilicifolium* and *S.polycystum* were  $9.7 \pm 5.0$ ,  $8.2 \pm 1.1$  and  $13.3 \pm 6.6\%$ , respectively. Statistical analysis showed that there was no significant difference (p > 0.05).

Table 3-22 Tyrosinase inhibition activity of Indonesian *Sargassum* enzymatic extracts.

Samples	Tyrosinase inhibition activity (% of inhibition)		
Kojic Acid	$83.3 \pm 5.5$		
Quecertin	$46.7 \pm 3.9$		
	Viscozyme	<b>Protamex</b>	
S. aquifolium	$11.5 \pm 9.7$	$9.7 \pm 5.0$	
S. ilicifolium	n.a.	$8.2 \pm 1.1$	
S. polycystum	$10.2 \pm 4.1$	$13.3 \pm 6.6$	

n.a.: no activity

## Conclusion

Tyrosinase and elastase inhibition activity of *S. muticum* polyphenol extracted with enzymes might show a weak activity, same case with Indonesian *Sargassum*. However, these results revealed the potential of enzymatic extract that needs further study as whitening and anti-aging agent.

## 3.6 Antimicrobial activity

#### 3.6.1 Antibacterial activity

Based on the antibacterial activity of *S. muticum* enzymatic extract against four pathogenic bacteria, i.e *S. hominis*, *B. subtilis*, *P. aeruginosa* and *E. coli*, all samples showed no remarkable activity (data was not shown). We assumed that the presence

of polysaccharides in our raw enzymatic extracts provoked the growth of the bacteria instead of suppressing their growth.

Antibacterial activity of *S. muticum* extracts from two methods of extractions was evaluated against *E. coli*, *B. subtilis* and *P. aeruginosa*. Less appealing results were obtained from the enzymatic extracts of *S. muticum*. They did not show any activity against the three bacteria (data was not shown); as a consequence, further analysis was not performed for the other enzymatic extracts.

Table 3-24 presents solely the antibacterial activity of *Sargassum* extracted using different solvents *against E. coli, B. subtilis* and *P. aeruginosa. S. aquifolium* in aqueous methanol 50% and *S. ilicifolium* in both methanolic extracts showed no activity against *B. subtilis* and *P. aeruginosa*, respectively.

Table 3-23 Antibacterial activity of *S. aquifolium*, *S. ilicifolium*, *S. polycystum* and *S. muticum* against *E. coli*, *B. subtilis* and *P. aeruginosa*.

C	G-14	Antik	ivity (cm)	
Samples	Solvents	E. coli	B. subtilis	P. aeruginosa
Phosphomycin		$2.2 \pm 0.1$	$1.7 \pm 0.1$	$0.9 \pm 0.1$
Ampicilin		$1.7 \pm 0.1$	$1.6 \pm 0.1$	$0.5 \pm 0.0$
Streptomycin		$1.8 \pm 0.2$	$1.1 \pm 0.2$	$2.7 \pm 0.1$
	Water	+	+	+
C	MeOH	+	+	+
S. aquifolium	MeOH 50%	+	n.a.	+
	EtOH 75%	+	+	+
	Water	+	+	+
G :1: :C 11:	MeOH	+	+	n.a.
S. ilicifollium	MeOH 50%	+	+	n.a.
	EtOH 75%	+	+	+
	Water	+	+	+
C 1	MeOH	+	+	+
S. polycystum	MeOH 50%	+	+	+
	EtOH 75%	+	+	+
	Water	+	+	+
a	MeOH	+	+	+
S. muticum	MeOH 50%	+	+	+
	EtOH 75%	+	+	+

n.a. = no activity

### 3.6.2 Biofilm inhibition activity

## 3.6.2.1 Screening of enzymes and optimization

For the biofilm generated from P. aeuriginosa, there was no sign of inhibition from the control, Alcalase, Celluclast and AMG extracts. Ultraflo and Termamyl extracts presented a weak inhibition with  $3.7 \pm 2.0\%$  and  $12.4 \pm 3.9\%$ , respectively. Highest activity was shown by S. muticum enzymatic extracts Viscozyme with  $43.7 \pm 4.2\%$  of inhibition. All S. muticum enzymatic extracts were less effective in inhibiting the formation of biofilm formed by P. aeruginosa compared to E. coli, as can be seen in Table 3-25.

Strong biofilm inhibition activity formed by  $E.\ coli$  was performed by Shearzyme extract followed by Termamyl extract with  $64.3 \pm 10.3\%$  and  $49.2 \pm 13.3\%$ , respectively. In addition, their activity was significantly better than the control,  $14.7 \pm 0.9\%$ . Based on the biofilm inhibition activity of  $S.\ muticum$  enzymatic extract, the value presented might indicate a promising potential of  $S.\ muticum$  enzymatic extracts as an antifouling agent.

Table 3-24 Biofilm inhibition of *S. muticum* enzymatic extract (expressed in % of inhibition, different superscript letters indicates a significant difference (p < 0.05)).

-	_	
Samples	P. aeruginosa	E. coli
Control	n.a	$14.7 \pm 0.9^{a}$
Neutrase	$36.1 \pm 7.8^{a}$	$29.1 \pm 10.0^{ac}$
Ultraflo	$3.7 \pm 2.0^{b}$	$32.8 \pm 3.7^{ac}$
Alcalase	n.a	$38.0 \pm 7.9^{b}$
Celluclast	n.a	$33.8 \pm 7.7^{a}$
AMG	n.a	$43.2 \pm 5.9^{be}$
Shearzyme	$26.9 \pm 2.1^{b}$	$64.3 \pm 10.3^{de}$
Viscozyme	$43.7 \pm 4.2^{a}$	$28.8 \pm 9.4^{a}$
Termamyl	$12.4 \pm 3.9^{b}$	$49.2 \pm 13.3^{\text{be}}$

Table 3-26 describes biofilm inhibition of *S. muticum* enzymatic extracts after the optimization. In this work, *B. subtilis* was added as biofilm-forming bacteria that were tested against *S. muticum* enzymatic extracts. Interestingly, 5% of enzymes managed to inhibit the biofilm formation generated by *E. coli* and *B. subtilis* until

 $40.3 \pm 0.3$  and  $59.0 \pm 0.4\%$  of inhibition, respectively. In the contrary with *E. coli* and *B. subtilis*, the enzymatic extracts were less effective against *P. aeruginosa* as they had inhibition activity ranging only from  $13.3 \pm 0.1$  to  $26 \pm 0.2\%$  of inhibition. Unfortunately, due to time limits, further bioactivities analysis of *S. muticum* extracted with 10% of enzymes could not be conducted.

Table 3-25 Biofilm inhibition activity of *S. muticum* enzymatic extracts (40°C, 3h, without denaturation) (expressed in % of inhibition).

Samples	Concentration (%)	P. aeruginosa	E. coli	B. subtilis
Viscozyme		$13.3 \pm 0.1$	$40.3 \pm 0.3$	$59.0 \pm 0.4$
Protamex	5	$26.3 \pm 0.2$	$29.8 \pm 0.6$	$40.0 \pm 0.4$
Termamyl		$16.3 \pm 0.2$	$36.0 \pm 0.5$	$42.3\pm0.2$
Viscozyme		n.d	n.d	n.d
Protamex	10	n.d	n.d	n.d
Termamyl		n.d	n.d	n.d
Water		n.d	n.d	n.d

n.d; not determined

## 3.6.2.2 Enzymatic extract of Indonesian Sargassum

Biofilm inhibition activity of Indonesian Sargassum enzymatic extracts formed by P. aeruginosa, E. coli and B. subtilis was evaluated as can be seen on Table 3-27. S. aquifolium, S. ilicifolium and S. polycystum extracted with Viscozyme and Protamex were more effective against the film formation generated by B. subtilis compared to others. S. ilicifolium exhibited the highest inhibition activity with  $49.0 \pm 6.8\%$ . However, Protamex extract of S. polycystum was also showed a promising inhibition against the film formation of E. coli with  $45.8 \pm 3.8\%$ . The lowest inhibition activity was exhibited by the Viscozyme extract of S. ilicifolium against the film formation of P. aeruginosa with  $9.1 \pm 3.0\%$  of inhibition. Based on the statistical analysis, there was a significant difference between the three species tested and between the types of extractant that effected the biofilm inhibition activity.

Table 3-26. Biofilm inhibition activity of Indonesian Sargassum enzymatic extracts (expressed in % of inhibition).

Samples	P. aeruginosa		P. aeruginosa E. coli		B. subtilis	
	Viscozyme	Protamex	Viscozyme	Protamex	Viscozyme	Protamex
S. aquifolium	$11.6 \pm 2.3$	$16.3 \pm 4.5$	$26.3 \pm 11.0$	$9.1 \pm 5.3$	$42.7 \pm 9.6$	$44.5 \pm 11.0$
S. ilicifolium	$9.1 \pm 3.0$	$9.9 \pm 4.8$	$20.0 \pm 11.8$	$18.2 \pm 3.1$	$25.2 \pm 14.8$	$49.0 \pm 6.8$
S. polycystum	$11.5\pm6.1$	$12.0\pm2.7$	$14.5 \pm 4.4$	$45.8 \pm 3.8$	$36.9 \pm 12.9$	$40.7 \pm 13.5$
	Sig. $(p<0.05)$ of Species .207 (df = 2; F = 1.647)		Sig. $(p<0.05)$ .025 (df = 1	•	~ <b>.</b>	5) of Bacteria 2 ; F = 51.724)

## 3.6.3 Cytotoxicity and Antiviral activity of S. muticum enzymatic extract

Cytotoxicity test had revealed that they did not have a toxic effect towards the tested cell, *i.e.* Vero cell kidney of green monkey). As for their potential as an antiviral agent against the HSV-1, only *S. muticum* Neutrase and Alcalase extracts that showed a potential antiviral activity with EC<sub>50</sub> at 430.1  $\pm$  16.3 and 225.1  $\pm$  23.3 µg mL<sup>-1</sup>. Zovirax showed no toxicity (CC<sub>50</sub>>500) and its EC<sub>50</sub> was at 0.7  $\pm$  0.3 µg mL<sup>-1</sup>, results are presented in Table 3-28.

Table 3-27 Cytotoxicity and antiviral activity of S. muticum enzymatic extracts

Samples	CC <sub>50</sub> (µg mL <sup>-1</sup> )	EC <sub>50</sub> (μg mL <sup>-1</sup> )
Zovirax	>500	$0.7 \pm 0.3$
Control	>500	>500
Neutrase	>500	$430.1 \pm 16.3$
Ultraflo	>500	>500
Alcalase	>500	$225.1 \pm 23.3$
Celluclast	>500	>500
AMG	>500	>500
Shearzyme	>500	>500
Viscozyme	>500	>500
Termamyl	>500	>500

 $CC_{50}$ : Cytotoxic Concentration 50% - a concentration of product generating 50% of cell viability compared to control.  $EC_{50}$ : Antiviral Effective Concentration – a concentration of product causing 50% of inhibition of HSV-induced destruction towards the cells. n.a = no activity detected.

## Conclusion

We acknowledged that the enzymatic extracts of *Sargassum* might have the ability as a probiotic agent instead of antibacterial agent. In comparison with the organic solvent extract, enzymatic extract showed no sign of inhibition. Leading to the assumption that sugar constituents in the enzymatic extracts serve as source of nutrition for the bacteria. However, the enzymatic extracts of *Sargassum* could inhibit the formation of biofilm. Some extracts had promising performance in inhibiting the biofilm with high percentage of inhibition, yet some showed less appealing activity. Considering that the enzymatic extracts of *S. muticum* had no

potential of toxicity towards the Vero cell, it implies the possibilities to develop this method further in *Sargassum*. Additionally, *Sargassum* is marine brown algal with a wide range of pharmacological properties. Present study unveiled the potential of *Sargassum* enzymatic extracts as antifouling and antiviral agents.

# 3.7 Polyphenol content, and bioactivities of purified fraction

Extraction yield of SPE from the aqueous methanol fraction of Viscozyme extract was  $0.5 \pm 0.0\%$  of dry algal material containing  $51.5 \pm 0.6\%$  polyphenol. It showed that there were 10 fold polyphenol content compared to the raw extract  $(6.4 \pm 0.5\%$  dry extract). Antioxidant activity of this fraction was also improved. Value of DPPH IC<sub>50</sub> was  $0.012 \pm 0.6$  from  $0.6 \pm 0.1$  mg mL<sup>-1</sup>. Reducing power capacity of this fraction improved as well with  $62.4 \pm 2.0$  from  $48.7 \pm 0.9$   $\mu$ M of Fe<sup>2+</sup>.

Table 3-29 Polyphenol content and bioactivities of purified fraction.

S. muticum Viscozyme extract	Polyphenol (% of dry extract)	DPPH (IC <sub>50</sub> (mg mL <sup>-1</sup> )	FRAP (µM of Fe <sup>2+</sup> )	Tyrosinase inhibition activity (% of inhibition)	CC <sub>50</sub> (µg mL <sup>-1</sup> )	EC <sub>50</sub> (μg ◊)
Purified fraction	$51.5 \pm 0.6$	$12 \times 10^{-3}$ $\pm 0.6$	$62.4 \pm 2.0$	$83.2 \pm 7.9$	>500	259.3 ± 9.6
Raw extract	$5.2 \pm 0.1$	$0.6 \pm 0.1$	$48.7 \pm 0.9$	$23.5 \pm 2.6$	>500	>500

#### Conclusion

Combination of enzyme-assisted and solid-phase extraction might improve the recovery of polyphenol compounds in brown algae. Working with purified samples has helped to learn about the bioactivities of target compound, in this case, phlorotannins. It was not only enhanced the quantity of polyphenol content but also improved its bioactivities.

## 3.8 Discussion

## 3.8.1 Biochemical composition of Sargassum

Valorization of marine algae at any significant scale will become a new endeavor and oblige a comprehensive insight as preliminary assessment for a sustain mass production. Although algae have been recognized as a source of variability regarding its chemical profiles, the natural availability or biomass, the impact of natural environmental variation on the presence, form and level of its chemical profile requires attention. Most of algae exhibit great plasticity regarding to biochemical composition in response to a single or combinations of multiple environmental factors (Stengel *et al.*, 2011). This plasticity basically helps them to survive, grow and reproduce in facing numerous important environmental variables like temperature, salinity, desiccation, hydrodynamics and wave exposure, nutrients, carbon dioxide and pH (Harley *et al.*, 2012). It obviously explains the importance of environmental towards the chemical, biological and physiological of algae. Thus, there have been quite studies that concerns about the effect of environmental in the biochemical composition of algae and so did in this study.

By evaluating the biochemical compositions of algae as well as its seasonal variation, it provides several advantages for further application. First, the biochemical compositions of algae represent its nutritional value which is commercially valuable for food industries. Therefore, they have been considered as food supplement for this century as they contain proteins, lipids, polysaccharides, minerals, vitamins and enzymes (Kilinç *et al.*, 2013). Second, due its plasticity towards its surrounding, it was without any doubt the quantity of algae biochemical composition strictly depended on its environmental variables. In some instances, these environmental effects have been previously described to be influential as well to the bioactivity level of algae as reported by many studies (Marechal *et al.*, 2004; Stirk *et al.*, 2007; Kulanthaiyesu and Sivakumar, 2012; Tanniou *et al.*, 2013b). For that reason, it becomes crucial to collect algae at best

period when their chemical constituents are produced in their optimal condition. A seasonal study on algae biochemical composition allows providing the required information particularly for the valorization point of view.

In one of our work, we attempted to evaluate the biochemical composition of tropical brown algae represented by *S. aquifolium*, *S. ilicifolium* and *S. polycystum*. Additionally, seasonal variation of *Sargassum* biochemical composition was also evaluated by using *S. muticum* as a model. Unlike the previous three species, *S. muticum* that was analyzed in this study came from the temperate region.

## 3.8.1.1 Indonesian Sargassum

In general, brown algae contain 5-15 % of proteins, ~ 4% of lipids, ~ 40% of ash, ~ 60% of polysaccharides, and 1-15 % of polyphenols (Park *et al.*, 2000; Burtin, 2003; Lee, 2008; Harnedy and Fitzgerald, 2011; Kim, 2012). Handayani *et al.* (2004) reported that the average of protein content of *S. crassifolium* was 5%. Supirman *et al.* (2012) obtained almost the same value of protein content of *S. filipendula*, i.e. 5%. Kawaroe *et al.* (2013) stated that the protein content of *S. crassifolium* was 8 % of dry weight. *S. ilicifolium* contained 9% of protein (Ganapathi *et al.*, 2013). In this study, the analyzed protein content in all Indonesian *Sargassum*, *S. aquifolium*, *S. ilicifolium* and *S. polycystum*, was higher. However, hot-water extract of *S. hemiphyllum* had 38.5% of proteins (Hwang *et al.*, 2010). It was even higher than our findings. Unfortunately, the authors did not explain further regarding this high protein content. It is probably due to method of extraction. Authors used hot-water extraction at 100 °C for 30 min.

Total sugar detected in *S. aquifolium, S. ilicifolium* and *S. polycystum* was lower than that of the previous studies with same species. *S. ilicifolium* contained 38% of carbohydrates (Ganapathi *et al.*, 2013). *S. polycystum* contained 34% of carbohydrates (Ahmad *et al.*, 2012). Same content of carbohydrates was also reported by Matanjun *et al.* (2009) from *S. polycystum* with 34%. *S. polycystum* 

had 5% of (Matanjun *et al.*, 2009). Shao *et al.* (2014) studied the chemical composition of different sulfated polysaccharides fractions from *S. horneri* in order to analyze their antioxidant and antitumor activity. Based on the result, the total sugars from three fractions ranged from 19-69% and 11-19% for sulfates. *S. filipendula* crude fucoidan was reported to contain 1.2% of uronic acid, 8.9% of total sugars and 5.3% of sulfates groups (García-Ríos *et al.*, 2012). Uronic acid and sulfates groups content found of Indonesian *Sargassum* in this study were higher.

Ash of *Sargassum sp.* is 34.6% (Yunizal, 2004). In *S. vulgare*, there are 18.5% of ash content (Marinho-Soriano *et al.*, 2006). In recent study, ash content of *S. aquifolium*, *S. ilicifolium* and *S. polycystum* ranged from 28 to 38% of dry algal material. Ash content of *S. polycystum* was 42% (Matanjun *et al.*, 2009) and 21% of dry algal material (Ahmad *et al.*, 2012).

Study by Ahmad *et al.* (2012) reported that *S. polycystum* contained 2.3 % of polyphenols. Budhiyanti *et al.* (2012) studied on antioxidant capacity of several *Sargassum* species collected from the coastline of Java. They reported that the total phenolic content of *Sargassum* sp. varied from 0.006-0.65% (in cytoplasmic) and 6.7-22% of dried extract (in membrane bound). This previous study showed slightly higher phenolic contents than in this study as it was extracted specifically from the membrane bound. Surprisingly, the polyphenols content found by Ganapathi *et al.* (2013) was similar with recent study, i.e. 2%.

As components of algae experience geographical, inter- and intra-individual variations due to fluctuations in the environmental conditions as well as season, these contents change, either lower or higher. Temperature, salinity, pH, light levels and quality, nutrient levels and composition, contaminants, CO<sub>2</sub> availability and biotic factors due to the distribution of grazers and endo- and epibionts are the fluctuations as mentioned by Stengel *et al.* (2011). These results implied that there were differences in *Sargassum* biochemical composition. Such differences might be due to the abiotic and biotic factors influencing the production of these

constituents (Manivannan *et al.*, 2008). Another hypothesis was that the method used to extract the chemical constituent of *Sargassum* was not the suitable one. Consequently, the obtained results did not represent the biochemical composition of *Sargassum* in general, especially for the polysaccharides. Thus, it would be interesting to determine the polysaccharides of brown algae by using the appropriate method of extraction in the next future.

#### Conclusion

Biochemical composition of Indonesian *Sargassum* shows inter-species and geographical variation. In addition, seasonal and environmental condition such as temperature, salinity, availability of nutrients and many more, might contribute to the variation of *Sargassum* biochemical composition. Indonesian *Sargassum*, like most *Sargassum* species, is characterized by its high content of ash. Among three species, *S. polycystum* has the highest ash content. Ash content reflects the total mineral of an individual and it is beneficial for the fodder or animal feed industry. Beside ash content, this species is also high in sulfates groups; it indicates the possibility of high fucoidan. Perhaps, this could become preliminary information that this species might contain interesting fucoidan yield that is potential for further application. As for *S. ilicifolium*, it has high uronic content that can be associated with its alginate. Probably, this species has its advantage for hydrocolloid industry.

## 3.8.1.2 Seasonal variation of *S. muticum* biochemical composition

Dry material of *S. muticum* revealed in this study was up to 25%. In previous study, the dry material of *S. muticum* ranged from 8 to 19% of total weight (Balboa *et al.*, 2016). Dry material of *S. muticum* was affected by season, in which, spring and summer had higher percentage of dry material than autumn and winter. Seasonal pattern of dry material in *Sargassum* did not show same pattern. In certain way, dry material of *S. muticum* collected in February to July 2014 was higher than the same months in 2015.

Biochemical composition of S. muticum had been significantly affected by the seasons. Protein, nitrogen and uronic acid content of S. muticum increased during winter and spring; while sugar, sulfates groups, and polyphenol increased in summer. Interestingly, the seasonal variation study of S. muticum biochemical composition in this study revealed the tendency as expected. Prior to analysis, we assumed that the seasonal variation would give a significant effect to the biochemical composition of S. muticum. Besides the seasons, it might be possible that the variation shown by S. muticum also followed its bio-physiological functions related to its life cycle. Early fall or autumn is known to be the period when S. muticum starts their reproduction followed by the growth of small individuals from October-November (Deysher, 1984; Loraine, 1989). It reaches its optimal growth in the spring then it stops in summer and begins to degrade during this season. This cycle is in accordance as our observation during the collection of S. muticum. From September-October, there were no long thalli found along the study site. Most of them were less than 5 cm long – also called as perennial primary shoot; therefore, it was hard to obtain a substantial biomass. Then, in November-February, the primary shoot started to grow and the length of thalli collected ranged from 5-15 cm, approximately. Early spring, which was on March, until May, the biomass of S. muticum was quite abundant and fertile as well. During this period of collection, the presence of receptacles on S. muticum thalli was observed. The thalli were ramified and their primary lateral reached to 20 cm long bearing numerous small blades and air bladder. Meanwhile, on June to August, the blades and air bladder of S. muticum began to fall out leading to a loss of buoyancy (see Appendix 1). A numerous presence of epiphytes such as gastropods and other algae were remarked on the thalli making them hard to wash and clean (see Appendix 1). This observation allows us to establish preliminary correlation between the seasonal variation of S. muticum biochemical composition and its life cycle. Unfortunately, the correlation made in this study was merely based on general description since we did not evaluate specifically the effect of S. muticum life cycle to the biochemical composition. This kind of correlation needs a comprehensive study involving the measurement of growth rate, morphological study, biochemical composition analysis and an *in situ* and *ex-situ* time-series environmental variables. In the meantime, it is most likely that the correlation between life cycle and biochemical composition of *S. muticum* is positively related. In theory, sugar content is high during summer and autumn.

Protein synthesis conducted by algae is through the fixation of atmospheric nitrogen and protein. Thus, the protein content in marine algae varies depending on the harvest period. Light, temperature and salinity are environmental factor influencing protein synthesis. Souchet (2004) in Kim (2012) reported the correlation between protein and water nitrogen content as high temperature decrease dissolved nitrogen content in water column. As consequence, the protein content in marine algae is lower during summer as the nitrogen fixation declines in this season. Higher temperature increases consumption energy due to algal respiration. Protein content is relatively lower in summer and higher in winter (Kim, 2012). The protein of S. muticum in this study showed the same tendency as well as nitrogen. In addition, the correlation between S. muticum protein and sea surface temperature was a significantly inversed yet weak correlation. Balboa et al. (2016) reported that the protein content of S. muticum collected in May to June contained 7% of protein and from November to January contained 11% of protein. Additionally, the greater protein content occurred during the decrease of temperature. In other words, there was an inverse correlation between protein content and temperature (Balboa et al., 2016).

In contrary with protein and nitrogen, the content of *S. muticum* polysaccharides was higher in summer than in winter and spring. It suggested that from winter to spring, the low content of polysaccharides was because *S. muticum* utilized them for its growth. In the early summer when it stopped to grow, the polysaccharides served as a reserve. Our result was in accordance with Schiener *et al.* (Schiener *et al.*, 2014) revealing that the carbohydrates in Laminariales were found to accumulate during summer to fall and decrease in winter as they were used as an energy source for new tissue growth. In addition, this accumulation

might also be due to the presence of herbivore or other environmental stress since polysaccharides serve as defense system in marine algae.

Hardouin *et al.* (2013) noted 29.3% of protein content of *S. muticum* was yielded from hydrolysis extracts. Same authors also reported that the compositions of ash, total sugars, uronic acid, sulfates groups and total phenol of *S. muticum* were 27.4%, 9.8%, 8.0%, 5.6% and 2.1%. These results were slightly in accordance with our finding in this study, except for uronic acid and sulfates group. Meanwhile, Jard *et al.* (2013) observed that *S. muticum* had 16.6% of total sugars, 13.6%, 8.4% of protein, 3% of sulfates and 1.9% of phenols.

As for the uronic acid and sulfates groups, these chemical entities highly related to the alginates and fucoidan content in brown algae, respectively. We found that the seasonal variation was not significantly different. Despite of this discouraging result, there was a slight increase of uronic acid and sulfates group in summer. Hypothetically, this tendency was in line with polysaccharides content. Analysis of correlation between the sulfates groups showed that there was a moderate correlation with the sea surface temperature, air temperature and sunshine. Unlike the sulfates, the uronic acid had a weak correlation with these variables.

In Saccharina japonica, S. pallidum, and Stephanocystis crassipes, their fucoidans were mainly influenced by the development of the reproductive structures of the algae itself. It was suggested that the fucoidans did not have a direct relationship with illumination regime, salinity and concentrations of biogenic substances in water (Skriptsova, 2016). Later, they discovered that fucoidan of S. pallidum was found in its maximum yield in July (summer) (Skriptsova, 2016). Skriptsova et al. (2010) also found that fucoidan yield of Undaria pinnatifida increased from April to July. As for the alginate content, represented with its uronic acid, Rodrigues-Montesinos et al. (2008) observed an increase of alginate in the cell wall and intercellular matrix of S. sinicola in spring as they matured. Then, they were low in alginate during winter since they were

still in juveniles phase. In 2004, study by the same authors reported that the alginate content in whole plant of *U. pinnatifida* increased from March to July followed by a decrease in August. In addition, alginate content of *U. pinnatifida* blades was maximal during spring with 51% and lowest in July with 34%. Increase of alginate content coincides with plant maturation and sporophyte development (Skriptsova *et al.*, 2004). Sugar, and sulfate groups content of *S. muticum* showed quite the same pattern as fucoidan and alginate from other species reported by previous studies. These entities showed an increase in spring-summer then followed by a decrease in fall-winter.

Phenols of S. muticum exhibited a significant effect given by the season, i.e. between summer-fall-winter and spring with the highest amount of polyphenols was recorded during fall. Looking further to the correlation analysis, we obtained that the polyphenols of S. muticum showed a weak correlation with the sea surface temperature, sunshine, air temperature and precipitation. It brought an assumption that polyphenols might have a more complex relationship. Theoretically, polyphenols in brown algae are mostly induced by the presence of predators, the exposure to sunlight, the fluctuation of environmental factors (temperature, lights, nutrient, etc.) and water pollution (Toth and Pavia, 2000; Jormalainen et al., 2003; Koivikko, 2008; Connan and Stengel, 2011). However, it is important to note that they also have primary role as one of the compounds constituting the cell wall of brown algae (Lee, 2008). Thus, they also play a crucial role in algae metabolism and there is a trade-off between polyphenols and annual growth of algae. Accordingly, there are many factors contributing to the production of polyphenols in brown algae. Regardless of these matters, the current study concerned only to the seasonal variation.

Phenols of *S. muticum* from the Callot Island and Point de Diable, coast of Brittany (France) showed an increase occurring between winter and spring in 2003 and 2004. The highest phenols were recorded in spring 2003 and the lowest one was in winter and fall. Meanwhile, at the Point de Diable, the highest phenols were observed in summer 2004 and a decrease to a minimal value was remarked

in winter 2003 (Plouguerné *et al.*, 2006). In this study, the highest phenols were found in fall season. Our assumption suggested that the difference shown in this study was due to different location and perhaps different method of extraction.

The ash content of brown algae is higher compared to most terrestrial vegetables containing from 5 to 10% (Rupérez, 2002). In addition, between the three groups of algae, the ash content of brown algae surpasses the green and red algae. Variations in ash content usually related to the seasonal variations of inorganic compounds and salts in the aquatic environments such as salinity (Kumar *et al.*, 2015). Same authors reported the ash content of *S. wightii* was higher in July and lower in November. In contrary with study conducted by Kumar *et al.* (2015), *S. muticum* ash content was noticed to be high from June till October, while the lowest was from March till May (Balboa *et al.*, 2016). In Japan, *S. horneri* contained 32% of ash in May and 25% in March (Murakami *et al.*, 2011). Present study revealed that the best ash content of *S. muticum* was found during fall and winter then it decreased from spring to summer. Such pattern was quite the same as presented by Balboa *et al.* (Balboa *et al.*, 2016) even though there was a time deviation assuming due to the different locality of *S. muticum*.

Secondary analysis using the Fourier Transform Infrared spectroscopy – abbreviated as FTIR – was conducted as a complementary analysis of colorimetric method. This qualitative analysis helped to describe clearly the seasonal distribution pattern of *S. muticum* biochemical composition and its seasonal grouping as presented on the results. The seasonal grouping managed to distinguish the tendency between the cold (fall and winter) and hot (spring and summer) seasons. However, it is crucial to carefully select the wavelengths corresponding to the target compounds. Even though our samples have been dried with the freeze-drying but they still contain some water.

Water absorbs mostly at 3700-3100 cm<sup>-1</sup> based on the table of infrared or to be precise at 3720-2950 cm<sup>-1</sup> - absorption of -OH of pure water with strong

intensity. Rodrigues *et al.* (2015) suggested to choose a global analysis based on the spectra at 1800-600 cm<sup>-1</sup>. This range of spectra is assumed to be the most informative. Moreover, there is also a noise area that related to the crystal of the FTIR at 2200-2000 cm<sup>-1</sup>. The global analysis mentioned here was related to the PCA analysis. Result revealed that the grouping of PCA using the spectra at 1800-600 cm<sup>-1</sup> distinguished the season into two groups, *i.e.* cold and hot season (Appendix 6). Such grouping was similar to the grouping of spectra particularly selected for different biochemical composition of *S. muticum*.

Detected peak at 3270-3250, 2940-2925, 2300-1900, 1660-1600, 1500-1200 and 1100-1000 cm<sup>-1</sup> were wavelengths showing interesting peak based on the global spectra of *S. muticum*. 3270 and 3250 cm<sup>-1</sup> suggested the presence of two hydrogens bonded (SO<sub>2</sub>N-H) (Gennari *et al.*, 1998). Wavelengths at 3600-3450 cm<sup>-1</sup> is assigned as the majority of hydroxyl absorption bands in the region intra-molecular hydrogen bonds (Rodriguez, 2010) and the hydroxyl peak usually occurs at 3545 cm<sup>-1</sup> (Matthews, 2007). Chen *et al.* (2015) adds that absorbance peak at 3600-3450 cm<sup>-1</sup> ascribed to the release of HCN. Free amine functionalities produce bands either at 3500-3300, 1650-1580, or 1622 cm<sup>-1</sup>, while N-H bonds absorb at 3350-3070, 1650-1550 or 1594 cm<sup>-1</sup> (Hamid, 2005). Strong bands at 1647 and 1540 cm<sup>-1</sup> are assigned to the amide I and II from protein (Petrone *et al.*, 2011). These bonds show the presence of complexes of amine phenols ligands in our samples. It is not surprising since the polyphenols of brown algae, namely phlorotannins, is known to have strong bond with protein (Stern *et al.*, 1996).

In order to detect the presence of phlorotannins in our samples – type of polyphenols in brown algae – we passed the commercial standard of phloroglucinol as comparison. Peaks at 3213.73, 1620.31, 1503.87, 1332.46, 1305.86, 1150.82, 1005.31, 996.28, 839.52, 812.39, 799.22, 664.07, 600.28, 578.01 and 517.02 cm<sup>-1</sup> were detected from the standard. There were several peaks detected in our samples corresponding with phloroglucinol, *i.e.* 1619-1617 and 1326-1324. Range of wavelengths at 2940-2925, 1660-1600, 1500-1400, 1100-1000 cm<sup>-1</sup> observed in our samples also showed to be in accordance with the

spectra of gallic acid – commercial standard used to detect the polyphenols – at 1639.70, 1509.58, 140.99, 1153.78, and 1097.08 cm<sup>-1</sup> by Meenakshi *et al.* (2011). Wavelength at 1620 cm<sup>-1</sup> indicated that there was aromatic ring as shown in the spectra of our samples (Lim *et al.*, 2002). Rajauria *et al.* (2013) also reported the same spectra at around 1624, 15505 and 1465 cm<sup>-1</sup> detected in brown algae *Himanthalia elongata* suggesting the presence of aromatic ring. The aromatics groups with C-C stretch absorb at 1500-1400 and 1600-1585 cm<sup>-1</sup>. Aromatic amines with C-N stretch are functional groups absorbing at 1335-1250 cm<sup>-1</sup>. Then the wide peak at around 3260 was assumed to correspond with the detection of carboxylic acid with O-H stretch. In general, the maximum transformation of hydroxyl group is at 3300-3500 cm<sup>-1</sup> region as for the aromatic rings are normally detected between 1400 to 1680 cm<sup>-1</sup> (Rajauria *et al.*, 2013).

Polysaccharides detection in FTIR are mainly characterized by its C-C and C-O stretching bonds in the pyranose ring as well as to C-O-C stretching of the glycosidics bonds between 1200-970 cm<sup>-1</sup> (Gomez-Ordonez and Ruperez, 2011). Other author emphasized to the 1800-650 cm<sup>-1</sup> showing several vibrational modes that are sensitive to the polysaccharides and glycosidic linkages in algae (Shekhar *et al.*, 2012). A strong peak at around 1023 cm<sup>-1</sup> revealed that our samples might be rich in guluronic acid since the 1025 cm<sup>-1</sup> is ascribed to the -OH bending of guluronate (Sakugawa *et al.*, 2004). The peak at 1023 and 890 cm<sup>-1</sup> detected in our samples were the same as those in *Fucus vesiculosus* indicating the presence of alginic acid and fucoidan (Shekhar *et al.*, 2012). Weak peak at 1420 cm<sup>-1</sup> shown in the spectra of *S. muticum* might be due to the symmetric stretch vibration of C-O-C of uronic acid (Singthong *et al.*, 2004).

Biochemical constituents as a representation of algae's nutritional value have been extensively studies in different time, place and species. Accordingly, many studies have reported the health benefits contained in this marine plant and it seems that algae have been long consumed in the Asia. One of its notoriously unique components is polysaccharides proven to have a wide array functions for food, health and beauty. Additionally, other constituents as protein, polyphenols,

ash, lipids and pigments are also surprisingly interesting because they are economically potential as well.

This chapter serves as a preliminary report about the biochemical composition of different *Sargassum* species representing the tropical and temperate region. In addition, the seasonal allows us to comprehend the effect of season towards the biochemical composition of brown algae. Then, the application of FTIR analysis as a secondary tool used in this study can facilitate the qualitative analysis by using directly the dry algal material.

Overall, the biochemical compositions of *Sargassum* in this study are in accordance as the previous study. The difference values of biochemical contents have been previously expected as algae show great plasticity to their environmental variables as their way to adapt and survive. FTIR analysis facilitates the preliminary evaluation of algae biochemical composition since this tool is rapid, easy to use and efficient, as it does not need substantial amount of dry material. By using this method, the seasonal pattern of a species can be observed and it is also possible to determine their correlation with the environmental parameter.

#### Conclusion

Seasons play important role in marine algae biochemical composition. Fluctuated environmental conditions enforce algae to adapt their physiological system in order to survive. Certain biochemical entities have their own season by evaluating its seasonal variation will provide information about the most favorable collection time. For example, for hydrocolloid industries that needs high yield of alginate, summer will be the best time to collect algal biomass, in this case *S. muticum*. For fucoidan, it is best to have algal biomass from spring and summer.

This study informs the correlation of sea surface temperature, air temperature, period of sunlight and precipitation to the production of biochemical composition of *S. muticum*. Some shows linear correlation while other

demonstrates inversed correlation. Besides seasons, life cycle of marine algae has been mentioned as contributing factor to the variation of its biochemical composition, particularly to its reproductive cycle. Such tendency can also be seen in this study. Monthly evaluation of *S. muticum* biochemical composition has shown that life cycle contributes to the biochemical profile of this alga. It may also be concluded that in most marine brown algae from the temperate regions, the seasonal variation of their biochemical compositions have the same pattern without considering the other external factor such as grazing, pollution, etc.

## 3.8.2 Extraction of *Sargassum* polyphenol

## 3.8.2.1 Seasonal variation of S. muticum polyphenol

Polyphenol serves multiple functions for marine brown algae, namely primary and secondary metabolites. As a consequence, the production of polyphenol is determined by an interaction of biotic and abiotic factors (Pavia et al., 1997) and a study related to its variation, in particular the seasonal variation, will give insight into their natural adaptative functions and become basic information for commercial interest. The most favorable collection period of algal material is important to consider. Present study shows that season becomes one of the important factors influencing the production of polyphenols content in S. muticum. Results presented in this study reveal that the polyphenols content of S. muticum achieved its highest quantity in summer and it decreased during winter. Previous study with S. muticum from Point de Diable revealed that its polyphenols content increased between winter and spring 2003, spring and summer 2004 (Plouguerné et al., 2006). Interestingly, same authors also discovered that S. muticum collected from Callot Island contained the highest polyphenols during winter and spring in 2003 and 2004. Authors used aqueous methanol extract of S. muticum. Polyphenols content of brown algae Ascophyllum nodosum also increased during summer, June to September ranging from 0.5-1% of fresh weight (Parys et al., 2009). Mannino et al. (2014) reported the same tendency happened with Cystoseira amentacea. Extract of 96% aqueous ethanol of this species

contained the highest polyphenols content between spring and summer, with 0.17  $\pm$  0.0% and 0.57 $\pm$ 0.1% of dry weight, respectively. *Cystoseira tamariscifolia* exhibited an exponential increase in its polyphenols from February-June and decreased from June to November (Abdala-Diaz *et al.*, 2006). It can be said that this result is in agreement with several cases from previous studies reporting the variation of polyphenols content in brown algae that occurred in different seasons.

Maximum polyphenols content of S. muticum occurring in summer, as presented by this study, has been mentioned to be attributable to the implication of this compounds in the photoprotection of the species (Plouguerné et al., 2006; Le Lann et al., 2012). UV-B radiations provoked the increase of polyphenol content in A. nodosum (Pavia et al., 1997), in Desmarestia spp. (Fairhead et al., 2006) and in C. baccata. In C. tamariscifolia, it was demonstrated that the exudation of phlorotannins from its tissue showed a significant correlation with increasing dosage of irradiation (Abdala-Diaz et al., 2006). Light intensity was not measured in this study. Instead, an observation was made for sunlight also known as daylight period in a day, air temperature, sea surface temperature and precipitation. The correlation analysis has described that all the environmental parameters except for precipitation have given significantly positive effect to polyphenols content. As intertidal species, S. muticum is often exposed to sunlight radiation during the low tide, particularly in summer where the sun intensity is stronger and the temperature is higher than the other seasons. To prevent themselves from oxidative stress, they produce more polyphenols, i.e. phlorotannins. This explains its high content found in this study during summer in all extracts.

In winter, the phlorotannins of *S. muticum* tend to be in its minimum level. It might be explained by their life cycle. The growth period of *S. muticum* occurs between winter and early spring. Thus, it is assumed that there is a trade-off in phlorotannins production of *S. muticum* that is related to its growth. Phlorotannins of brown algae are produced in the physodes, thus it makes them extractable and reactive. Along their life cycle, these physodes exude phlorotannins to algal cell

wall. They are then complexed to alginic acid and deposited at the cell wall. Within this rate, the decrease content of extractable phlorotannins occurs when rates of plant growth, which is cell wall formation, are high. In the opposite, the increase concentration of extractable phlorotannins happens when ever their synthesis exceeds the incorporation into algal cell wall (Arnold and Targett, 2003). As reported in *A. nodosum*, its phlorotannins concentration was high when they experience slow growth rate. Annual shoots of *A. nodosum* having high phlorotannins levels (>9% of dry weight) grew 25% and 54% less than those with a relatively low phlorotannins level (<6% of dry weight) (Pavia *et al.*, 1999). A negative correlation between growth rate and phlorotannins level was reported in *S. patens* as well. *S. patens* that was cultured at 10 °C contained high level of phlorotannins but it had slow growth rate. On the other hand, an individual kept at 30 °C experienced fast growth rate and a decrease in phlorotannins level (Endo *et al.*, 2013). It seems that reproduction cycle of algae contributes indirectly to polyphenol production.

Polyphenol production of marine brown algae shows putative role against the presence of herbivores. This extrinsic factor can significantly induce changes in phlorotannin production of brown algae inhabited nearshore marine communities since it will affect the palatability (Pavia and Brock, 2000). As reviewed by Amsler and Fairhead (2006), snail *Littorina sitkana* avoid the wounded areas of *Fucus gardneri* which showed phlorotannin accumulation, and preferred to feed on undamaged algae with lower phlorotannin contents. Snail *L. obtusata* consumed less *A. nodosum* containing high phlorotannins contents. Turban shell *Turbo cornutus* is marine herbivorous gastropod that prefers eating dead brown algae than live one. This is due to the inhibitory effect of phlorotannins on glycosidase activity, a digestive enzyme found in viscera of *T. cornutus*. Such activity leads to the feed preference of this species (Shibata *et al.*, 2002).

#### Conclusion

Inhabiting the temperate region, *S. muticum* has been mentioned to contain polyphenol that is potential for further application. As most species, *S. muticum* contains high polyphenol in summer. Most favorable organic solvent to extract *S. muticum* polyphenol as proven in this study is using 50% aqueous methanol solvent. Therefore, summer will be the best period to collect *S. muticum* for its polyphenol content.

As secondary metabolites in marine brown algae, polyphenol production is highly influenced by the surrounding conditions. One of conditions that have a significant effect towards the production of polyphenol is season as proven in *S. muticum*. This compound is high during spring-summer with maximum content in summer. It suggests the role of this compound as photoprotective agent in *S. muticum* especially when this species is over-exposed to sunlight during the lowest tide. The presence of long interconnected aromatic ring in brown algal polyphenol is the defensive agent mostly responsible generating protection against oxidative stress in brown algae.

Having strong antioxidant activity, polyphenol of brown algae might become an important source of antioxidative agent. As a consequence, polyphenol compounds from marine brown algal have been extensively explored and studied since it is valuable for commercial interest. For commercial purpose, it is crucial to acknowledge the most optimum period of polyphenol production. This study provides the required information. In addition, different type of solvents applied in this study helps to acknowledge best-fitted solvent to extract polyphenol.

# 3.8.2.2 Solid-liquid extraction of *Sargassum* polyphenol

Many studies have attempted to optimize the extraction process by applying different types of solvent, temperature, extraction time, drying process and many more. SLE has become the most used method in extracting polyphenol (Michalak and Chojnacka, 2015). This method mostly relies on the polarity of the solvents used due to the chemical nature of target compounds (Bart, 2011). In general, the

process in SLE method can be categorized in three parts; (1) changing phase of the solute, in this case the target compounds, as it dissolve in the solvent, (2) its diffusion through the solvent in the pores of the solid to the outside of the particles, and (3) the transfer of the solute from the solution in contact with the particles to the main bulk of the solution (Richardson et al., 2002a). The more dispersed the solute in the solid material, the more difficult the extraction will be and the extraction rate will fall. This is because the solvent will have to penetrate further into the layer of solid material in order to reach the solute (Richardson et al., 2002a). Furthermore, the efficiency of SLE highly depends on the particle size, temperature, and agitation during the process (Richardson et al., 2002b). In the extraction of phenolic compounds, the storage time of dry materials and conditions, as well as the presence of interfering substances are other important factors to consider, influencing the extraction efficiency (Naczk and Shahidi, 2004). In addition, the polarity of target compound becomes the most important factor in choosing the solvent (Azmir et al., 2013). Methanol, ethanol, acetone, ethyl acetate, hexane (Koivikko et al., 2007; Kim et al., 2011; Girija et al., 2013; Sathya et al., 2013; Le Lann et al., 2016) are some examples of organic solvents mostly used for phlorotannins extraction.

Ethanol extract of *S. filipendula, S. duplicatum, S. crassifolium, S. binderi,* and *Padina* sp. showed to contain higher polyphenol with 12.9, 7.9, 6.8, 9.1 and 10.2 mg g<sup>-1</sup> of extracts than ethyl acetate and hexane extracts. Among ethanol and ethyl acetate, hexane extracted less polyphenol in *S. filipendula, S. duplicatum, S. crassifolium, S. binderi,* and *Padina* sp. (Bambang *et al.*, 2013). Methanol extracts of *S. hystrix, S. siliquastrum,* and *S. mcclurei* had approximately 20, 17 and 15 g 100 g<sup>-1</sup> extract of polyphenol. *S. polyceratium, S. angustifolium* and *S. filipendula* extracted with methanol showed less polyphenol content, ranging from 5 to 10 g 100 g<sup>-1</sup> of extract, than the three previous species (Budhiyanti *et al.*, 2012). By applying different temperature and extraction time for optimizing the extraction of polyphenols in *S. muticum,* Tanniou (2014) also revealed that 40 °C and 3 h of extraction resulted better content of polyphenols. Further, polyphenol extraction

of *S. muticum* was more effective with 50% aqueous ethyl acetate than the mixture of 88% hexane- 12% ethanol and 75% of aqueous ethanol as it contained the highest polyphenol content, 18.6% of dry weight. Meanwhile, the lowest polyphenol content in *S. muticum* was obtained from the hexane-ethanol mixture with only 3.7% of dry weight (Tanniou *et al.*, 2013a). In this study, *S. muticum* extracted in methanol 50% had higher polyphenol content than water, methanol and ethanol 75%, same case with *S. polycystum*. On the other hand, *S. aquifolium* and *S. ilicifolium* contained the highest polyphenol content in methanol extract. These findings have proven that type of solvents determine the efficiency of polyphenol extraction. In addition, temperature also play critical role during the extraction of polyphenol. Bambang *et al.* (2013) selected room temperature, meanwhile Budhiyanti *et al.* (2012) and Tanniou *et al.* (2013a) used 40 °C as their extraction condition, same with our study.

Other study highlighted in their findings that 40 °C, 180 min, and 40 mL g <sup>1</sup> of solid-liquid ratio were the most efficient conditions for the extraction of polyphenols in Sargassum sp. (Ismail and Abdullah, 2015). Polyphenol compounds are most soluble in less polar solvents than water (Koivikko et al., 2005). Most recommended solvent is a mixture of water with methanol, ethanol or acetone, for example, brown algae Fucus vesiculosus have been commonly extracted with aqueous mixtures of ethanol and acetone (Ragan and Glombitza, 1986; Waterman and Mole, 1994; Koivikko et al., 2005). Lopez et al. (2011) explained that solvent polarity had no drastic effect to the total amounts of a phenolic groups, but more to the phenolic profile. Regardless of the amount, since this study has proven the contrary, their theory about difference in phenolic profile due to solvent polarity is in agreement with recent result. As can be seen from the chromatogram data (Appendix 8), the spectra of methanol, methanol 50%, water and ethanol 75% extracts are not the same. Water and methanol 50% extracts displayed the most distinguished profile of peaks between the other two solvents.

Most of the temperature suggested for the extraction of terrestrial and marine vegetal polyphenols compound was below 60 °C (Le Lann *et al.*, 2008; Dai and Mumper, 2010; Yoon *et al.*, 2010; Tanniou *et al.*, 2013a; Andrade *et al.*, 2015; Ismail and Abdullah, 2015). Organic solvents such as methanol, acetone, and many others tend to evaporate at high temperatures (approximately > 50°C), thus their efficiency decreases due to the shifting of solvents-to-water ratio leading to the change of sample-to-solvent ratio (Tan *et al.*, 2013). As reported by Dai & Mumper (2010), besides temperature, the sample-to-solvent ratio is another one of the crucial factors that influences the yield of chemical extraction.

Nevertheless, there was several studies that tested the extraction of polyphenols compound using temperature above 60 °C depending in the method and type of solvents applied for the extraction (Siriwardhana *et al.*, 2008; Park *et al.*, 2012; Shitanda *et al.*, 2014). For marine algae, their maceration into a high level of heat might accelerate the extraction process as the heating soften the tissues and facilitate the depolymerisation of polysaccharides (Siriwardhana *et al.*, 2008), phenol-protein, and phenol-polysaccharides linkages (Tan *et al.*, 2013). As proven by Siriwardhana *et al.* (2008), brown algae *Hizikia fusiformis* extracted in water at 100 °C for 60 minutes yielded 4.65 % of polyphenols content while at 25, 50 and 75 °C had only 1.40, 2.02 and 2.74 % of polyphenol, respectively.

#### Conclusion

Various factors have given significant effect to the extraction of polyphenol in marine brown algae; two of them are type of solvents and temperature. In this study, extraction yield, polyphenol content and chromatographic profiling of *Sargassum* species exhibit variation due to type of solvents and species. Some studies have reported that chemical profile of brown algal polyphenol in one individual is various depend on the polarity of solvents. Interest of this study is to acknowledge what is the most optimum solvent to extract polyphenol in different species of *Sargassum* and. It seems that each species displays its own preference of solvent that might be related to the polarity of polyphenol produced in *Sargassum*.

# 3.8.2.3 Enzyme-assisted extraction of *S. muticum* polyphenol

In marine algae, the main drawback during the extraction of phenolic compounds occurs due to the presence of complexes polysaccharides as the main component of algal cell wall. Brown algal phlorotannins are known to be incorporated in its cell wall (Schoenwaelder and Clayton, 1999) covalently bonded to the polysaccharides (Koivikko, 2008) and proteins (Stern *et al.*, 1996). Therefore, it requires a strong condition to degrade these bonds and extract the phlorotannins. The SLE method does not meet this requirement since it does not degrade such complex bonds since it works based on the solubility of target compound in to the solvents (Richardson *et al.*, 2002a).

One part of this study was to evaluate the extraction yield and the polyphenol content *S. muticum* digested by various commercial enzymes. Enzyme-assisted extraction has been relatively well-known for the past few years as an alternative method to extract the bioactive compound in marine algae (Heo *et al.*, 2003a, 2003b; Ahn *et al.*, 2004a; Siriwardhana *et al.*, 2004; Park *et al.*, 2005; Athukorala *et al.*, 2006a, 2006b; Je *et al.*, 2009; Charoensiddhi *et al.*, 2014; Sanchez-Camargo *et al.*, 2015). This method has been mostly used to recover bioactive compounds, such as polyphenol that is highly potential as antioxidant (Heo *et al.*, 2003b; Ahn *et al.*, 2004a; Sanchez-Camargo *et al.*, 2015), or in certain case, polysaccharides for its anticoagulant and protein for its antihypertensive effects (Sato *et al.*, 2002).

In order to optimize the extraction yield, optimal conditions of pH and temperature have been chosen for each selected enzymes. Carbohydrase and protease, two main types of enzymes mostly applied in bioactive extraction of marine brown algae, work optimally at pH ranging from 4.5-8.0 and temperature at 40-60 °C (Athukorala *et al.*, 2006a; Zou *et al.*, 2008; Je *et al.*, 2009; Charoensiddhi *et al.*, 2014). To achieve optimum degree of hydrolysis, in other word yield of extraction, some studies established 12 h of extraction (Heo *et al.*, 2003a, 2003b; Athukorala *et al.*, 2006b; Je *et al.*, 2009) and the other study chose

a one-day extraction (24 h) (Charoensiddhi *et al.*, 2014). As an example, AMG (carbohydrase) and Alcalase (protease) extract of *Ecklonia cava* showed higher extraction yield, 41.5% and 42.7%, than the other carbohydrase (Viscozyme, Celluclast, Termamyl and Ultraflo) and protease (Protamex, Kojizyme, Neutrase, Flavourzyme) in 12 h of extraction (Heo *et al.*, 2003a). Still from the same author, AMG was also shown to yield higher degree of hydrolysis than other carbohydrase in *S. fulvellum* (27.9%), *S. coreanum* (33.2%), and *S thunbergii* (33.9%). Then, Heo et al (2003a) reported that Alcalase yielded high degree of hydrolysis in *Ishige okamurae*, *S. horneri*, *S. coreanum*, *S. thunbergii* and *Scytosiphon momentaria*, with 29.9%, 40.0%, 40.7%, 30.5% and 30.9%, respectively. *E. radiate* extracted with carbohydrase (Ultraflo) and protease (Alcalase and Flavourzyme) for 24 h increased the extraction yield by 5-20% compared to aqueous extraction. Almost 70% of algal dry material was achieved when introducing Ultraflo, Alcalase and Flavourzyme to the extraction of *E. radiata* (Charoensiddhi *et al.*, 2014).

Eight commercial enzymes consisted of two proteases (Neutrase and Alcalase) and six carbohydrases (Ultraflo, Celluclast, AMG, Termamyl, Shearzyme and Viscozyme) used in this study were proven to be effective in increasing the yield of extraction in comparison with water extraction, except for Shearzyme. Highest yield was obtained from *S. muticum* digested with Neutrase, one of proteolytic enzymes, though the value was not significantly different from the other enzymes and water extraction. Enzymes from the protease group have been reported from several studies as an effective enzyme in enhancing the extraction yield in algae (Heo *et al.*, 2003a; Wang *et al.*, 2010; Hardouin *et al.*, 2013). Opposite result was obtained from previous study. Extraction of *S. muticum* using Alcalase for 2 and 4 h yielded 13.6 and 17.8% of dry weight. Viscozyme extract of *S. muticum* showed higher extraction yield than Alcalase extract. For 2 h extraction, the extraction yield was 20.6% of dry weight and 4 h extraction resulted 23.5% of dry weight (Sanchez-Camargo *et al.*, 2015). In present study, 3 h extraction of *S. muticum* yielded 28.7%, this value was slightly

higher than the previous study. Such contradictive results might be due to different concentration of enzyme used in the extraction since proportion of substrate to enzyme is one of the most considerable variables (Wijesinghe and Jeon, 2012). Previous study used 0.1% of enzyme while in present study added 2.5% of enzyme in algal solution. It is assumed that the enzyme concentration is too low, as a consequence, the degradation process is not optimum since the enzyme reaches its limiting concentration but there are still plenty of substrate available to be bound by the enzyme. When the substrate concentration is high, the addition of enzyme can enhance the rate of reaction, till the substrate concentration becomes limiting (Sowbhagya and Chitra, 2010). Suitable amount of enzyme addition is important to consider for achieving the limiting point of substrate for optimum result. In general, our finding regarding the extraction yield of brown algae is in accordance with certain previous studies even though in particular case we had the opposite result as mentioned above.

Beside extraction yield, polyphenol content of *S. muticum* was also enhanced by enzyme-assisted extraction compared to the aqueous extraction. The highest polyphenol content was observed in Viscozyme extract of *S. muticum* containing 6% dry algal material of polyphenol. *S. coreanum* extracted with Protamex contained 9.9 mg g<sup>-1</sup> dry algal material of polyphenols content, but with Viscozyme, its polyphenols content was higher with 11 mg g<sup>-1</sup> dry algal material (Heo *et al.*, 2003a). Study by Sanchez-Camargo also obtained higher polyphenol content in Viscozyme extracts of *S. muticum* from 2 and 4 h of extraction (10.3 and 10 mg g<sup>-1</sup> dry algal material) than the Alcalase extract (7.91 and 6.1 mg g<sup>-1</sup> dry algal material).

The efficiency of Viscozyme in extracting polyphenol content from the brown algae works differently in every species. Siriwardhana *et al.* (2004) studied the effect of different commercial enzymes in antioxidant activity of *H. fusiformis*. Based on their result, Alcalase and Neutrase showed the highest content of polyphenol compound. Alcalase also enhanced the polyphenols content of *S. thunbergii*, *S. horneri*, *S. fullvelum* and *Ishige okamurae* (Heo *et al.*, 2005).

This study reveals that the polyphenols content of S. muticum extracted with carbohydrases was higher compared to the proteases. It leads to an assumption that the carbohydrases disrupt algal cell wall and release the metabolites linked to the polysaccharides, in this case the polyphenols compounds. It is also possible that the polyphenols compounds of brown algae are mostly attached to the alginates instead of proteins. As mentioned by previous studies, the polyphenols compounds are not only covalently attached to the proteins but also to the alginates (Bitton et al., 2006; Salgado et al., 2009; Deniaud-Bouet et al., 2014). The enzymatic degradation of alginate fractions always release a significant amount of polyphenol since polyphenols substances form high-molecular weight complexes with alginates due to oxidation with haloperoxidases (Bitton et al., 2006; Salgado et al., 2009). This type of linkage serves as structural component constituting the brown algal cell wall. Polyphenols cross-linking of alginate may be an integral part of initial wall formation as shown in the Fucus zygote. The linkages between polyphenol compounds and algal cell wall polysaccharides occurs immediately after the secretion of polyphenols substances into the cell wall, complexing finally with alginic acid (Arnold and Targett, 2003; Koivikko *et al.*, 2005).

From the results of the evaluation of biological activities presented previously, it is apparent that the eight enzymatic extracts of *S. muticum* showed different levels of potential. It indicates that the extracted polyphenols compound might function differently in responding to the test applied. In addition, it is also important to consider the possibility of an intervention from other compound as in this study the enzymatic extracts tested were all crude material.

#### Conclusion

Based on the results presented in this study, enzyme-assisted extraction enhances the extraction yield and polyphenol content to certain extent compared to the aqueous extracts in *S. muticum*. Some studies applied 12 h of extraction in order to optimize the polyphenol extraction, but here only by applying 3 h of extraction;

we can see an increase of polyphenol content to 2 fold than the aqueous extraction.

## 3.8.2.4 Optimization of Sargassum polyphenol enzyme-assisted extraction

Based on the screening of *S. muticum* polyphenol extraction in different commercial enzymes, Viscozyme and Termamyl extracts came out with their high polyphenol content. As a result, these two enzymes were selected for the next step, which was the optimization part to evaluate the efficiency of these enzymes in extracting polyphenols with adjusted conditions. In addition, Protamex was also used. As mentioned by many previous studies, 40 °C has been the most favourable temperature to extract polyphenols, thus, this temperature was selected.

It seems that 40°C is not favourable for polyphenol extraction of S. muticum, S. ilicifolium, and S. polycystum using Viscozyme. In the contrary, Protamex extract of these species showed better polyphenol content ranging from 3 to 6% of dry algal material. Such result is not too surprising as 40°C is the most optimum condition for Protamex. Perhaps, in this soft extraction condition, Protamex manages to break protein-phlorotannin linkage of algal cell wall and release more polyphenols. Other possibility is, instead of polyphenol compounds, the protein compound extracted by this proteolytic enzyme is more likely reacted with Folin-Ciocalteu reagent during the determination of polyphenol content. This reagent does not merely measure polyphenol since this it measures the total reducing capacity of a sample. It can also react to any reducing substances such as thiols, nitrogen-containing compound and some in organic ions (Ikawa et al., 2003; Everette et al., 2010). It might explain partially regarding the high polyphenol content in Protamex extracts of S. muticum, S. ilicifolium and S. polycystum, that there was interference from other compounds, especially when the extracts were not purified. However, it does not exclude the presence of polyphenol, *i.e.* phlorotannins, in those enzymatic extracts.

As previously mention (see 4.2.2), some studies applied high temperature to extract polyphenol in brown algae. An elevated temperature helps the solvents to reach the samples matrices as the viscosity and the surface tension of the solvents are decreased. In addition, it facilitates the solubility of analytes and promotes higher mass transfer rate (Dai and Mumper, 2010). Using high temperature will be mostly compatible with water as solvent. Furthermore, in relation to algae, the introduction of the enzyme followed by heat treatment can further hydrolyze proteins and laminarine (Siriwardhana *et al.*, 2008). As a consequence, this process results in liberating more bioactive compounds to the solvents. However, due to the instability of polyphenol compounds as the elevated temperature stimulate a degradative reaction which is the oxidations of compound resulting in a decrease of polyphenols yields (Akowuah and Zhari, 2010; Dai and Mumper, 2010). As a result, high temperature is not really applicable to polyphenol extraction.

#### **Conclusion**

It is unfortunate because polyphenol extraction of *Sargassum* using enzymeassisted extraction at 40°C does not enhance the polyphenol content compared to the previous screening at 50°C. Even though, the Protamex extract of *Sargassum* indeed shows higher polyphenol content than the Viscozyme extracts. It is not surprising as 40°C is the optimum temperature for Protamex.

# 3.8.2.5 Purification and characterization of *S. muticum* polyphenol enzymatic extract

FT-IR also detects other peaks that appear to be correlated with polyphenol bonds characterized by its aromatic and hydroxyl groups with the detection of C-H aromatics, C-O-C vibration, C-O stretch, and C=C. However, a protein bond is also exposed through this analysis by the presence of amides bond either in the raw or semi-purified enzymatic extracts. In contrary, the purification method has successfully removed the cellulose and mannuronic acid. Later, the HPLC

analysis has also detected interesting peak that is assumed to be the phloroglucinol-like compound.

Due to the lack of semi-purified extracts, it was impossible to characterize them using mass-spectrometry. This will be performed in future work to help to identify the group of phlorotannins existed in *S. muticum* extracted with enzyme.

## 3.8.3 Evaluation of biological activities of Sargassum polyphenol extracts

#### 3.8.3.1 Antioxidant activities

Polyphenols compounds could be monomeric, oligomeric or polymeric with an aromatic ring having one or more hydroxyl substituents (Balboa *et al.*, 2013). These compounds have been known as a promising and strong antioxidant agent since they have up to eight interconnected rings of hydroxyl groups (Wang *et al.*, 2009). The polyphenols group is known as an excellent hydrogen donor forming hydrogen bonds with the carboxyl group of protein (Ozdal *et al.*, 2013). Later the existence of these hydroxyl groups provides the molecule with the ability of donating a proton to radical leading to its role as a possible chain breaking molecule or antioxidant upon secondary oxidation (Franco *et al.*, 2008).

Therefore, the antioxidant activity of brown algae is strongly related to their polyphenols content. As in this study, *S. muticum* digested with Viscozyme had strongest antiradical activity indicated by the low value of IC<sub>50</sub>. Reducing the particle size can increase the extraction rate and the yield of polyphenols compounds that can be achieved by grinding and enzyme-assisted extraction (Franco *et al.*, 2008).

S. horneri digested with AMG, Celluclast, Termamyl, Ultraflo and Viscozyme for 12 h of extraction was reported to show DPPH radical scavenging activities at 25 μg mL<sup>-1</sup> (Park *et al.*, 2004, 2005). Heo *et al.* (2005) also studied the antioxidant activity via DPPH assay of S. horneri enzymatic extracts. However, their results showed that the Viscozyme, Celluclast, AMG and Ultraflo extracts with final concentration 2 mg mL<sup>-1</sup> did not perform any inhibition.

Instead, the Termamyl extract of *S. horneri* showed 11.2% of inhibition. This activity was poor compared to the commercial standard BHA and BHT with 87% and 56% of inhibition, respectively. Same authors also reported that the antiradical activity of *S. fullvelum*, *S. coreanum*, and *S. thunbergii* extracted with carbohydrase also showed weak activity ranging from 4-38% compared to the standards. It might imply that there is an inter-individual and interspecies variation affecting the performance of antioxidant activity.

Antioxidant effects of S. muticum enzymatic extracts might also be due to the presence of polysaccharides and protein. Sugar content in S. muticum enzymatic extracts in this study ranged from 6 to 10% of dry algal material as shown by the biochemical analysis. Moreover, there was approximately 7% of sulfate groups and 0.6% of uronic acid content in S. muticum enzymatic extracts. It is possible that mostly the sulfate groups and alginic acid provoke the antioxidative effect of S. muticum in this study since it is the sulfated polysaccharides in brown algae known to possess wide range of pharmacological properties. Heterofucans of S. filipendula had been reported to show antioxidant activity such as hydroxyl and superoxide radical scavenging, chelating effects on ferrous ions and reducing powers in active concentration at 2 mg mL<sup>-1</sup> (Costa et al., 2011). Lim et al. (2014) in their study concerning the antioxidant effects of S. binderi also reported that the antiradical effect of this sample was at 2 mg mL<sup>-1</sup>. It indicated that their antioxidant activity was weaker in comparison to the commercial standard, BHA, BHT and ascorbic acid with 6.0 x 10<sup>-4</sup>, 9.6 x 10<sup>-4</sup> and 3.5 x 10<sup>-4</sup> mg mL<sup>-1</sup>. However, it had performed high superoxide (compared to commercial food-grade fucoidan, BHA and BHT) and hydroxycal radical scavenging activities (compared to BHT) (Lim et al., 2014). The ability of S. graminifolium polysaccharides to inhibit 50% of superoxide anions scavenging and antiradical activities was at 1.9 and 0.6 mg mL<sup>-1</sup>, respectively (Zhang et al., 2012).

## 3.8.3.2 Tyrosinase and elastase inhibition activities

Skin aging causes many changes, including thinning, dryness, laxity, fragility, enlarged pores, skin pigmentation, fine lines and wrinkles (Wang *et al.*, 2015). These alterations in skin structure result from pathomechanism such as enzymatic degradation and disintegration of dermal elastin fibers, which maintain resilience of the skin; the proteolytic breakdown and disorganization of dermal collagen networks which provide the skin with strength and resistance; and the continuous exposure of free radicals against cellular compartments of the skin especially mitochondrial DNA (mtDNA) and excessive synthesis of melanin (Jablonski and Chaplin, 2010; Makrantonaki and Zouboulis, 2010; Dupont *et al.*, 2013).

Tyrosinase is a metallo-enzyme oxidase catalyzing two distinct reactions of melanin synthesis. Melanin helps to protect skin from the damaging ultraviolet radiation of the sun. However, elevated concentrations of melanin in the skin can result in hypigmentation such as freckles and moles (Kang et al., 2012). While tyrosinase is responsible for skin pigmentation, elastase is a proteinase enzyme capable of degrading elastin. Elastin is an extracellular matrix protein providing elasticity to the connective tissues. It forms elastic fiber in skin dermis having an influence on skin elasticity (Kim et al., 2009; Moon et al., 2010). Therefore, inhibition of tyrosinase and elastase activity become a method to protect against skin aging (Kim et al., 2009; Chang, 2012). Many have concerned and interested to the promising potential of marine algal bioactive compound in cosmeceutical industries related to their tyrosinase and elastase inhibition activities. Hexane fraction of S. polycystum contained potential skin-whitening agents that could inhibit the melanoma cells in dose-dependent way. The higher concentration of fraction, the stronger activity will be (Chan al., 2011). etDichloromethane:methanol extract of Saccorhiza polyschides, Cystoseira humilis, Bifurcaria bifurcatan C. ericoides and Laminaria ochroleuca exhibited an important elastase inhibition activity greater than 70% of inhibition for 100 µg (Oumaskour et al., 2016). In the case of phlorotannins, many studied have reported their potential in inhibiting the tyrosinase activity. Dieckol, isolated

phlorotannins from the brown alga E. cava has been examined for its inhibitory effects on mushroom tyrosinase. Dieckol was capable to inhibit the mushroom tyrosinase with IC<sub>50</sub> 20 µM and it showed no cytotoxicity (Kang et al., 2012). 7phloroekol of E. cava had been observed to have an inhibitory effect on mushroom tyrosinase at 50 µg/ml. This compound was isolated from the ethanol extract of E. cava (Yoon et al., 2009). These examples provide knowledge concerning the tyrosinase inhibition activity from marine brown algal phlorotannins. Some results presented in present study are in accordance with previous studies as previously mentioned. Polyphenol extracts of S. muticum extracted in different commercial enzymes showed interesting result regarding the tyrosinase and elastase inhibition activities. Attention was given to the tyrosinase inhibition activity; S. muticum Neutrase extract demonstrated the highest activity at 41% compare to other enzymes. Later, Protamex extract of S. ilicifolium and S. polycystum also showed the potential to inhibit the tyrosinase, slightly better than, the Viscozyme extract. To our best knowledge, there are not any studies yet that evaluates the tyrosinase and elastase inhibition activity of brown algal polyphenol extracted with commercial enzymes. This finding could serve as basic information regarding the potential of polyphenol enzymatic extract from brown algae in cosmeceutical industries.

## 3.8.3.3 Antimicrobial activities

As reviewed by Singh (2013), phloroglucinol, eckol, phlorofucofuroeckol (deacetylated 8), dieckol, and 8,8"-bieckol have been tested for antibacterial activity against pathogenic gram positive and negative, such as *Escherichia coli*, *Staphylococcus aureus*, *Streptococcus pyogenes*, *Vibrio parahaemolyticus*, *Propionibacterium acnes*, and many more. All other compounds, besides phloroglucinol, have shown bactericidal activity. Bactericidal effect tends to increase with the degree of polymerization of phloroglucinol. Further, phlorotannins interaction with bacterial proteins is assumed to play a role in the bactericidal action (Nagayama *et al.*, 2002; Shibata *et al.*, 2002). In this study, polyphenol extracts of *Sargassum* extracted using SLE showed positive activity

against *E. coli*, *B. subtilis* and *P. aeruginosa*. Unfortunately, the enzymatic extracts of *Sargassum* polyphenol did not show expected results as they did not demonstrate any antibacterial activity. It is suspected that the presence of sugar could contribute to the growth of bacteria in the media. However, this assumption will need further verification by evaluating the purified enzymatic extract of *Sargassum* polyphenol.

Even though the enzymatic extract of Sargassum polyphenol did not show antibacterial activity, biofilm inhibition activity demonstrated the opposite result. It seems that polyphenol of Sargassum enzymatic extracts was more active against biofilm formation. This contradictive result is caused by different mechanism of inhibition. In antibacterial activity, the mechanisms of inhibition are most likely thought to rely on inhibition of extracellular microbial enzymes, deprivation of substrates required for microbial growth, or direct inhibition of oxidative phosphorylation (Scalbert, 1991). Regarding biofilm inhibition activity, two main mechanisms have been proposed, i.e. direct inhibition of bacterial settlement or survival of the macrofoulers larvae and regulation of macrofoulers growth, which in turn affects larval settlement of the macrofoulers (Avelin Mary et al., 1993). Some studies have reported the promising potential of brown algal polyphenol compound as antifouling agent. Water extract of S. vulgare polyphenol from its pneumatocysts presented 75 to 99% inhibition of mussel Perna perna attachment. Polyphenol from leaflets of S. vulgare had 67 to 80% inhibition of byssal adhesion (Plouguerné et al., 2012). Phlorotannin extract of S. tenerrimum had EC<sub>50</sub> of larval settlement inhibition of Hydroides elegans at 0.526 ppm. It indicated that this extract was effective in inhibiting the settlement of H. elegans larval attachement (Lau and Qian, 1997). Phlorotannins have been reported as rapidly induced compounds toward grazing. One of the studies reporting such result is an induction phlorotannins content in Stypopodium zonale (Ank et al., 2013). Authors observed that there was an increase of phlorotannins content in S. zonale submitted to stimulate herbivory for 2 days compared with uninjured algae. However, after 4 days of exposure to herbivory, drastic decline of phlorotannins content was remarked. A possible explanation made by authors that phlorotannins induction is likely related to the healing process of the wound, and not an anti-herbivore response.

Antiviral effects of brown algae are usually shown by its polysaccharides as explained by previous studies with active concentration from  $0.5 - > 500 \,\mu\text{g/ml}$ (Ponce et al., 2003; Lee et al., 2004, 2011; Ahn et al., 2006; Queiroz et al., 2008; Sinha et al., 2010; Elizondo-Gonzalez et al., 2012). The efficiencies of polysaccharides against virus are based on the following mode of actions (1) inhibition of viral reverse transcriptase; (2) the interference with viral adsorption; and (3) enhancement of immune function of host cells (Xianliang et al., 2000; Xin et al., 2000; Zhu et al., 2004; Talarico et al., 2005; Queiroz et al., 2008). However, for the past few years, several studies concerning the antiviral effects of brown algae phlorotannins have appeared. Ahn et al. (2004b) reported an inhibitory effect of 8,8'-bieckol and 8-4"-dieckol isolated from E. cava on HIV-1 reverse transcriptase and protease in vitro. Phloroglucinol, eckol, 7-phloroeckol, phlorofucoroeckol and dieckol isolated from E. cava were effective against porcine epidemic diarrhea coronavirus at concentration ranging from 10.8 – 14.6 μM (Wang et al., 2008). Ethyl acetate fractions of phlorotannins from brown algae Eisinia bicyclis have been reported to show antiviral effects against genital human papilloma virus (HPV) at 50 µg/ml (Kim and Kwak, 2015). Dichloromethane: methanol extract of S. cymosum, S. polyceratium, S. vulgare, Padina gymnospora, Lobophore variegate, Dictyopteris delicatula, Sictyota menstrualis and Stypodium zonale demonstrated herpes simplex virus-1 (HSV-1) inhibition activity with percentage of inhibition ranging from 20 to 96% at 400 µg/ml. Spectroscopy analysis of these crude extracts detected the presence of polyphenol compounds along with other active compound like halogenated sesquiterpenes obtusol, elatol, meroditerpenoid atomaric acid, triacylglycerols and fatty acids (Soares et al., 2012).

Result in this study regarding to the antiviral effect of *S. muticum* enzymatic extracts seems to be in the contrary to the previous studies. Out of

eight, there were only two enzymatic extracts that were quite effective against the HSV-1; they were *S. muticum samples* digested with Neutrase and Alcalase. However, the antiviral effects of these two enzymatic extracts might not be provoked only by the phlorotannins. It is possible that the presences of other compounds such as protein and polysaccharides that are present in enzymatic extracts are involved in disrupting the viral cells. Based on the colorimetric analysis, the biochemical composition of *S. muticum* enzymatic extracts contained 30-55% of protein and 6-12% sugar.

#### Conclusion

Polyphenol extraction of *Sargassum* indeed demonstrates interesting antioxidant activities as reported by many studies. In addition, this study also shows diverse biological activities performed by *Sargassum* polyphenol extracts such as tyrosinase, and biofilm inhibition activities. In the case of *S. muticum*, polyphenol extracts shows quite encouraging result for its elastase inhibition and antiviral activities with no cytotoxicities. However, enzymatic extract of *Sargassum* polyphenol does not show any positive antibacterial activity. Despite of the fact that polyphenol is quite remarkable for its antibacterial activity. This is possible that the sugar content in enzymatic extract of *Sargassum* polyphenol contributes to bacterial growth by providing more nutrients.

Various results obtained from this study imply the vast array of polyphenol potential for commercial interest from *Sargassum* species. Moreover, the bioactivities of polyphenol from this species display between species variation. It is important to consider type of species that has more active polyphenol and method used to extract these compounds. Enzyme-assisted extraction of *Sargassum* polyphenol surely shows remarkable bioactivity in certain extent in comparison to the solid-liquid extraction. One example, reducing power activity of *S. aquifolium* polyphenol in Viscozyme extract is better than the methanol, ethanol or even aqueous extracts. Certain results in this study have proven the efficiency of enzyme-assisted extraction in improving the bioactivities of bioactive compounds, especially polyphenols from marine brown algae.

## 3.8.3.4 Polyphenol content and bioactivities of purified fraction

Semi-purification fraction of S. muticum Viscozyme had yielded an important percentage of polyphenols compounds assuming to be phlorotannins. Antiradical and reducing power capacity of the semi-purified fraction were improved compared to the raw enzymatic extracts. Later, the antiviral activity of polyphenol seems to be enhanced as well with no cytotoxicity remarked. This result suggests that the antioxidant activity showed by S. muticum enzymatic extracts is generated by the presence of polyphenols compound. It is supported by FT-IR and HPLC analysis that detects the presence of peaks corresponding to the aromatic rings of phloroglucinol. Chromatography has been used in some previous studies as a procedure to purify phlorotannins in brown algae. Fucophlorethols, fuhalols from S. spinuligerum had been successfully isolated by means of flash chromatography and HPLC followed by <sup>1</sup>H NMR for further identification (Glombitza and Keusgen, 1995; Glombitza et al., 1997). Phlorethol fucodiphlorethol, and Fucophlorethols from E. cava was isolated using a flash chromatography (Ham et al., 2007; Li et al., 2011). Column chromatography and RP HPLC were applied to isolate the dieckol and phlorofucofuroeckol-a from E. cava (Choi et al., 2014).

#### Conclusion

This part of study might only succeed to perform semi-purification step, but the results have at least verify the role of polyphenol-rich (phlorotannins) fraction of *S. muticum* as active compound. Bioactivity of this fraction is not limited to antioxidant but also to antiviral activity. It also gives no toxic effect to the tested cell. Preliminary characterization has managed to identify the presence of aromatic groups in purified fraction and the purification procedure has eliminated the presence the polysaccharides.

Working with purified polyphenol fraction of *S. muticum* helps to further comprehend the bioactivity efficiency of this compound. Selection of appropriate method is crucial to obtain sufficient fraction for various assays in order to verify the bioactivity of target compound. Moreover, by purifying the polyphenol of *S.* 

*muticum* will assure that the bioactivity performed indeed come from the polyphenol without any interference from other contaminant compounds such as polysaccharides, proteins, or lipid.

## 3.8.4 Characterization of Indonesian *Sargassum* species using Fourier-Transform Infrared Spectroscopy

Based on early detection of the spectroscopy analysis, several spectra assumed to correspondent with phloroglucinols were detected. The characteristic of phenolic compounds lies on its aromatic ring (1260, 1620, 2970, and 2850-2970 cm<sup>-1</sup>) and hydroxyl group (3415 cm<sup>-1</sup>) (Lim et al., 2002; Ganapathi et al., 2013). Besides phenolic compounds, the detected spectra showed that there was also the presence of other compounds. Broad spectrums between 3500-3000 cm<sup>-1</sup> usually relate to the presence of polysaccharides and proteins (Gomez-Ordonez and Ruperez, 2011; Kannan, 2014). Spectra ranging from 1200-970 cm<sup>-1</sup> assumed to be the bonds of C-C and C-O pyranose ring and C-O-C stretching of glycosidic bonds (Gomez-Ordonez and Ruperez, 2011). Medium peak at 1636 cm<sup>-1</sup> and 1428 cm<sup>-1</sup> as shown at Figure 1 corresponded to the carbonyl (C=O) and carboxyl (COOH) that might indicate the presence of alginic acid (Ganapathi et al., 2013). Peak signal observed around 648 cm<sup>-1</sup> might be due to C-Br bond and near 600-670 cm<sup>-1</sup> <sup>1</sup> due to C-S and C=S stretching vibrations (sulfides) (Girija et al., 2013; Kannan, 2014). As the enzymatic extracts tested in this study were crude extracts, then the presence of other molecules besides phenolic compounds had been expected. Therefore, in polyphenol extract of Indonesian Sargassum, polysaccharides and polyphenols in the same time.

### Conclusion

FT-IR analysis definitely helps in identifying the biochemical entities of algal material through the detection of its functional groups. Enzymatic extract of Indonesian *Sargassum* polyphenol, polysaccharides and proteins are two mostly identified compounds as expected. As enzymes degrade algal cell wall, they release any bioactive compounds that are present in the matrix of algal cell wall.

Without any purification of designated compounds, interferences from other compounds than polyphenol in bioactivity assays must not be excluded.

## **Chapter 4 Conclusions**

Marine brown algal have attracted many attention as they show infinite applications with valuable economic value due to their complexes and distinctive polysaccharides and bioactive compounds. In addition, brown algae have interesting nutritional profiles that are beneficial for food and fertilizer industries. Before going further to commercial exploitation, it is highly suggested to apprehend thoroughly about the variation of algal biochemical compositions and bioactive compositions, in this case, polyphenol. Determination of the best period to collect algal biomass having the highest content of biochemical compositions for certain application is important. Seasonal study of S. muticum (collected at Saint Gildas, France) biochemical composition for two years reveals that in certain extent, this marine alga tends to show quite similar pattern. Protein and ash content of S. muticum is lower during summer and increases in autumn to spring. On the opposite, sugar content is higher in summer and decreases from autumn to spring. In addition, the reproductive cycle also contributes to the biochemical composition in S. muticum. In comparison with other Sargassum species, such variation exhibits similar tendency. The seasonal variation study indicates that there is chemodiversity in Sargassum species driven by the fluctuated conditions of the surroundings. Present study serves as fundamental information for further possible valorization of algal material in food industries as source of natural nutrition.

Beside the nutritional value of *Sargassum*, this species shows a promising potential for biotechnological application due to the presence of its phlorotannins. This compound is widely known to possess strong antioxidant activity than other polyphenol compounds generated by its long polymers of hydroxyl groups. In addition, this compound also contributes for tyrosinase, elastase and biofilm inhibition, and antiviral activity. It is also proven to be non toxic. Most of the activity showed by *Sargassum* polyphenol originates from the enzymatic extracts. Enzymatic extraction is meant to improve the extraction yield and polyphenol

content in *Sargassum*, and to enhance the bioactivities since it is expected to liberate more polyphenol than the conventional extraction, *i.e.* solid-liquid extraction.

This study has demonstrated that the application of enzymes in extracting the phenolic compound in *Sargassum* is quite effective. In addition, the enzymatic extracts of Sargassum demonstrate a vast array of bioactivities. Based on the results presented in this study, different enzymes used give various results concerning the total phenolic content and their bioactivity. Among the eight enzymes, Viscozyme releases higher content of phenolic compounds and it is correlated with its strong antiradical activity. This might be supported by the optimum conditions provided, i.e. 50 °C for 3 H, during the extraction process, which is favorable for the Viscozyme, extract activity. Since phenolic compounds are sensitive to heat and easily oxidized by the exposure of light, it is strongly recommended to apply a mild temperature between 40 to 60 °C. Besides the extraction condition, in the case of phenolic compound, it is important to take the period of collection into further consideration. As commonly known, the production of phenolic compound is mostly stimulated by their environmental condition, such as the presence of grazer and direct exposition to sunlight. As a result, the content will vary according to the season.

Another important note to take is that by using enzymes, the active molecules released might not exclusively come from the phenolic groups. It is possible that it also extracts other molecules since the objective of this method is to degrade algae cell walls. Therefore, it will not be surprising that the fucoidan and protein might be responsible for the bioactivity in *Sargassum*. As a consequence, it will be interesting to further work on the purified samples pretreated by the enzyme-assisted extraction in the future. In addition, the characterization of *Sargassum* phenolic compound might help in understanding more about the mechanisms of their bioactivity.

Exploitation of marine natural bioactive compounds involves the collection of living resources for commercial purposes, sustainable factor must be considered along the process. Some bioactive compounds particularly those induced by the surrounding, in other word, exist as secondary metabolites, often present in low concentrations. In the search of marine bioactive compounds for commercial purposes, a substantial amount of samples is important to consider in providing an adequate amount of target compounds to complete the whole process. In addition, a sustainable resource also becomes important factor in the search of marine natural bioactive compound.

Regarding the exploitation of brown algal phlorotannins derived from Sargassum, its massive biomass influx offers certain advantage in terms of valorization. However, considering that the content of phlorotannins is merely up to 20%, it is strongly suggested to determine and evaluate the supply chain management of this species for future exploitation. As an alternative if the natural population could not fulfill the required supply of biomass, cultivation approach might be one of the possible propositions in order to provide algal biomass in substantial amount and sustainably. Furthermore, governmental support as well as reliable and integrated collaboration of inter-institutional research is also required. understanding and reliable collaboration between An government, universities/institutes and stakeholders - private sectors or industries, will bring remarkable output for the interest of marine algal natural compound.

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## **Appendix 1 Documentations**





A. Left: S. muticum collected in August 2014 Right: S. muticum in August 2014 covered with epiphytes



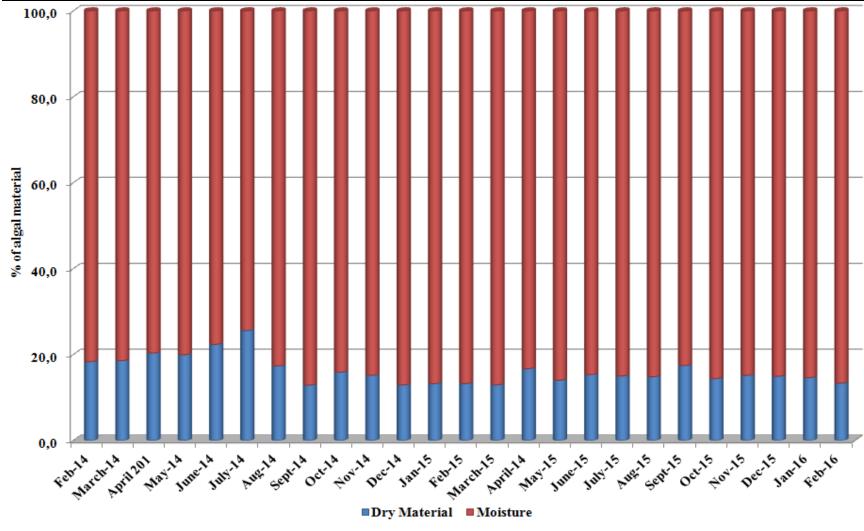


B. Left: S. muticum from September 2014 Right: S. muticum from January 2015



C. S. muticum collected in January 2015 (winter) (A), S. muticum in May 2015 (Spring) (B), S. muticum in June 2015 (Summer) (C), and S. muticum November 2015 (Autumn) (D).

## Appendix 2 S. muticum moisture content and dry material



D Histogram of S. muticum moisture content and dry material expressed in % of dry material.

## Appendix 3 Meteorological data of Vannes, Morbihan

Meteorological data from www.meteociel.fr

Months	Sunlight (h/mois)	Tmax (°C) Vannes	Tmin (°C) Vannes	Precipitations (mm/mois)	Sea surface temperature (°C)
February 2014	75,2	11,8	5,4	202,4	10
March 2014	166,9	14,4	3,7	52,5	11
April 2014	179,6	16,8	7,2	56,7	12
May 2014	213,6	18,1	8,6	52,8	13
June 2014	270,7	23,2	12,4	43,0	17
July 2014	242,1	25,0	14,1	46,2	18
August 2014	198,1	21,7	12,4	119,2	18
September 2014	250,1	24,9	12,5	4,8	19
October 2014	138,1	19,5	10,3	90,0	16
November 2014	61,5	14,8	7,9	173,9	14
December 2014	57,8	10,8	4,7	58,0	12
January 2015	72,7	10,6	3,4	105,1	9
February 2015	127,7	9,8	1,3	42,2	9
March 2015	129,0	12,9	4,8	18,4	9
April 2015	204,2	18,0	7,3	57,4	13
May 2015	171,4	18,0	9,3	64,3	13
June 2015	290,7	22,9	11,4	4,2	16
July 2015	181,8	23,5	14,0	68,7	17
August 2015	179,6	23,0	13,4	101,2	18
September 2015	217,0	20,2	10,0	80,7	17
October 2015	123,8	17,2	8,5	40,0	14
November 2015	66,9	15,8	9,9	68,6	14
December 2015	47,6	14,2	9,6	76,4	12
January 2016	50,2	11,0	4,4	114,5	9
February 2016	84,9	11,1	4,5	95,8	9

# Appendix 4 Analysis of variance of *S. muticum* biochemical composition – seasonal variation

#### **Descriptive Statistics**

Dependent Variable: DryMat

Seasons	Mean	Std. Deviation	N
Autumn	15,1461	1,61266	18
Spring	17,0818	3,01028	18
Summer	18,3994	4,48324	18
Winter	14,3590	1,93085	21
Total	16,1711	3,30169	75

#### Levene's Test of Equality of Error Variances<sup>a</sup>

Dependent Variable: DryMat

F	df1	df2	Sig.	
10,080	3	71	,000	

Tests the null hypothesis that the error variance of the dependent variable is equal across groups.

a. Design: Intercept + Seasons

#### **Tests of Between-Subjects Effects**

Dependent Variable: DryMat

	Type III Sum of				
Source	Squares	df	Mean Square	F	Sig.
Corrected Model	192,172 <sup>a</sup>	3	64,057	7,401	,000
Intercept	19708,405	1	19708,405	2277,074	,000
Seasons	192,172	3	64,057	7,401	,000
Error	614,515	71	8,655		
Total	20419,507	75			
Corrected Total	806,688	74			

a. R Squared = ,238 (Adjusted R Squared = ,206)

## **Estimated Marginal Means**

#### 1. Grand Mean

Dependent Variable: DryMat

		95% Confidence Interval			
Mean	Std. Error	Lower Bound	Upper Bound		
16,247	,340	15,568	16,925		

#### 2. Seasons

Dependent Variable: DryMat

			95% Confidence Interval		
Seasons	Mean	Std. Error	Lower Bound	Upper Bound	
Autumn	15,146	,693	13,763	16,529	
Spring	17,082	,693	15,699	18,464	
Summer	18,399	,693	17,017	19,782	
Winter	14,359	,642	13,079	15,639	

#### Post Hoc Tests Seasons

#### **Multiple Comparisons**

Dependent Variable: DryMat

Tukev HSD

Tukey HSD						
	-	Mean Difference			95% Confid	ence Interval
(I) Seasons	(J) Seasons	(I-J)	Std. Error	Sig.	Lower Bound	Upper Bound
Autumn	Spring	-1,9357	,98065	,208	-4,5157	,6444
	Summer	-3,2533 <sup>*</sup>	,98065	,008	-5,8334	-,6733
	Winter	,7871	,94498	,839	-1,6991	3,2733
Spring	Autumn	1,9357	,98065	,208	-,6444	4,5157
	Summer	-1,3177	,98065	,539	-3,8977	1,2624
	Winter	2,7227*	,94498	,026	,2365	5,2089
Summer	Autumn	3,2533 <sup>*</sup>	,98065	,008	,6733	5,8334
	Spring	1,3177	,98065	,539	-1,2624	3,8977
	Winter	4,0404*	,94498	,000	1,5542	6,5266
Winter	Autumn	-,7871	,94498	,839	-3,2733	1,6991
	Spring	-2,7227 <sup>*</sup>	,94498	,026	-5,2089	-,2365
	Summer	-4,0404 <sup>*</sup>	,94498	,000	-6,5266	-1,5542

Based on observed means.

The error term is Mean Square(Error) = 8,655.

<sup>\*.</sup> The mean difference is significant at the ,05 level.

## **Homogeneous Subsets**

DryMat

Tukey HSD<sup>a,b,c</sup>

rakey riel							
		Subset					
Seasons	N	1	2	3			
Winter	21	14,3590					
Autumn	18	15,1461	15,1461				
Spring	18		17,0818	17,0818			
Summer	18			18,3994			
Sig.		,846	,194	,523			

Means for groups in homogeneous subsets are displayed.

Based on observed means.

The error term is Mean Square(Error) = 8,655.

- a. Uses Harmonic Mean Sample Size = 18,667.
- b. The group sizes are unequal. The harmonic mean of the group sizes is used. Type I error levels are not guaranteed.
- c. Alpha = ,05.

#### **Descriptives**

Dependent Variable: Biochemical

		N	Mana	Std.	Std.		lence Interval Mean	Minimo	Massinassea
		N	Mean	Deviation	Error	Lower Bound	Upper Bound	Minimum	Maximum
Uronic Acid	Spring	18	2,4033	1,27131	,29965	1,7711	3,0355	1,37	6,88
	Summer	18	2,3067	,77705	,18315	1,9203	2,6931	1,36	4,10
	Autumn	18	2,2472	,37386	,08812	2,0613	2,4331	1,57	3,08
	Winter	21	2,1567	,49771	,10861	1,9301	2,3832	1,50	3,05
	Total	75	2,2736	,78580	,09074	2,0928	2,4544	1,36	6,88
Protein	Spring	18	27,0628	3,39211	,79953	25,3759	28,7496	22,35	34,88
	Summer	18	21,8428	6,36032	1,49914	18,6799	25,0057	9,64	30,71
	Autumn	18	27,1700	10,39115	2,44922	22,0026	32,3374	11,69	59,71
	Winter	21	27,4352	2,62323	,57244	26,2412	28,6293	20,70	32,99
	Total	75	25,9400	6,63283	,76589	24,4139	27,4661	9,64	59,71
Sugar	Spring	18	10,8828	3,19887	,75398	9,2920	12,4735	5,57	17,22
	Summer	18	14,7939	2,39312	,56406	13,6038	15,9840	11,69	19,94
	Autumn	18	14,3428	3,38238	,79723	12,6608	16,0248	7,19	19,54
	Winter	21	11,6095	3,67370	,80167	9,9373	13,2818	5,60	17,19
	Total	75	12,8553	3,57511	,41282	12,0328	13,6779	5,57	19,94
Sulfates	Spring	18	10,1683	1,97514	,46554	9,1861	11,1505	7,95	12,43
Groups	Summer	18	10,6561	3,42244	,80668	8,9542	12,3580	7,01	15,26
	Autumn	18	9,5972	2,42427	,57141	8,3917	10,8028	6,90	14,65
	Winter	21	9,2757	2,03859	,44486	8,3478	10,2037	1,84	11,77
	Total	75	9,8984	2,51990	,29097	9,3186	10,4782	1,84	15,26
Polyphenol	Spring	18	2,1550	,34658	,08169	1,9826	2,3274	1,38	2,76
	Summer	18	2,3511	,95636	,22542	1,8755	2,8267	,79	3,87
	Autumn	18	3,1517	1,73673	,40935	2,2880	4,0153	,69	6,43
	Winter	21	2,5462	,64325	,14037	2,2534	2,8390	1,78	3,44
	Total	75	2,5508	1,08540	,12533	2,3011	2,8005	,69	6,43

Ash	Spring Summer Autumn Winter Total	18 18 18 21 75	23,5256 22,3694 25,4306 25,8014 24,3425	2,90156 3,42822 3,93399 2,25975 3,40159	,68390 ,80804 ,92725 ,49312 ,39278	22,0826 20,6646 23,4742 24,7728 23,5599	24,0743	17,09 16,57 19,00 21,14 16,57	28,04 27,69 31,16 30,00 31,16
Nitrogen	Spring	18	2,2906	,49929	,11768	2,0423	2,5388	1,25	2,83
	Summer	18	1,6728	,69131	,16294	1,3290	2,0166	1,10	3,25
	Autumn	18	1,9517	,59993	,14141	1,6533	2,2500	1,37	3,25
	Winter	21	2,0090	,35058	,07650	1,8495	2,1686	1,40	2,59
	Total	75	1,9821	,57429	,06631	1,8500	2,1143	1,10	3,25

**Test of Homogeneity of Variances** 

	Levene Statistic	df1	df2	Sig.
UronicAcid	3,002	3	71	,036
Protein	5,081	3	71	,003
Sugar	,858,	3	71	,467
SulfatesGroups	10,229	3	71	,000
Polyphenol	17,410	3	71	,000
Ash	3,663	3	71	,016
Nitrogen	1,554	3	71	,208

#### **ANOVA**

		Sum of Squares	df	Mean Square	F	Sig.
Uronic Acid	Between Groups	,622	3	,207	,327	,806
	Within Groups	45,071	71	,635		
	Total	45,693	74			
Protein	Between Groups	399,044	3	133,015	3,306	,025
	Within Groups	2856,540	71	40,233		
	Total	3255,585	74			
Sugar	Between Groups	210,099	3	70,033	6,758	,000
	Within Groups	735,726	71	10,362		
	Total	945,825	74			
Sulfates Groups	Between Groups	21,421	3	7,140	1,130	,343
	Within Groups	448,470	71	6,316		
	Total	469,891	74			
Polyphenol	Between Groups	10,037	3	3,346	3,079	,033
	Within Groups	77,142	71	1,087		
	Total	87,179	74			
Ash	Between Groups	148,094	3	49,365	4,949	,004
	Within Groups	708,146	71	9,974		
	Total	856,240	74			
Nitrogen	Between Groups	3,467	3	1,156	3,918	,012
	Within Groups	20,939	71	,295		
	Total	24,406	74			

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## **Post Hoc Tests**

### **Multiple Comparisons**

Tukey HSD

Tukey HSD		_	Mean			95% Confider	ce Interval
Dependent			Difference	Std.		Lower	Upper
Variable	(I) Season	(J) Season	(I-J)	Error	Sig.	Bound	Bound
Uronic Acid	Spring	Summer	,09667	,26558	,983	-,6021	,7954
		Autumn	,15611	,26558	,935	-,5426	,8548
		Winter	,24667	,25592	,770	-,4266	,9200
	Summer	Spring	-,09667	,26558	,983	-,7954	,6021
		Autumn	,05944	,26558	,996	-,6393	,7582
		Winter	,15000	,25592	,936	-,5233	,8233
	Autumn	Spring	-,15611	,26558	,935	-,8548	,5426
		Summer	-,05944	,26558	,996	-,7582	,6393
		Winter	,09056	,25592	,985	-,5828	,7639
	Winter	Spring	-,24667	,25592	,770	-,9200	,4266
		Summer	-,15000	,25592	,936	-,8233	,5233
		Autumn	-,09056	,25592	,985	-,7639	,5828
Protein	Spring	Summer	5,22000	2,11432	,074	-,3426	10,7826
	-1 3	Autumn	-,10722	2,11432	1,000	-5,6699	5,4554
		Winter	-,37246	2,03741	,998	-5,7327	4,9878
	Summer	Spring	-5,22000	2,11432	,074	-10,7826	,3426
		Autumn	-5,32722	2,11432	,065	-10,8899	,2354
		Winter	-5,59246 <sup>*</sup>	2,03741	,038	-10,9527	-,2322
	Autumn	Spring	,10722	2,11432	1,000	-5,4554	5,6699
		Summer	5,32722	2,11432	,065	-,2354	10,8899
		Winter	-,26524	2,03741	,999	-5,6255	5,0950
	Winter	Spring	,37246	2,03741	,998	-4,9878	5,7327
		Summer	5,59246	2,03741	,038	,2322	10,9527
0	0	Autumn	,26524	2,03741	,999	-5,0950	5,6255
Sugar	Spring	Summer Autumn	-3,91111 <sup>*</sup>	1,07302	,003	-6,7342	-1,0881
		Winter	-3,46000 -,72675	1,07302 1,03399	,010, 896,	-6,2830 -3,4471	-,6370 1,9936
	Summer		3,91111 <sup>*</sup>				
	Summer	Spring Autumn	,45111	1,07302 1,07302	,003 ,975	1,0881 -2,3719	6,7342 3,2742
		Winter	3,18437 <sup>*</sup>	1,07302	,975	,4640	5,9047
	Autumn	Spring	3,46000°	1,07302	,010	,6370	6,2830
	/ tatariiri	Summer	-,45111	1,07302	,975	-3,2742	2,3719
		Winter	2,73325	1,03399	,048	,0129	5,4536
	Winter	Spring	,72675	1,03399	,896	-1,9936	3,4471
		Summer	-3,18437 <sup>*</sup>	1,03399	,015	-5,9047	-,4640
		Autumn	-2,73325 <sup>*</sup>	1,03399	,048	-5,4536	-,0129
Sulfates	Spring	Summer	-,48778	,83775	,937	-2,6919	1,7163
Groups		Autumn	,57111	,83775	,904	-1,6330	2,7752
		Winter	,89262	,80728	,687	-1,2313	3,0165
	Summer	Spring	,48778	,83775	,937	-1,7163	2,6919
		Autumn	1,05889	,83775	,589	-1,1452	3,2630
		Winter	1,38040	,80728	,326	-,7435	3,5043
	Autumn	Spring	-,57111	,83775	,904	-2,7752	1,6330
ĺ		Summer	-1,05889	,83775	,589	-3,2630	1,1452
	Mintor	Winter	,32151	,80728	,978	-1,8024	2,4454
	Winter	Spring	-,89262 -1 38040	,80728	,687 326	-3,0165 -3,5043	1,2313
		Summer	-1,38040 -,32151	,80728 ,80728	,326	-3,5043 -2,4454	,7435, 1,8024
		Autumn	-,32131	,00120	,978	-2, <del>44</del> 04	1,0024

Polyphenol	Spring	Summer	-,19611	,34745	,942	-1,1102	,7180
		Autumn	-,99667 <sup>*</sup>	,34745	,027	-1,9108	-,0825
		Winter	-,39119	,33481	,649	-1,2721	,4897
	Summer	Spring	,19611	,34745	,942	-,7180	1,1102
		Autumn	-,80056	,34745	,107	-1,7147	,1136
		Winter	-,19508	,33481	,937	-1,0760	,6858
	Autumn	Spring	,99667	,34745	,027	,0825	1,9108
		Summer	,80056	,34745	,107	-,1136	1,7147
		Winter	,60548	,33481	,278	-,2754	1,4863
	Winter	Spring	,39119	,33481	,649	-,4897	1,2721
		Summer	,19508	,33481	,937	-,6858	1,0760
		Autumn	-,60548	,33481	,278	-1,4863	,2754
Ash	Spring	Summer	1,15611	1,05272	,692	-1,6135	3,9257
		Autumn	-1,90500	1,05272	,277	-4,6746	,8646
		Winter	-2,27587	1,01442	,122	-4,9448	,3930
	Summer	Spring	-1,15611	1,05272	,692	-3,9257	1,6135
		Autumn	-3,06111 <sup>*</sup>	1,05272	,024	-5,8307	-,2915
		Winter	-3,43198 <sup>*</sup>	1,01442	,006	-6,1009	-,7631
	Autumn	Spring	1,90500	1,05272	,277	-,8646	4,6746
		Summer	3,06111*	1,05272	,024	,2915	5,8307
		Winter	-,37087	1,01442	,983	-3,0398	2,2980
	Winter	Spring	2,27587	1,01442	,122	-,3930	4,9448
		Summer	3,43198 <sup>*</sup>	1,01442	,006	,7631	6,1009
		Autumn	,37087	1,01442	,983	-2,2980	3,0398
Nitrogen	Spring	Summer	,61778 <sup>^</sup>	,18102	,006	,1415	1,0940
		Autumn	,33889	,18102	,249	-,1374	,8151
		Winter	,28151	,17444	,378	-,1774	,7404
	Summer	Spring	-,61778 <sup>*</sup>	,18102	,006	-1,0940	-,1415
		Autumn	-,27889	,18102	,419	-,7551	,1974
		Winter	-,33627	,17444	,226	-,7952	,1227
	Autumn	Spring	-,33889	,18102	,249	-,8151	,1374
		Summer	,27889	,18102	,419	-,1974	,7551
		Winter	-,05738	,17444	,988	-,5163	,4016
	Winter		-,03738	,17444	,378	-,7404	,4010
	vviiitei	Spring	· ·			*	
		Summer	,33627	,17444	,226	-,1227	,7952
		Autumn	,05738	,17444	,988	-,4016	,5163

 $<sup>^{\</sup>star}.$  The mean difference is significant at the 0.05 level.

## **Homogeneous Subsets**

Tukey HSD<sup>a,b</sup> Uronic Acid

		Subset for alpha = 0.05
Season	N	1
Winter	21	2,1567
Autumn	18	2,2472
Summer	18	2,3067
Spring	18	2,4033
Sig.		,780

a b		Protein
Tukey HSD <sup>a,o</sup>	Tukey HSD <sup>a,b</sup>	

		Subset for alpha = 0.05	
Season	N	1	2
Summer	18	21,8428	
Spring Autumn	18 18	27,0628 27,1700	27,0628 27,1700
Winter	21		27,4352
Sig.		,059	,998

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Sugar

Tukey HSD <sup>a,b</sup>					
		Subset for alpha = 0.05			
Season	N	1	2	3	
Spring	18	10,8828			
Winter	21	11,6095	11,6095		
Autumn	18		14,3428	14,3428	
Summer	18			14,7939	
Sig.		,901	,055	,973	

**Sulfates Groups** 

Tukey HSD<sup>a,b</sup>

		Subset for alpha = 0.05
Season	Ν	1
Winter	21	9,2757
Autumn	18	9,5972
Spring	18	10,1683
Summer	18	10,6561
Sig.		,343

Ash

Tukey HSD<sup>a,b</sup>

		Subset for a	lpha = 0.05
Season	N	1	2
Summer	18	22,3694	
Spring	18	23,5256	23,5256
Autumn	18		25,4306
Winter	21		25,8014
Sig.		,679	,133

## Nitrogen

Tukey HSD <sup>a,</sup>	Nit b

		Subset for a	alpha = 0.05
Season	N	1	2
Summer	18	1,6728	
Autumn	18	1,9517	1,9517
Winter	21	2,0090	2,0090
Spring	18		2,2906
Sig.		,241	,235

## **Polyphenol** Tukey HSD<sup>a,b</sup>

		Subset for alpha = 0.05	
Season	N	1	2
Spring	18	2,1550	
Summer	18	2,3511	2,3511
Winter	21	2,5462	2,5462
Autumn	18		3,1517
Sig.		,662	,097

Means for groups in homogeneous subsets are displayed.

- a. Uses Harmonic Mean Sample Size = 18,667.
- b. The group sizes are unequal. The harmonic mean of the group sizes is used. Type I error levels are not guaranteed.

### Appendix 5 Pearson's Analysis of S. muticum biochemical composition – seasonal variation

Polyphenol

1.635

,022

75

Cross-products

Covariance

N

319,387

4,316

75

102,673

1,387

75

-121.134

-1,637

75

#### **Correlations** Uronic Sulfates SST Protein Sugar Polyphenol Ash Nitrogen Sunlight Tmax Tmin Precipitation Acid Groups Pearson Correlation -,169 -,317<sup>\*</sup> ,248\* .075 ,024 -,016 -,180 -,050 1 .053 ,026 ,050 Sig. (2-tailed) ,654 ,147 ,006 ,825 ,668 ,032 ,521 .838 ,892 ,123 ,668 Sum of Squares and Uronic Acid 45,693 20,291 -35,188 8,293 319,000 -46,482 1,635 9,956 6,816 -3,373 -476,006 -9,661 Cross-products Covariance ,617 ,274 -,476 -,628 ,022 ,135 .112 4,311 .092 -,046 -,131 -6,433 N 75 75 75 75 75 75 75 75 75 75 75 75 -,257\* ,600\* -,314 -,257 -**,**250<sup>\*</sup> -,220 Pearson Correlation ,053 ,177 ,181 -,162 ,128 Sig. (2-tailed) ,129 ,026 .030 .057 ,654 ,000, ,121 ,006 ,165 ,026 ,272 Sum of Squares and 20,291 3255,585 310,459 -317,395 319,387 301,538 -88,467 -5805,906 -613,701 -446,013 2870,621 -356,740 Protein Cross-products Covariance ,274 43,994 4,195 -4,289 4,075 -1,195 -6,027 -4,821 4,316 -78,458 -8,293 38,792 75 75 75 75 75 75 75 75 75 75 75 75 N Pearson Correlation ,177 -,030 ,358 -,030 -,502 ,267 ,336 ,271 ,350<sup>\*</sup> -,169 .099 Sig. (2-tailed) .147 ,129 .800 ,002 ,799 ,000 ,399 ,020 ,003 ,018 ,002 Sum of Squares and -35,188 310,459 945,825 -19,783 102,673 -26,899 -76,304 1906,394 344,780 322,423 3271,073 304,966 Sugar Cross-products -,476 1,387 Covariance 4,195 12,781 -,267 -,364 -1,031 25,762 4,659 4,357 44,204 4,121 75 75 75 75 75 N 75 75 75 75 75 75 75 .378\* Pearson Correlation -.317 -,257 -,598 .235 .298\* .298\* -.030 -.065 .195 -.031 Sig. (2-tailed) ,006 ,026 ,800 000, ,582 ,042 ,001 ,010 ,094 ,792 ,009 Sum of Squares and Sulfates Groups -46,482 -317.395 -19,783 469,891 -121.134-40,925 25,215 5142,651 270,416 131.614 -263,165 183,357 Cross-products Covariance -,628 -4,289 -,267 6,350 -1,637 -,553 ,341 69,495 3,654 1,779 2,478 -3,556 N 75 75 75 75 75 75 75 75 75 75 75 75 ,600<sup>\*</sup> .358\* -,598\* .077 -,452 Pearson Correlation .026 -.162 -.046 .067 .054 .000 1 Sig. (2-tailed) ,825 ,000 ,002 ,000 .509 ,000, ,164 ,696 .568 .999 ,646 Sum of Squares and

87,179

1,178

75

21.150

,286

75

-20,838

-,282

75

-950,945

-12,851

75

-17.935

-,242

75

19,526

,264

75

197,019

2,662

.026

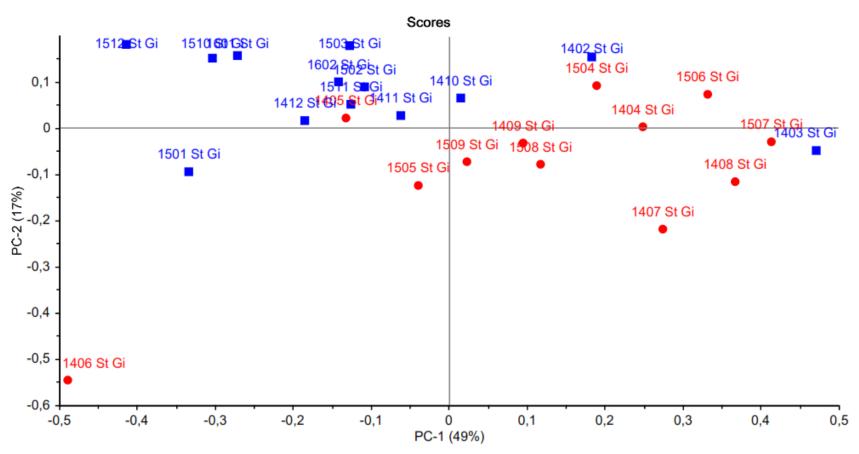
000,

75

		Uronic Acid	Protein	Sugar	Sulfates Groups	Polyphenol	Ash	Nitrogen	Sunlight	Tmax	Tmin	Precipitation	SST
	Pearson Correlation	,050	,181	-,030	-,065	,077	1	-,089	-,509**	-,454**	-,377**	,237*	-,331**
	Sig. (2-tailed)	,668	,121	,799	,582	,509		,450	,000	,000	,001	,040	,004
Ash	Sum of Squares and Cross-products	9,956	301,538	-26,899	-40,925	21,150	856,240	-12,806	-9340,351	-556,718	-344,421	2720,728	-274,476
	Covariance	,135	4,075	-,364	-,553	,286	11,571	-,173	-126,221	-7,523	-4,654	36,767	-3,709
	N	75	75	75	75	75	75	75	75	75	75	75	75
	Pearson Correlation	,248*	-,314**	-,502**	,235*	-,452**	-,089	1	,113	-,099	-,190	-,172	-,203
	Sig. (2-tailed)	,032	,006	,000	,042	,000	,450		,333	,398	,102	,139	,081
Nitrogen	Sum of Squares and Cross-products	8,293	-88,467	-76,304	25,215	-20,838	-12,806	24,406	351,738	-20,499	-29,366	-333,662	-28,450
	Covariance	,112	-1,195	-1,031	,341	-,282	-,173	,330	4,753	-,277	-,397	-4,509	-,384
	N	75	75	75	75	75	75	75	75	75	75	75	75

<sup>\*\*.</sup> Correlation is significant at the 0.01 level (2-tailed).

\*. Correlation is significant at the 0.05 level (2-tailed).



E FT-IR spectra of *S. muticum* biochemical composition at 1800-600 cm<sup>-1</sup>; Red color: spring-summer (hot season), Blue color: autumnwinter (cold season).

# Appendix 7 Analysis of variance of *S. muticum* polyphenol content – seasonal variation with solid-liquid extraction

Levene's Test of Equality of Error Variances<sup>a</sup>

	F	df1	df2	Sig.
Water	40,221	3	56	,000
MeOH	14,900	3	56	,000
MeOH50	17,337	3	56	,000
EtOH75	7,862	3	56	,000

Tests the null hypothesis that the error variance of the dependent variable is equal across groups.

a. Design: Intercept + Season

Tests of Between-Subjects Effects

	Dependent	Type III Sum of				
Source	Variable	Squares	df	Mean Square	F	Sig.
Corrected Model	Water	50,375 <sup>a</sup>	3	16,792	17,617	,000
	MeOH	18,804 <sup>b</sup>	3	6,268	2,728	,052
	MeOH50	71,240 <sup>c</sup>	3	23,747	8,594	,000
	EtOH75	20,919 <sup>d</sup>	3	6,973	5,701	,002
Intercept	Water	399,614	1	399,614	419,256	,000
	MeOH	337,561	1	337,561	146,937	,000
	MeOH50	777,601	1	777,601	281,402	,000
	EtOH75	385,020	1	385,020	314,783	,000
Season	Water	50,375	3	16,792	17,617	,000
	MeOH	18,804	3	6,268	2,728	,052
	MeOH50	71,240	3	23,747	8,594	,000
	EtOH75	20,919	3	6,973	5,701	,002
Error	Water	53,376	56	,953		
	MeOH	128,650	56	2,297		
	MeOH50	154,745	56	2,763		
	EtOH75	68,495	56	1,223		
Total	Water	541,908	60			
	MeOH	514,943	60			
	MeOH50	1047,977	60			
	EtOH75	503,955	60			
Corrected Total	Water	103,751	59			
	MeOH	147,454	59			
	MeOH50	225,985	59			
	EtOH75	89,414	59			

a. R Squared = ,486 (Adjusted R Squared = ,458)

- b. R Squared = ,128 (Adjusted R Squared = ,081)
- c. R Squared = ,315 (Adjusted R Squared = ,279)
- d. R Squared = ,234 (Adjusted R Squared = ,193)

# **Estimated Marginal Means**

#### Season

				95% Confide	ence Interval
Dependent Variable	Season	Mean	Std. Error	Lower Bound	Upper Bound
Water	Autumn	3,108	,282	2,543	3,672
	Spring	2,099	,230	1,638	2,560
	Summer	3,853	,230	3,392	4,314
	Winter	1,477	,282	,912	2,041
MeOH	Autumn	2,890	,438	2,013	3,767
	Spring	2,487	,357	1,772	3,203
	Summer	2,894	,357	2,179	3,610
	Winter	1,412	,438	,535	2,288
MeOH50	Autumn	4,235	,480	3,274	5,196
	Spring	2,534	,392	1,750	3,319
	Summer	5,085	,392	4,300	5,870
	Winter	2,843	,480	1,881	3,804
EtOH75	Autumn	2,881	,319	2,241	3,520
	Spring	2,224	,261	1,702	2,746
	Summer	3,378	,261	2,856	3,900
	Winter	1,859	,319	1,220	2,499

#### Post Hoc Tests Season

#### **Multiple Comparisons**

Tukey HSD

Tukey HSD			Mean			95% Confid	dence Interval
Dependent		(J)	Difference			Lower	
Variable	(I) Season	Season	(I-J)	Std. Error	Sig.	Bound	Upper Bound
Water	Autumn	Spring	1,0086	,36384	,037	,0452	1,9720
		Summer	-,7453	,36384	,183	-1,7087	,2181
		Winter	1,6308 <sup>*</sup>	,39857	,001	,5755	2,6862
	Spring	Autumn	-1,0086 <sup>*</sup>	,36384	,037	-1,9720	-,0452
	Opinig	Summer	-1,7539 <sup>*</sup>	,32543	,000	-2,6156	-,8922
		Winter	,6222	,36384	,328	-,3412	1,5856
	Cummor		,				
	Summer	Autumn	,7453	,36384	,183	-,2181	1,7087
		Spring	1,7539	,32543	,000	,8922	2,6156
		Winter	2,3761*	,36384	,000	1,4127	3,3395
	Winter	Autumn	-1,6308 <sup>*</sup>	,39857	,001	-2,6862	-,5755
		Spring	-,6222	,36384	,328	-1,5856	,3412
		Summer	-2,3761 <sup>*</sup>	,36384	,000	-3,3395	-1,4127
MeOH	Autumn	Spring	,4028	,56487	,892	-1,0929	1,8985
		Summer	-,0044	,56487	1,000	-1,5001	1,4913
		Winter	1,4783	,61878	,091	-,1601	3,1168
	Spring	Autumn	-,4028	,56487	,892	-1,8985	1,0929
		Summer	-,4072	,50523	,851	-1,7450	,9306
		Winter	1,0756	,56487	,238	-,4201	2,5713
	Summer	Autumn	,0044	,56487	1,000	-1,4913	1,5001
		Spring	,4072	,50523	,851	-,9306	1,7450
		Winter	1,4828	,56487	,053	-,0129	2,9785
	Winter	Autumn	-1,4783	,61878	,091	-3,1168	,1601
		Spring	-1,0756	,56487	,238	-2,5713	,4201
		Summer	-1,4828	,56487	,053	-2,9785	,0129

MeOH50	Autumn	Spring	1,7006	,61951	,039	,0602	3,3409
		Summer	-,8500	,61951	,522	-2,4904	,7904
		Winter	1,3925	,67864	,182	-,4045	3,1895
	Spring	Autumn	-1,7006 <sup>*</sup>	,61951	,039	-3,3409	-,0602
		Summer	-2,5506 <sup>*</sup>	,55411	,000	-4,0178	-1,0833
		Winter	-,3081	,61951	,959	-1,9484	1,3323
	Summer	Autumn	,8500	,61951	,522	-,7904	2,4904
		Spring	2,5506 <sup>*</sup>	,55411	,000	1,0833	4,0178
		Winter	2,2425*	,61951	,003	,6021	3,8829
	Winter	Autumn	-1,3925	,67864	,182	-3,1895	,4045
		Spring	,3081	,61951	,959	-1,3323	1,9484
		Summer	-2,2425 <sup>*</sup>	,61951	,003	-3,8829	-,6021
EtOH75	Autumn	Spring	,6569	,41216	,390	-,4344	1,7483
		Summer	-,4969	,41216	,626	-1,5883	,5944
		Winter	1,0217	,45150	,119	-,1739	2,2172
	Spring	Autumn	-,6569	,41216	,390	-1,7483	,4344
		Summer	-1,1539 <sup>*</sup>	,36865	,014	-2,1300	-,1777
		Winter	,3647	,41216	,813	-,7266	1,4561
	Summer	Autumn	,4969	,41216	,626	-,5944	1,5883
		Spring	1,1539 <sup>*</sup>	,36865	,014	,1777	2,1300
		Winter	1,5186 <sup>*</sup>	,41216	,003	,4272	2,6100
	Winter	Autumn	-1,0217	,45150	,119	-2,2172	,1739
		Spring	-,3647	,41216	,813	-1,4561	,7266
		Summer	-1,5186 <sup>*</sup>	,41216	,003	-2,6100	-,4272

<sup>\*.</sup> The mean difference is significant at the ,05 level.

#### **Homogeneous Subsets**

#### Water

Tukev HSD<sup>a,b,c</sup>

Tukey HSD			
		Sub	set
Season	N	1	2
Winter	12	1,4767	
Spring	18	2,0989	
Autumn	12		3,1075
Summer	18		3,8528
Sig.		,328	,183

Means for groups in homogeneous subsets are displayed.

Based on observed means.

The error term is Mean Square(Error) = ,953.

- a. Uses Harmonic Mean Sample Size = 14,400.
- b. The group sizes are unequal. The harmonic mean of the group sizes is used.
   Type I error levels are not guaranteed.
- c. Alpha = ,05.

#### MeOH

Tukev HSD<sup>a,b,c</sup>

Tukey HSD		
		Subset
Season	N	1
Winter	12	1,4117
Spring	18	2,4872
Autumn	12	2,8900
Summer	18	2,8944
Sig.		,053

Means for groups in homogeneous subsets are displayed.

Based on observed means.

The error term is Mean Square(Error) = 2,297.

- a. Uses Harmonic Mean Sample Size = 14,400.
- b. The group sizes are unequal. The harmonic mean of the group sizes is used. Type I error levels are not guaranteed.
- c. Alpha = ,05.

#### MeOH50

Tukey HSD<sup>a,b,c</sup>

		Subset					
Season	N	1	2	3			
Spring	18	2,5344					
Winter	12	2,8425	2,8425				
Autumn	12		4,2350	4,2350			
Summer	18			5,0850			
Sig.		,959	,123	,522			

Means for groups in homogeneous subsets are displayed.

Based on observed means.

The error term is Mean Square(Error) = 2,763.

- a. Uses Harmonic Mean Sample Size = 14,400.
- b. The group sizes are unequal. The harmonic mean of the group sizes is used. Type I error levels are not guaranteed.

c. Alpha = ,05.

EtOH75

Tukey HSD<sup>a,b,c</sup>

randy ridb						
		Subset				
Season	N	1	2			
Winter	12	1,8592				
Spring	18	2,2239				
Autumn	12	2,8808	2,8808			
Summer	18		3,3778			
Sig.		,074	,626			

Means for groups in homogeneous subsets are displayed.

Based on observed means.

The error term is Mean Square(Error) = 1,223.

b. The group sizes are unequal. The harmonic mean of the group sizes is used. Type I error levels are not guaranteed.

c. Alpha = ,05.

#### **Correlations**

**Descriptive Statistics** 

	Mean	Std. Deviation	N
Water	2,7023	1,32608	60
MeOH	2,4748	1,58089	60
MeOH50	3,7013	1,95710	60
EtOH75	2,6285	1,23105	60
Sunlight	152,0480	72,96261	75
Tmax	17,1680	4,87348	75
Tmin	8,4400	3,62760	75
Precipitation	73,4800	45,55026	75
SST	13,5600	3,29717	75

#### Correlations

				Correlation	5					
		Water	MeOH	MeOH50	EtOH75	Sunlight	Tmax	Tmin	Precipitation	SST
Water	Pearson Correlation	1	,672**	,653**	,691	,700**	,777**	,757**	-,336**	,763**
	Sig. (2-tailed)		,000	,000	,000	,000	,000	,000	,009	,000
	N	60	60	60	60	60	60	60	60	60
MeOH	Pearson Correlation	,672**	1	,692**	,747**	,370**	,324*	,318 <sup>*</sup>	-,202	,272*
	Sig. (2-tailed)	,000		,000	,000	,004	,012	,013	,122	,036
	N	60	60	60	60		60	60	60	60
MeOH50	Pearson Correlation	,653**	,692**	1	,698**	,451**	,492**	,516 <sup>**</sup>	-,216	,469**
	Sig. (2-tailed)	,000	,000		,000	,000	,000	,000	,098	,000
	N	60	60	60	60		60	60	60	60
EtOH75	Pearson Correlation	,691**	,747**	,698**	1	,288*	,432**	,483**	-,110	,409**
	Sig. (2-tailed)	,000	,000	,000		,026	,001	,000	,404	,001
	N	60	60	60	60	60	60	60	60	60
Sunlight	Pearson Correlation	,700**	,370**	,451**	,288*	1	,826**	,602	-,551**	,672**
	Sig. (2-tailed)	,000	,004	,000	,026		,000	,000	,000	,000
	N	60	60	60	60		75	75	75	75
Tmax	Pearson Correlation	,777**	,324*	,492**	,432**	,826**	1	,934**	-,341**	,940**
	Sig. (2-tailed)	,000	,012	,000	,001	,000		,000	,003	,000
	N	60	60	60	60		75	75	75	75
Tmin	Pearson Correlation	,757**	,318 <sup>*</sup>	,516 <sup>**</sup>	,483	,602**	,934**	1	-,167	,931**
	Sig. (2-tailed)	,000	,013	,000	,000	,000	,000		,152	,000
	N	60	60	60	60	75	75	75	75	75
Precipitation	Pearson Correlation	-,336**	-,202	-,216	-,110	-,551 <sup>**</sup>	-,341**	-,167	1	-,200
	Sig. (2-tailed)	,009	,122	,098	,404	,000	,003	,152		,086
	N	60	60	60	60	75	75	75	75	75
SST	Pearson Correlation	,763**	,272*	,469 <sup>**</sup>	,409**	,672 <sup>**</sup>	,940**	,931**	-,200	1
	Sig. (2-tailed)	,000	,036	,000	,001	,000	,000	,000	,086	
	N	60	60	60	60	75	75	75	75	75

<sup>\*\*.</sup> Correlation is significant at the 0.01 level (2-tailed).

<sup>\*.</sup> Correlation is significant at the 0.05 level (2-tailed).

# Appendix 8 Analysis of variance of Indonesian Sargassum – extraction yield and polyphenol content of soli-liquid extraction

#### **Descriptive Statistics**

Dependent Variable: Yield

Solvent	Algae	Mean	Std. Deviation	N
EtOH75	S. aquifolium	12,2667	,40415	3
	S. ilicifolium	7,4600	1,25012	3
	S. polycystum	12,5333	,50332	3
	Total	10,7533	2,57082	9
MeOH	S. aquifolium	5,7000	1,75214	3
	S. ilicifolium	3,0667	,70238	3
	S. polycystum	8,2667	3,16439	3
	Total	5,6778	2,90937	9
MeOH50	S. aquifolium	15,7667	1,50444	3
	S. ilicifolium	7,2700	,95692	3
	S. polycystum	13,4333	1,92959	3
	Total	12,1567	4,02227	9
Water	S. aquifolium	17,2667	2,19393	3
	S. ilicifolium	14,6667	,76376	3
	S. polycystum	22,7667	1,12398	3
	Total	18,2333	3,80690	9
Total	S. aquifolium	12,7500	4,85171	12
	S. ilicifolium	8,1158	4,42893	12
	S. polycystum	14,2500	5,77101	12
	Total	11,7053	5,57152	36

#### Levene's Test of Equality of Error Variances<sup>a</sup>

Dependent Variable: Yield

F	df1	df2	Sig.
2,310	11	24	,042

Tests the null hypothesis that the error variance of the dependent variable is equal across groups.

a. Design: Intercept + Solvent + Algae + Solvent \*Algae

**Tests of Between-Subjects Effects** 

Dependent Variable: Yield

Source	Type III Sum of Squares	df	Mean Square	F	Sig.
Corrected Model	1028,226 <sup>a</sup>	11	93,475	38,522	,000
Intercept	4932,487	1	4932,487	2032,723	,000
Solvent	720,506	3	240,169	98,976	,000
Algae	245,414	2	122,707	50,569	,000
Solvent * Algae	62,307	6	10,384	4,280	,005
Error	58,237	24	2,427		
Total	6018,951	36			
Corrected Total	1086,463	35			

a. R Squared = ,946 (Adjusted R Squared = ,922)

#### **Estimated Marginal Means**

#### 1. Grand Mean

Dependent Variable: Yield

		95% Confidence Interval		
Mean	Std. Error	Lower Bound	Upper Bound	
11,705	,260	11,169	12,241	

#### 2. Solvent

Dependent Variable: Yield

			95% Confidence Interval		
Solvent	Mean	Std. Error	Lower Bound	Upper Bound	
EtOH75	10,753	,519	9,682	11,825	
MeOH	5,678	,519	4,606	6,749	
MeOH50	12,157	,519	11,085	13,228	
Water	18,233	,519	17,162	19,305	

#### 3. Algae

Dependent Variable: Yield

			95% Confidence Interval		
Algae	Mean	Std. Error	Lower Bound	Upper Bound	
S. aquifolium	12,750	,450	11,822	13,678	
S. ilicifolium	8,116	,450	7,188	9,044	
S. polycystum	14,250	,450	13,322	15,178	

#### 4. Solvent \* Algae

Dependent Variable: Yield

Dependent	variable. Held				
	<del>-</del>			95% Confidence Interval	
Solvent	Algae	Mean	Std. Error	Lower Bound	Upper Bound

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EtOH75	S. aquifolium	12,267	,899	10,410	14,123
	S. ilicifolium	7,460	,899	5,604	9,316
	S. polycystum	12,533	,899	10,677	14,390
MeOH	S. aquifolium	5,700	,899	3,844	7,556
	S. ilicifolium	3,067	,899	1,210	4,923
	S. polycystum	8,267	,899	6,410	10,123
MeOH50	S. aquifolium	15,767	,899	13,910	17,623
	S. ilicifolium	7,270	,899	5,414	9,126
	S. polycystum	13,433	,899	11,577	15,290
Water	S. aquifolium	17,267	,899	15,410	19,123
	S. ilicifolium	14,667	,899	12,810	16,523
	S. polycystum	22,767	,899	20,910	24,623

#### Post Hoc Tests Solvent

#### **Multiple Comparisons**

Dependent Variable: Yield

Tukey HSD

Takey Flob						
					95% Con	
		Mean Difference			Lower	Upper
(I) Solvent	(J) Solvent	(I-J)	Std. Error	Sig.	Bound	Bound
EtOH75	MeOH	5,0756 <sup>*</sup>	,73432	,000	3,0498	7,1013
	MeOH50	-1,4033	,73432	,250	-3,4290	,6224
	Water	-7,4800 <sup>*</sup>	,73432	,000	-9,5057	-5,4543
MeOH	EtOH75	-5,0756 <sup>*</sup>	,73432	,000	-7,1013	-3,0498
	MeOH50	-6,4789 <sup>*</sup>	,73432	,000	-8,5046	-4,4532
	Water	-12,5556 <sup>*</sup>	,73432	,000	-14,5813	-10,5298
MeOH50	EtOH75	1,4033	,73432	,250	-,6224	3,4290
	MeOH	6,4789 <sup>*</sup>	,73432	,000	4,4532	8,5046
	Water	-6,0767 <sup>*</sup>	,73432	,000	-8,1024	-4,0510
Water	EtOH75	7,4800 <sup>*</sup>	,73432	,000	5,4543	9,5057
	MeOH	12,5556 <sup>*</sup>	,73432	,000	10,5298	14,5813
	MeOH50	6,0767 <sup>*</sup>	,73432	,000	4,0510	8,1024

Based on observed means.

The error term is Mean Square(Error) = 2,427.

<sup>\*.</sup> The mean difference is significant at the ,05 level.

### **Homogeneous Subsets**

#### Yield

Tukey HSD<sup>a,b</sup>

		Subset		
Solvent	N	1	2	3
MeOH	9	5,6778		
EtOH75	9		10,7533	
MeOH50	9		12,1567	
Water	9			18,2333
Sig.		1,000	,250	1,000

Means for groups in homogeneous subsets are displayed.

Based on observed means.

The error term is Mean Square(Error) = 2,427.

a. Uses Harmonic Mean Sample Size = 9,000.

b. Alpha = ,05.

### Algae

#### **Multiple Comparisons**

Dependent Variable: Yield

Tukev HSD

Tukey HSD						
		Mean			95% Confide	nce Interval
		Difference			Lower	Upper
(I) Algae	(J) Algae	(I-J)	Std. Error	Sig.	Bound	Bound
S. aquifolium	S. ilicifolium	4,6342 <sup>*</sup>	,63594	,000	3,0460	6,2223
	S. polycystum	-1,5000	,63594	,067	-3,0881	,0881
S. ilicifolium	S. aquifolium	-4,6342 <sup>*</sup>	,63594	,000	-6,2223	-3,0460
	S. polycystum	-6,1342 <sup>*</sup>	,63594	,000	-7,7223	-4,5460
S. polycystum	S. aquifolium	1,5000	,63594	,067	-,0881	3,0881
	S. ilicifolium	6,1342 <sup>*</sup>	,63594	,000	4,5460	7,7223

Based on observed means.

The error term is Mean Square(Error) = 2,427.

<sup>\*.</sup> The mean difference is significant at the ,05 level.

## **Homogeneous Subsets**

#### Yield

Tukey HSD<sup>a,b</sup>

	-		
			Subset
Algae	N	1	2
S. ilicifolium	12	8,1158	
S. aquifolium	12		12,7500
S. polycystum	12		14,2500
Sig.		1,000	,067

Means for groups in homogeneous subsets are displayed.

Based on observed means.

The error term is Mean Square(Error) = 2,427.

a. Uses Harmonic Mean Sample Size = 12,000. b. Alpha = ,05.

#### **Descriptive Statistics**

Dependent Variable: TPC

Solvent	Algae	Mean	Std. Deviation	N
EtOH75	S. aquifolium	3,4367	,33606	3
	S. ilicifolium	3,5667	,23288	3
	S. polycystum	3,9033	1,42732	3
	Total	3,6356	,77110	9
MeOH	S. aquifolium	3,7933	,90007	3
	S. ilicifolium	4,4800	1,43447	3
	S. polycystum	2,8767	,19399	3
	Total	3,7167	1,10076	9
MeOH50	S. aquifolium	2,1767	,11372	3
	S. ilicifolium	3,1300	,16523	3
	S. polycystum	2,3833	,21127	3
	Total	2,5633	,45809	9
Water	S. aquifolium	1,7696	,05977	3
	S. ilicifolium	2,0888	,10726	3
	S. polycystum	1,4358	,01938	3
	Total	1,7647	,28950	9
Total	S. aquifolium	2,7941	,97262	12
	S. ilicifolium	3,3164	1,09449	12
	S. polycystum	2,6498	1,11772	12
	Total	2,9201	1,07274	36

#### Levene's Test of Equality of Error Variances<sup>a</sup>

Dependent Variable: TPC

F	df1	df2	Sig.
5,845	11	24	,000

Tests the null hypothesis that the error variance of the dependent variable is equal across groups.

a. Design: Intercept + Solvent + Algae + Solvent \* Algae

#### **Tests of Between-Subjects Effects**

Dependent Variable: TPC

Source	Type III Sum of Squares	df	Mean Square	F	Sig.
	•				
Corrected Model	29,856 <sup>a</sup>	11	2,714	6,251	,000
Intercept	000.005	_	306,96	700 000	000
	306,965	1	5	706,996	,000
Solvent	23,477	3	7,826	18,024	,000
Algae	2,952	2	1,476	3,399	,050
Solvent * Algae	3,427	6	,571	1,316	,288
Error	10,420	24	,434		
Total	347,242	36			
Corrected Total	40,277	35			

a. R Squared = ,741 (Adjusted R Squared = ,623)

# **Estimated Marginal Means**

#### 1. Grand Mean

Dependent Variable: TPC

		95% Confidence Interval		
Mean	Std. Error	Lower Bound	Upper Bound	
2,920	,110	2,693	3,147	

#### 2. Solvent

Dependent Variable: TPC

Depondent variable. If e					
			95% Confidence Interval		
Solvent	Mean	Std. Error	Lower Bound	Upper Bound	
EtOH75	3,636	,220	3,182	4,089	
MeOH	3,717	,220	3,263	4,170	
MeOH50	2,563	,220	2,110	3,017	
Water	1,765	,220	1,311	2,218	

3. Algae

Dependent Variable: TPC

			95% Confidence Interval		
Algae	Mean	Std. Error	Lower Bound	Upper Bound	
S. aquifolium	2,794	,190	2,401	3,187	
S. ilicifolium	3,316	,190	2,924	3,709	
S. polycystum	2,650	,190	2,257	3,042	

#### 4. Solvent \* Algae

Dependent Variable: TPC

	-			95% Confidence Interval	
Solvent	Algae	Mean	Std. Error	Lower Bound	Upper Bound
EtOH75	S. aquifolium	3,437	,380	2,651	4,222
	S. ilicifolium	3,567	,380	2,781	4,352
	S. polycystum	3,903	,380	3,118	4,689
MeOH	S. aquifolium	3,793	,380	3,008	4,579
	S. ilicifolium	4,480	,380	3,695	5,265
	S. polycystum	2,877	,380	2,091	3,662
MeOH50	S. aquifolium	2,177	,380	1,391	2,962
	S. ilicifolium	3,130	,380	2,345	3,915
	S. polycystum	2,383	,380	1,598	3,169
Water	S. aquifolium	1,770	,380	,984	2,555
	S. ilicifolium	2,089	,380	1,304	2,874
	S. polycystum	1,436	,380	,651	2,221

#### Post Hoc Tests Solvent

#### **Multiple Comparisons**

Dependent Variable: TPC

Tukey HSD

		Mean Difference			95% Confide	ence Interval
(I) Solvent	(J) Solvent	(I-J)	Std. Error	Sig.	Lower Bound	Upper Bound
EtOH75	MeOH	-,0811	,31062	,994	-,9380	,7758
	MeOH50	1,0722*	,31062	,010	,2153	1,9291
	Water	1,8708	,31062	,000	1,0140	2,7277
MeOH	EtOH75	,0811	,31062	,994	-,7758	,9380
	MeOH50	1,1533 <sup>*</sup>	,31062	,006	,2965	2,0102
	Water	1,9519 <sup>*</sup>	,31062	,000	1,0951	2,8088

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MeOH50	EtOH75	-1,0722 <sup>*</sup>	,31062	,010	-1,9291	-,2153
	MeOH	-1,1533 <sup>*</sup>	,31062	,006	-2,0102	-,2965
	Water	,7986	,31062	,074	-,0583	1,6555
Water	EtOH75	-1,8708 <sup>*</sup>	,31062	,000	-2,7277	-1,0140
	MeOH	-1,9519 <sup>*</sup>	,31062	,000	-2,8088	-1,0951
	MeOH50	-,7986	,31062	,074	-1,6555	,0583

<sup>\*.</sup> The mean difference is significant at the .05 level.

#### **Homogeneous Subsets**

**TPC** 

Tukey HSD<sup>a,b</sup>

		Subset		
Solvent	N	1	2	
Water	9	1,7647		
MeOH50	9	2,5633		
EtOH75	9		3,6356	
MeOH	9		3,7167	
Sig.		,074	,994	

Means for groups in homogeneous subsets are

displayed. Based on observed means.

The error term is Mean Square(Error) = ,434.

a. Uses Harmonic Mean Sample Size = 9,000.

b. Alpha = ,05.

#### **Algae**

#### **Multiple Comparisons**

Dependent Variable: TPC

Tukey HSD

		Mean			95% Confide	nce Interval
		Difference			Lower	Upper
(I) Algae	(J) Algae	(I-J)	Std. Error	Sig.	Bound	Bound
S. aquifolium	S. ilicifolium	-,5223	,26901	,149	-1,1941	,1495
	S. polycystum	,1443	,26901	,854	-,5275	,8161
S. ilicifolium	S. aquifolium	,5223	,26901	,149	-,1495	1,1941
	S. polycystum	,6666	,26901	,052	-,0052	1,3383
S. polycystum	S. aquifolium	-,1443	,26901	,854	-,8161	,5275
	S. ilicifolium	-,6666	,26901	,052	-1,3383	,0052

272

Based on observed means.

The error term is Mean Square(Error) = ,434.

## **Homogeneous Subsets**

TPC

Tukey HSD<sup>a,b</sup>

		Subset
Algae	N	1
S. polycystum	12	2,6498
S. aquifolium	12	2,7941
S. ilicifolium	12	3,3164
Sig.		,052

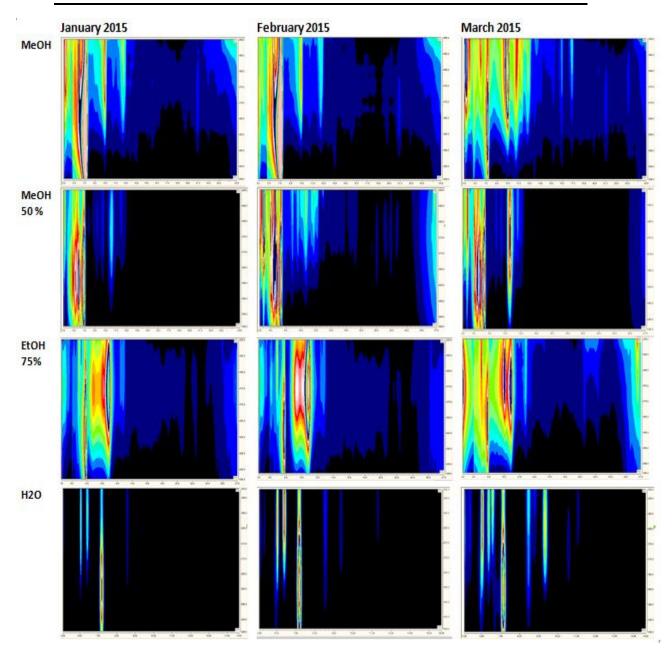
Means for groups in homogeneous subsets are displayed.

Based on observed means. The error term is Mean Square(Error) = ,434.

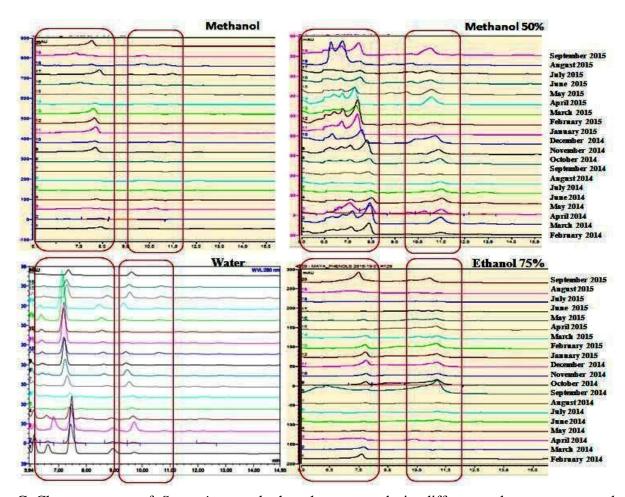
a. Uses Harmonic Mean Sample Size = 12,000.

b. Alpha = ,05.

Appendix 8 3D Image of HPLC chromatogram of *S. muticum* polyphenol compound – seasonal variations in solid-liquid extraction



F. Seasonal variation of *S. muticum* polyphenols in different solvents in 3D image from HPLC analysis. All peaks were detected at wavelength ranging at 260 - 280 nm.



G Chromatogram of S. muticum polyphenol compounds in different solvents – seasonal variation profile

Appendix 9 Analysis of variance of *S. muticum* – screening of enzyme assisted extraction for extraction yield, polyphenol content and bioactivities

Multivariate Tests <sup>a</sup>							
Effect		Value	F	Hypothesis df	Error df	Sig.	
Intercept	Pillai's Trace	1,000	4868,315 <sup>b</sup>	8,000	11,000	,000	
	Wilks' Lambda	,000	4868,315 <sup>b</sup>	8,000	11,000	,000	
	Hotelling's Trace	3540,593	4868,315 <sup>b</sup>	8,000	11,000	,000	
	Roy's Largest Root	3540,593	4868,315 <sup>b</sup>	8,000	11,000	,000	
Sample	Pillai's Trace	4,260	2,562	64,000	144,000	,000	
	Wilks' Lambda	,000	4,766	64,000	69,938	,000	
	Hotelling's Trace	41,087	5,938	64,000	74,000	,000	
	Roy's Largest Root	20,064	45,145 <sup>c</sup>	8,000	18,000	,000	

a. Design: Intercept + Sample

b. Exact statistic

c. The statistic is an upper bound on F that yields a lower bound on the significance level.

Levene's Test of Equality of Error Variances<sup>a</sup>

	F	df1	df2	Sig.
Yield	3,449	8	18	,014
TPC	1,895	8	18	,124
DPPH	6,882	8	18	,000
FRAP	2,967	8	18	,026
Antityrosinase	,706	8	18	,683
Antielastase	3,724	8	18	,010
PAOIBiofilm	5,742	8	18	,001
EcoliBiofilm	2,233	8	18	,075

Tests the null hypothesis that the error variance of the dependent variable is equal across groups.

a. Design: Intercept + Sample

# **Estimated Marginal Means**

1. Grand Mean

			95% Confidence Interval	
Dependent Variable	Mean	Std. Error	Lower Bound	Upper Bound
Yield	29,011	,701	27,538	30,484
TPC	4,282	,121	4,028	4,536
DPPH	1067,722	68,806	923,166	1212,279
FRAP	58,043	,284	57,447	58,638
Antityrosinase	17,103	1,334	14,300	19,906
Antielastase	27,152	,663	25,758	28,546
PAOIBiofilm	13,181	1,768	9,467	16,895
EcoliBiofilm	28,051	3,953	19,746	36,356

2. Sample

	_			95% Confide	ence Interval
Dependent Variable	Sample	Mean	Std. Error	Lower Bound	Upper Bound
Yield	ALC	28,700	2,103	24,282	33,118
	AMG	30,200	2,103	25,782	34,618
	CEL	29,867	2,103	25,448	34,285
	CTRL	26,500	2,103	22,082	30,918
	NEU	32,600	2,103	28,182	37,018
	SHE	26,200	2,103	21,782	30,618
	TER	27,333	2,103	22,915	31,752
	ULT	31,567	2,103	27,148	35,985
	VIS	28,133	2,103	23,715	32,552
TPC	ALC	3,797	,362	3,036	4,558
	AMG	4,808	,362	4,047	5,569
	CEL	4,482	,362	3,720	5,243
	CTRL	2,586	,362	1,825	3,347
	NEU	3,819	,362	3,058	4,581
	SHE	4,686	,362	3,925	5,448
	TER	5,045	,362	4,284	5,806
	ULT	4,163	,362	3,402	4,925
	VIS	5,149	,362	4,388	5,911
DPPH	ALC	1980,082	206,419	1546,412	2413,753
	AMG	728,609	206,419	294,938	1162,279

	CEL	602,167	206,419	168,497	1035,838
	CTRL	720,057	206,419	286,387	1153,728
	NEU	900,175	206,419	466,504	1333,846
	SHE	1020,250	206,419	586,579	1453,920
	TER	810,362	206,419	376,691	1244,033
	ULT	2233,168	206,419	1799,498	2666,839
	VIS	614,631	206,419	180,961	1048,302
FRAP	ALC	52,651	,851	50,863	54,438
	AMG	62,405	,851	60,618	64,193
	CEL	59,783	,851	57,996	61,571
	CTRL	65,926	,851	64,139	67,713
	NEU	59,253	,851	57,466	61,041
	SHE	58,830	,851	57,042	60,617
	TER	57,355	,851	55,567	59,142
	ULT	57,530	,851	55,743	59,317
	VIS	48,650	,851	46,863	50,437
Antityrosinase	ALC	14,032	4,002	5,624	22,441
	AMG	18,759	4,002	10,351	27,168
	CEL	17,874	4,002	9,466	26,282
	CTRL	18,102	4,002	9,694	26,510
	NEU	9,974	4,002	1,565	18,382
	SHE	16,353	4,002	7,945	24,762
	TER	17,500	4,002	9,091	25,908
	ULT	21,131	4,002	12,723	29,539
	VIS	20,201	4,002	11,793	28,610
Antielastase	ALC	30,891	1,990	26,709	35,072
	AMG	21,643	1,990	17,461	25,825
	CEL	29,950	1,990	25,768	34,132
	CTRL	23,115	1,990	18,933	27,297
	NEU	23,422	1,990	19,240	27,604
	SHE	32,880	1,990	28,698	37,062
	TER	28,772	1,990	24,590	32,954
	ULT	24,008	1,990	19,826	28,189
	VIS	29,683	1,990	25,502	33,865
P. aeruginosa Biofilm	ALC	3,492	5,304	-7,650	14,635
	AMG	,000	5,304	-11,143	11,143
	CEL	14,271	5,304	3,129	25,414

	CTRL	-3,553E-15	5,304	-11,143	11,143
	NEU	36,090	5,304	24,947	47,232
	SHE	26,996	5,304	15,854	38,139
	TER	12,441	5,304	1,298	23,583
	ULT	3,678	5,304	-7,465	14,821
	VIS	21,661	5,304	10,518	32,803
<i>E.coli</i> Biofilm	ALC	28,333	11,859	3,418	53,248
	AMG	30,856	11,859	5,941	55,771
	CEL	23,807	11,859	-1,107	48,722
	CTRL	14,691	11,859	-10,224	39,606
	NEU	15,697	11,859	-9,218	40,612
	SHE	64,255	11,859	39,340	89,170
	TER	35,814	11,859	10,900	60,729
	ULT	22,885	11,859	-2,030	47,800
	VIS	16,121	11,859	-8,794	41,036

#### **Post Hoc Tests**

Yield

Tukey HSD<sup>a,b</sup>

Takey Hob		Subset
Sample	N	1
SHE	3	26,2000
CTRL	3	26,5000
TER	3	27,3333
VIS	3	28,1333
ALC	3	28,7000
CEL	3	29,8667
AMG	3	30,2000
ULT	3	31,5667
NEU	3	32,6000
Sig.		,472

Means for groups in homogeneous subsets are displayed.

Based on observed means.

The error term is Mean Square(Error) = 13,269.

a. Uses Harmonic Mean Sample Size = 3,000.b. Alpha = ,05.

#### TPC

Tukey HSD<sup>a,b</sup>

		Subset	
Sample	N	1	2
CTRL	3	2,5859	
ALC	3	3,7972	3,7972
NEU	3	3,8193	3,8193
ULT	3	4,1634	4,1634
CEL	3		4,4817
SHE	3		4,6864
AMG	3		4,8080
TER	3		5,0451
VIS	3		5,1493
Sig.		,112	,237

Means for groups in homogeneous subsets are displayed.

Based on observed means.

The error term is Mean Square(Error) = ,394.

a. Uses Harmonic Mean Sample Size = 3,000.

b. Alpha = ,05.

**DPPH** 

Tukey HSD<sup>a,b</sup>

		Subset		
Sample	N	1	2	3
CEL	3	602,1672		
VIS	3	614,6312		
CTRL	3	720,0574		
AMG	3	728,6087		
TER	3	810,3620		
NEU	3	900,1749		
SHE	3	1020,2496	1020,2496	
ALC	3		1980,0824	1980,0824
ULT	3			2233,1682
Sig.		,871	,076	,992

Means for groups in homogeneous subsets are displayed.

Based on observed means.

The error term is Mean Square(Error) = 127826,654.

- a. Uses Harmonic Mean Sample Size = 3,000.
- b. Alpha = ,05.

FRAP

Tukey HSD<sup>a,b</sup>

		Subset			
Sample	N	1	2	3	4
VIS	3	48,6500			
ALC	3	52,6506			
TER	3		57,3548		
ULT	3		57,5300		
SHE	3		58,8297	58,8297	
NEU	3		59,2533	59,2533	
CEL	3		59,7831	59,7831	
AMG	3			62,4054	62,4054
CTRL	3				65,9260
Sig.		,071	,551	,135	,147

Means for groups in homogeneous subsets are displayed.

Based on observed means.

The error term is Mean Square(Error) = 2,171.

a. Uses Harmonic Mean Sample Size = 3,000.

b. Alpha = ,05.

#### Antityrosinase

Tukey HSD<sup>a,b</sup>

		Subset
Sample	N	1
NEU	3	9,9735
ALC	3	14,0325
SHE	3	16,3534
TER	3	17,4996
CEL	3	17,8740
CTRL	3	18,1022
AMG	3	18,7594
VIS	3	20,2015
ULT	3	21,1311
Sig.		,579

Means for groups in homogeneous subsets are displayed.

Based on observed means.

The error term is Mean

Square(Error) = 48,051.

a. Uses Harmonic Mean Sample

Size = 3,000.

b. Alpha = ,05.

#### Antielastase

Tukey HSD<sup>a,b</sup>

		Subset	
Sample	N	1	2
AMG	3	21,6433	
CTRL	3	23,1150	23,1150
NEU	3	23,4221	23,4221
ULT	3	24,0075	24,0075
TER	3	28,7722	28,7722
VIS	3	29,6834	29,6834
CEL	3	29,9501	29,9501
ALC	3	30,8906	30,8906
SHE	3		32,8801
Sig.		,076	,054

Means for groups in homogeneous subsets are displayed.

Based on observed means.

The error term is Mean Square(Error) = 11,886.

- a. Uses Harmonic Mean Sample Size = 3,000.
- b. Alpha = ,05.

#### **PAOIBiofilm**

Tukey HSD<sup>a,b</sup>

		Subset		
Sample	N	1	2	3
AMG	3	,0000		
CTRL	3	,0000		
ALC	3	3,4924	3,4924	
ULT	3	3,6778	3,6778	
TER	3	12,4405	12,4405	12,4405
CEL	3	14,2713	14,2713	14,2713
VIS	3	21,6607	21,6607	21,6607
SHE	3		26,9964	26,9964
NEU	3			36,0896
Sig.		,157	,101	,098

Means for groups in homogeneous subsets are displayed.

Based on observed means.

The error term is Mean Square(Error) = 84,389.

a. Uses Harmonic Mean Sample Size = 3,000. b. Alpha = ,05.

# Appendix 10 Analysis of variance of Indonesian Sargassum – extraction yield of and polyphenol content of enzyme-assisted extraction

#### **Descriptive Statistics**

Dependent Variable: Yield

Enzyme	Algae	Mean	Std. Deviation	N
Protamex	S. aquifolium	26,2000	5,85918	3
	S. ilicifolium	21,4333	2,05020	3
	S. polycystum	38,1667	6,78994	3
	Total	28,6000	8,76926	9
Viscozyme	S. aquifolium	29,4667	5,40123	3
	S. ilicifolium	27,8667	1,45717	3
	S. polycystum	35,7667	2,65769	3
	Total	31,0333	4,76156	9
Total	S. aquifolium	27,8333	5,34815	6
	S. ilicifolium	24,6500	3,86614	6
	S. polycystum	36,9667	4,79528	6
	Total	29,8167	6,95881	18

#### Levene's Test of Equality of Error Variances<sup>a</sup>

Dependent Variable: Yield

F	df1	df2	Sig.
1,821	5	12	,183

Tests the null hypothesis that the error variance of the

dependent variable is equal across groups.

a. Design: Intercept + Enzyme + Algae + Enzyme \* Algae

#### **Tests of Between-Subjects Effects**

Dependent Variable: Yield

	Type III Sum of	.,		_	ċ
Source	Squares	df	Mean Square	F	Sig.
Corrected Model	577,232°	5	115,446	5,632	,007
Intercept	16002,605	1	16002,605	780,636	,000
Enzyme	26,645	1	26,645	1,300	,276
Algae	490,503	2	245,252	11,964	,001
Enzyme * Algae	60,083	2	30,042	1,465	,269
Error	245,993	12	20,499		
Total	16825,830	18			
Corrected Total	823,225	17			

#### a. R Squared = ,701 (Adjusted R Squared = ,577)

# **Estimated Marginal Means**

#### 1. Grand Mean

Dependent Variable: Yield

		95% Confidence Interval	
Mean	Std. Error	Lower Bound	Upper Bound
29,817	1,067	27,491	32,142

#### 2. Enzyme

Dependent Variable: Yield

			95% Confidence Interval	
Enzyme	Mean	Std. Error	Lower Bound	Upper Bound
Protamex	28,600	1,509	25,312	31,888
Viscozyme	31,033	1,509	27,745	34,322

#### 3. Algae

Dependent Variable: Yield

			95% Confidence Interval		
Algae	Mean	Std. Error	Lower Bound	Upper Bound	
S. aquifolium	27,833	1,848	23,806	31,861	
S. ilicifolium	24,650	1,848	20,623	28,677	
S. polycystum	36,967	1,848	32,939	40,994	

#### 4. Enzyme \* Algae

Dependent Variable: Yield

				95% Confidence Interval	
Enzyme	Algae	Mean	Std. Error	Lower Bound	Upper Bound
Protamex	S. aquifolium	26,200	2,614	20,505	31,895
	S. ilicifolium	21,433	2,614	15,738	27,129
	S. polycystum	38,167	2,614	32,471	43,862
Viscozyme	S. aquifolium	29,467	2,614	23,771	35,162
	S. ilicifolium	27,867	2,614	22,171	33,562
	S. polycystum	35,767	2,614	30,071	41,462

#### Post Hoc Tests Algae

**Multiple Comparisons** 

Dependent Variable: Yield

Tukey HSD

					95% Confidence Interval	
(I) Algae	(J) Algae	Mean Difference (I-J)	Std. Error	Sig.	Lower Bound	Upper Bound
S. aquifolium	S. ilicifolium	3,1833	2,61403	,466	-3,7905	10,1572
	S. polycystum	-9,1333 <sup>*</sup>	2,61403	,011	-16,1072	-2,1595
S. ilicifolium	S. aquifolium	-3,1833	2,61403	,466	-10,1572	3,7905
	S. polycystum	-12,3167 <sup>*</sup>	2,61403	,001	-19,2905	-5,3428
S. polycystum	S. aquifolium	9,1333 <sup>*</sup>	2,61403	,011	2,1595	16,1072
	S. ilicifolium	12,3167 <sup>*</sup>	2,61403	,001	5,3428	19,2905

Based on observed means.

The error term is Mean Square(Error) = 20,499.

#### **Homogeneous Subsets**

#### Yield

Tukey HSD<sup>a,b</sup>

		Subset		
Algae	N	1	2	
S. ilicifolium	6	24,6500		
S. aquifolium	6	27,8333		
S. polycystum	6		36,9667	
Sig.		,466	1,000	

Means for groups in homogeneous subsets are displayed. Based on observed means. The error term is Mean Square(Error) = 20,499.

- a. Uses Harmonic Mean Sample Size = 6,000.
- b. Alpha = ,05.

<sup>\*.</sup> The mean difference is significant at the ,05 level.

### Analysis of Variance Sargassum Indonesia TPC EAE

#### **Descriptive Statistics**

Dependent Variable: TPC

Enzyme	Algae	Mean	Std. Deviation	N
Protamex	S. aquifolium	4,1532	,01163	3
	S. ilicifolium	4,9753	,26753	3
	S. polycystum	6,0753	,13383	3
	Total	5,0679	,84847	9
Viscozyme	S. aquifolium	4,7333	,18356	3
	S. ilicifolium	3,5849	,05365	3
	S. polycystum	4,0328	,43502	3
	Total	4,1170	,55472	9
Total	S. aquifolium	4,4433	,33836	6
	S. ilicifolium	4,2801	,78082	6
	S. polycystum	5,0540	1,15515	6
	Total	4,5925	,85026	18

#### Levene's Test of Equality of Error Variances<sup>a</sup>

Dependent Variable: TPC

F	df1	df2	Sig.
6,970	5	12	,003

Tests the null hypothesis that the error variance of the dependent variable is equal across groups.

a. Design: Intercept + Enzyme + Algae + Enzyme \* Algae

#### **Tests of Between-Subjects Effects**

Dependent Variable: TPC

	Type III Sum of				
Source	Squares	df	Mean Square	F	Sig.
Corrected Model	11,659 <sup>a</sup>	5	2,332	44,355	,000
Intercept	379,635	1	379,635	7221,242	,000
Enzyme	4,069	1	4,069	77,397	,000
Algae	1,997	2	,999	18,995	,000
Enzyme * Algae	5,593	2	2,796	53,193	,000
Error	,631	12	,053		
Total	391,924	18			
Corrected Total	12,290	17			

#### a. R Squared = ,949 (Adjusted R Squared = ,927)

### **Estimated Marginal Means**

#### 1. Grand Mean

Dependent Variable: TPC

F '	1			
		95% Confidence Interval		
Mean	Std. Error	Lower Bound	Upper Bound	
4,592	,054	4,475	4,710	

#### 2. Enzyme

Dependent Variable: TPC

			95% Confidence Interval		
Enzyme	Mean	Std. Error	Lower Bound	Upper Bound	
Protamex	5,068	,076	4,901	5,234	
Viscozyme	4,117	,076	3,951	4,284	

#### 3. Algae

Dependent Variable: TPC

			95% Confidence Interval		
Algae	Mean	Std. Error	Lower Bound	Upper Bound	
S. aquifolium	4,443	,094	4,239	4,647	
S. ilicifolium	4,280	,094	4,076	4,484	
S. polycystum	5,054	,094	4,850	5,258	

#### **Post Hoc Tests**

### Algae

#### **Multiple Comparisons**

Dependent Variable: TPC

Tukey HSD

		Mean			95% Confidence Interval	
(I) Algae	(J) Algae	Difference (I-J)	Std. Error	Sig.	Lower Bound	Upper Bound
S. aquifolium	S. ilicifolium	,1632	,13238	,458	-,1900	,5163
	S. polycystum	-,6108 <sup>*</sup>	,13238	,002	-,9639	-,2576
S. ilicifolium	S. aquifolium	-,1632	,13238	,458	-,5163	,1900
	S. polycystum	-,7739 <sup>*</sup>	,13238	,000	-1,1271	-,4208
S. polycystum	S. aquifolium	,6108 <sup>*</sup>	,13238	,002	,2576	,9639
	S. ilicifolium	,7739 <sup>*</sup>	,13238	,000	,4208	1,1271

Based on observed means.

The error term is Mean Square(Error) = ,053.

 $^{\ast}.$  The mean difference is significant at the ,05 level.

#### **Homogeneous Subsets**

#### TPC

Tukey HSD<sup>a,b</sup>

		Subset		
Algae	N	1	2	
S. ilicifolium	6	4,2801		
S. aquifolium	6	4,4433		
S. polycystum	6		5,0540	
Sig.		,458	1,000	

Means for groups in homogeneous subsets are displayed.

Based on observed means.

The error term is Mean Square(Error) = ,053.

- a. Uses Harmonic Mean Sample Size = 6,000.
- b. Alpha = ,05.

# Appendix 11 Analysis of variance of Indonesian *Sargassum* polyphenol – antioxidant of solid-liquid and enzyme-assisted extraction

#### **Tests of Between-Subjects Effects**

Dependent Variable: DPPH

Source	Type III Sum of Squares	df	Mean Square	F	Sig.
Corrected Model	2558046971,177ª	17	150473351,246	2,677	,006
Intercept	1967350126,996	1	1967350126,99 6	34,994	,000
Sample	539919294,348	2	269959647,174	4,802	,014
Solvents	1037452000,309	5	207490400,062	3,691	,008
Sample * Methods	980675676,520	10	98067567,652	1,744	,108
Error	2023883344,430	36	56218981,790		
Total	6549280442,604	54			
Corrected Total	4581930315,608	53			

a. R Squared = ,558 (Adjusted R Squared = ,350)

#### **Estimated Marginal Means**

#### 1. Grand Mean

Dependent Variable: DPPH

		95% Confidence Interval		
Mean	Std. Error	Lower Bound	Upper Bound	
6035,927	1020,339	3966,583	8105,271	

#### 2. Sample

Dependent Variable: DPPH

Dependent Valiable. DFFH					
			95% Confidence Interval		
Sample	Mean	Std. Error	Lower Bound	Upper Bound	
S. aquifolium	2246,843	1767,279	-1337,366	5831,052	
S. ilicifolium	9987,139	1767,279	6402,930	13571,348	
S. polycystum	5873,798	1767,279	2289,589	9458,007	

#### 3. Methods

Dependent Variable: DPPH

Dependent variable	. DEFII		95% Confidence Interval		
Solvents	Mean	Std. Error	Lower Bound	Upper Bound	
EtOH 75%	5125,606	2499,311	56,769	10194,443	
Methanol	1132,185	2499,311	-3936,651	6201,022	
Methanol 50 %	13520,793	2499,311	8451,956	18589,630	

Protamex	10196,928	2499,311	5128,091	15265,765
Viscozyme	3285,777	2499,311	-1783,060	8354,614
Water	2954,270	2499,311	-2114,567	8023,107

4. Sample \* Methods

Dependent Variable: DPPH

Dependent variat	DIE: DPPH			95% Confidence Interval	
Sample	Solvents	Mean	Std. Error	Lower Bound	Upper Bound
S. aquifolium	EtOH 75%	,000	4328,933	-8779,483	8779,483
	Methanol	-9,095E-13	4328,933	-8779,483	8779,483
	Methanol 50 %	3202,615	4328,933	-5576,868	11982,098
	Protamex	1415,633	4328,933	-7363,850	10195,116
	Viscozyme	,000	4328,933	-8779,483	8779,483
	Water	8862,810	4328,933	83,327	17642,293
S. ilicifolium	EtOH 75%	9872,120	4328,933	1092,637	18651,603
	Methanol	3396,556	4328,933	-5382,927	12176,039
	Methanol 50 %	17080,624	4328,933	8301,141	25860,107
	Protamex	19716,200	4328,933	10936,717	28495,683
	Viscozyme	9857,332	4328,933	1077,849	18636,815
	Water	5,457E-12	4328,933	-8779,483	8779,483
S. polycystum	EtOH 75%	5504,697	4328,933	-3274,786	14284,180
	Methanol	5,002E-12	4328,933	-8779,483	8779,483
	Methanol 50 %	20279,139	4328,933	11499,656	29058,622
	Protamex	9458,951	4328,933	679,468	18238,434
	Viscozyme	4,547E-12	4328,933	-8779,483	8779,483
	Water	-9,095E-13	4328,933	-8779,483	8779,483

#### **Post Hoc Tests**

#### **Sample**

#### **Multiple Comparisons**

Dependent Variable: DPPH

Tukey HSD

rancy rice						
					95% Confidence Interval	
		Mean Difference				Upper
(I) Sample	(J) Sample	(I-J)	Std. Error	Sig.	Lower Bound	Bound
S. aquifolium	S. ilicifolium	-7740,2956 <sup>*</sup>	2499,31061	,010	-13849,3533	-1631,2379
	S. polycystum	-3626,9548	2499,31061	,326	-9736,0124	2482,1029
S. ilicifolium	S. aquifolium	7740,2956 <sup>*</sup>	2499,31061	,010	1631,2379	13849,3533
	S. polycystum	4113,3408	2499,31061	,240	-1995,7169	10222,3985
S. polycystum	S. aquifolium	3626,9548	2499,31061	,326	-2482,1029	9736,0124
	S. ilicifolium	-4113,3408	2499,31061	,240	-10222,3985	1995,7169

Based on observed means.

The error term is Mean Square(Error) = 56218981,790.

#### **Homogeneous Subsets**

#### **DPPH**

Tukey HSD<sup>a,b</sup>

		Subset		
Sample	N	1	2	
S. aquifolium	18	2246,8431		
S. polycystum	18	5873,7979	5873,7979	
S. ilicifolium	18		9987,1387	
Sig.		,326	,240	

Means for groups in homogeneous subsets are displayed.

Based on observed means.

The error term is Mean Square(Error) = 56218981,790.

a. Uses Harmonic Mean Sample Size = 18,000.

b. Alpha = ,05.

#### **Methods**

**Multiple Comparisons** 

Dependent Variable: DPPH

<sup>\*.</sup> The mean difference is significant at the ,05 level.

Tukey HSD

Tukey HSD	_	Mean Difference		95% Confidence Interval		ence Interval
(I) Solvents	(J) Methods	(I-J)	Std. Error	Sig.	Lower Bound	Upper Bound
EtOH 75%	Methanol	3993,4204	3534,55896	,866	-6640,5643	14627,4051
	Methanol 50 %	-8395,1870	3534,55896	,192	-19029,1717	2238,7977
	Protamex	-5071,3223	3534,55896	,706	-15705,3070	5562,6624
	Viscozyme	1839,8285	3534,55896	,995	-8794,1562	12473,8132
	Water	2171,3357	3534,55896	,989	-8462,6490	12805,3204
Methanol	EtOH 75%	-3993,4204	3534,55896	,866	-14627,4051	6640,5643
	Methanol 50 %	-12388,6073 <sup>*</sup>	3534,55896	,015	-23022,5920	-1754,6226
	Protamex	-9064,7427	3534,55896	,133	-19698,7274	1569,2420
	Viscozyme	-2153,5918	3534,55896	,990	-12787,5765	8480,3929
	Water	-1822,0847	3534,55896	,995	-12456,0694	8811,9000
Methanol 50	EtOH 75%	8395,1870	3534,55896	,192	-2238,7977	19029,1717
%	Methanol	12388,6073 <sup>*</sup>	3534,55896	,015	1754,6226	23022,5920
	Protamex	3323,8647	3534,55896	,933	-7310,1200	13957,8494
	Viscozyme	10235,0155	3534,55896	,065	-398,9692	20869,0002
	Water	10566,5227	3534,55896	,052	-67,4620	21200,5074
Protamex	EtOH 75%	5071,3223	3534,55896	,706	-5562,6624	15705,3070
	Methanol	9064,7427	3534,55896	,133	-1569,2420	19698,7274
	Methanol 50 %	-3323,8647	3534,55896	,933	-13957,8494	7310,1200
	Viscozyme	6911,1509	3534,55896	,387	-3722,8338	17545,1356
	Water	7242,6580	3534,55896	,336	-3391,3267	17876,6427
Viscozyme	EtOH 75%	-1839,8285	3534,55896	,995	-12473,8132	8794,1562
	Methanol	2153,5918	3534,55896	,990	-8480,3929	12787,5765
	Methanol 50 %	-10235,0155	3534,55896	,065	-20869,0002	398,9692
	Protamex	-6911,1509	3534,55896	,387	-17545,1356	3722,8338
	Water	331,5071	3534,55896	1,000	-10302,4776	10965,4918
Water	EtOH 75%	-2171,3357	3534,55896	,989	-12805,3204	8462,6490
	Methanol	1822,0847	3534,55896	,995	-8811,9000	12456,0694
	Methanol 50 %	-10566,5227	3534,55896	,052	-21200,5074	67,4620
	Protamex	-7242,6580	3534,55896	,336	-17876,6427	3391,3267
	Viscozyme	-331,5071	3534,55896	1,000	-10965,4918	10302,4776

Based on observed means.

The error term is Mean Square(Error) = 56218981,790.

<sup>\*.</sup> The mean difference is significant at the ,05 level.

#### **Homogeneous Subsets**

#### DPPH

Tukey HSD<sup>a,b</sup>

		Subset		
Solvents	N	1	2	
Methanol	9	1132,1855		
Water	9	2954,2701	2954,2701	
Viscozyme	9	3285,7773	3285,7773	
EtOH 75%	9	5125,6058	5125,6058	
Protamex	9	10196,9281	10196,9281	
Methanol 50 %	9		13520,7928	
Sig.		,133	,052	

Means for groups in homogeneous subsets are displayed.

Based on observed means.

The error term is Mean Square(Error) = 56218981,790.

- a. Uses Harmonic Mean Sample Size = 9,000.
- b. Alpha = 05.

#### **FRAP**

#### **Descriptive Statistics**

Dependent Variable: FRAP

Sample	Solvents	Mean	Std. Deviation	N
S. aquifolium	EtOH 75%	14,4353	,29421	3
	Methanol	12,6588	2,54162	3
	Methanol 50 %	22,4471	4,92682	3
	Protamex	16,1725	,24924	3
	Viscozyme	33,1961	2,46124	3
	Water	21,8089	1,72733	3
	Total	20,1198	7,38959	18
S. ilicifolium	EtOH 75%	10,1375	1,24743	3
	Methanol	11,4638	,79286	3
	Methanol 50 %	16,0303	,75736	3
	Protamex	8,6480	,56784	3
	Viscozyme	15,2844	,98845	3
	Water	24,3450	1,58457	3
	Total	14,3182	5,42148	18
S. polycystum	EtOH 75%	13,7451	1,97886	3
	Methanol	12,7451	3,07423	3
	Methanol 50 %	16,9137	2,60771	3
	Protamex	16,3030	,11792	3
	Viscozyme	27,6863	4,90511	3
	Water	46,4706	5,31729	3
	Total	22,3106	12,53955	18
Total	EtOH 75%	12,7726	2,32053	9
	Methanol	12,2893	2,12591	9
	Methanol 50 %	18,4637	4,12110	9
	Protamex	13,7078	3,80838	9
	Viscozyme	25,3889	8,42009	9
	Water	30,8748	12,10221	9
	Total	18,9162	9,43207	54

#### **Tests of Between-Subjects Effects**

Dependent Variable: FRAP

Dependent variable.	INAI				
	Type III Sum of				
Source	Squares	df	Mean Square	F	Sig.

#### Levene's Test of Equality of Error Variances<sup>a</sup>

Dependent Variable: FRAP

F	df1	df2	Sig.	
3,348	17	36	,001	

Tests the null hypothesis that the error variance of the dependent variable is equal across groups.

a. Design: Intercept + Sample + Solvents + Sample

#### **Tests of Between-Subjects Effects**

Dependent Variable: FRAP

Source	Type III Sum of Squares	df	Mean Square	F	Sig.
Corrected Model	4477,074 <sup>a</sup>	17	263,357	39,833	,000
Intercept	19322,399	1	19322,399	2922,554	,000
Sample	614,026	2	307,013	46,436	,000
Solvents	2645,063	5	529,013	80,014	,000
Sample * Solvents	1217,985	10	121,799	18,422	,000
Error	238,013	36	6,611		
Total	24037,486	54			
Corrected Total	4715,088	53			

a. R Squared = ,950 (Adjusted R Squared = ,926)

#### **Estimated Marginal Means**

#### 1. Grand Mean

Dependent Variable: FRAP

		95% Confidence Interval		
Mean	Std. Error	Lower Bound	Upper Bound	
18,916	,350	18,207	19,626	

#### 2. Sample

Dependent Variable: FRAP

			95% Confidence Interval		
Sample	Mean	Std. Error	Lower Bound	Upper Bound	
S. aquifolium	20,120	,606	18,891	21,349	
S. ilicifolium	14,318	,606	13,089	15,547	
S. polycystum	22,311	,606	21,081	23,540	

<sup>\*</sup> Solvents

3. Methods

Dependent Variable: FRAP

			95% Confidence Interval		
Solvents	Mean	Std. Error	Lower Bound	Upper Bound	
EtOH 75%	12,773	,857	11,034	14,511	
Methanol	12,289	,857	10,551	14,028	
Methanol 50 %	18,464	,857	16,725	20,202	
Protamex	13,708	,857	11,970	15,446	
Viscozyme	25,389	,857	23,651	27,127	
Water	30,875	,857	29,137	32,613	

#### 4. Sample \* Methods

Dependent Variable: FRAP

	-			95% Confidence Interval	
Sample	Methods	Mean	Std. Error	Lower Bound	Upper Bound
S. aquifolium	EtOH 75%	14,435	1,485	11,425	17,446
	Methanol	12,659	1,485	9,648	15,670
	Methanol 50 %	22,447	1,485	19,436	25,458
	Protamex	16,173	1,485	13,162	19,183
	Viscozyme	33,196	1,485	30,185	36,207
	Water	21,809	1,485	18,798	24,820
S. ilicifolium	EtOH 75%	10,138	1,485	7,127	13,148
	Methanol	11,464	1,485	8,453	14,475
	Methanol 50 %	16,030	1,485	13,020	19,041
	Protamex	8,648	1,485	5,637	11,659
	Viscozyme	15,284	1,485	12,274	18,295
	Water	24,345	1,485	21,334	27,356
S. polycystum	EtOH 75%	13,745	1,485	10,734	16,756
	Methanol	12,745	1,485	9,734	15,756
	Methanol 50 %	16,914	1,485	13,903	19,924
	Protamex	16,303	1,485	13,292	19,314
	Viscozyme	27,686	1,485	24,676	30,697
	Water	46,471	1,485	43,460	49,481

# Post Hoc Tests Sample

#### **Multiple Comparisons**

Dependent Variable: FRAP

Tukey HSD

•	-	Mean Difference			95% Confide	ence Interval
(I) Sample	(J) Sample	(I-J)	Std. Error	Sig.	Lower Bound	Upper Bound
S. aquifolium	S. ilicifolium	5,8016 <sup>*</sup>	,85709	,000	3,7066	7,8966
	S. polycystum	-2,1909 <sup>*</sup>	,85709	,039	-4,2859	-,0959
S. ilicifolium	S. aquifolium	-5,8016 <sup>*</sup>	,85709	,000	-7,8966	-3,7066
	S. polycystum	-7,9925 <sup>*</sup>	,85709	,000	-10,0874	-5,8975
S. polycystum	S. aquifolium	2,1909 <sup>*</sup>	,85709	,039	,0959	4,2859
	S. ilicifolium	7,9925*	,85709	,000	5,8975	10,0874

Based on observed means.

The error term is Mean Square(Error) = 6,611.

#### **Homogeneous Subsets**

FRAP

Tukey HSD<sup>a,b</sup>

		Subset		
Sample	N	1	2	3
S. ilicifolium	18	14,3182		
S. aquifolium	18		20,1198	
S. polycystum	18			22,3106
Sig.		1,000	1,000	1,000

Means for groups in homogeneous subsets are displayed.

Based on observed means.

The error term is Mean Square(Error) = 6,611.

a. Uses Harmonic Mean Sample Size = 18,000.

b. Alpha = ,05.

<sup>\*.</sup> The mean difference is significant at the ,05 level.

#### Methods

#### **Multiple Comparisons**

Dependent Variable: FRAP

Tukey HSD

•	-	Mean	Std.		95% Confid	ence Interval
(I) Solvents	(J) Methods	Difference (I-J)	Error	Sig.	Lower Bound	Upper Bound
EtOH 75%	Methanol	,4834	1,21211	,999	-3,1634	4,1301
	Methanol 50 %	-5,6911 <sup>*</sup>	1,21211	,001	-9,3378	-2,0443
	Protamex	-,9352	1,21211	,971	-4,5820	2,7115
	Viscozyme	-12,6163 <sup>*</sup>	1,21211	,000	-16,2630	-8,9695
	Water	-18,1022 <sup>*</sup>	1,21211	,000	-21,7489	-14,4555
Methanol	EtOH 75%	-,4834	1,21211	,999	-4,1301	3,1634
	Methanol 50 %	-6,1744 <sup>*</sup>	1,21211	,000	-9,8212	-2,5277
	Protamex	-1,4186	1,21211	,848	-5,0653	2,2281
	Viscozyme	-13,0996 <sup>*</sup>	1,21211	,000	-16,7464	-9,4529
	Water	-18,5856 <sup>*</sup>	1,21211	,000	-22,2323	-14,9388
Methanol 50	EtOH 75%	5,6911 <sup>*</sup>	1,21211	,001	2,0443	9,3378
%	Methanol	6,1744 <sup>*</sup>	1,21211	,000	2,5277	9,8212
	Protamex	4,7559 <sup>*</sup>	1,21211	,005	1,1091	8,4026
	Viscozyme	-6,9252 <sup>*</sup>	1,21211	,000	-10,5719	-3,2785
	Water	-12,4111 <sup>*</sup>	1,21211	,000	-16,0578	-8,7644
Protamex	EtOH 75%	,9352	1,21211	,971	-2,7115	4,5820
	Methanol	1,4186	1,21211	,848	-2,2281	5,0653
	Methanol 50 %	-4,7559 <sup>*</sup>	1,21211	,005	-8,4026	-1,1091
	Viscozyme	-11,6811 <sup>*</sup>	1,21211	,000	-15,3278	-8,0343
	Water	-17,1670 <sup>*</sup>	1,21211	,000	-20,8137	-13,5202
Viscozyme	EtOH 75%	12,6163 <sup>*</sup>	1,21211	,000	8,9695	16,2630
	Methanol	13,0996 <sup>*</sup>	1,21211	,000	9,4529	16,7464
	Methanol 50 %	6,9252 <sup>*</sup>	1,21211	,000	3,2785	10,5719
	Protamex	11,6811 <sup>*</sup>	1,21211	,000	8,0343	15,3278
	Water	-5,4859 <sup>*</sup>	1,21211	,001	-9,1326	-1,8392
Water	EtOH 75%	18,1022 <sup>*</sup>	1,21211	,000	14,4555	21,7489
	Methanol	18,5856 <sup>*</sup>	1,21211	,000	14,9388	22,2323
	Methanol 50 %	12,4111 <sup>*</sup>	1,21211	,000	8,7644	16,0578
	Protamex	17,1670 <sup>*</sup>	1,21211	,000	13,5202	20,8137
	Viscozyme	5,4859 <sup>*</sup>	1,21211	,001	1,8392	9,1326

Based on observed means.

The error term is Mean Square(Error) = 6,611.

\*. The mean difference is significant at the ,05 level.

#### **Homogeneous Subsets**

FRAP

Tukey HSD<sup>a,b</sup>

		Subset				
Methods	N	1	2	3	4	
Methanol	9	12,2893				
EtOH 75%	9	12,7726				
Protamex	9	13,7078				
Methanol 50 %	9		18,4637			
Viscozyme	9			25,3889		
Water	9				30,8748	
Sig.		,848,	1,000	1,000	1,000	

Means for groups in homogeneous subsets are displayed.

Based on observed means.

The error term is Mean Square(Error) = 6,611.

a. Uses Harmonic Mean Sample Size = 9,000.

b. Alpha = ,05.

Appendix 12 Analysis of variance of Indonesia *Sargassum* polyphenol – biofilm inhibition activity of enzyme-assisted extraction

#### **Descriptive Statistics**

Dependent Variable: Antibiofilm

Bacteria	Species	EAE	Mean	Std. Deviation	N
Bacillus subtilis	S. aquifolium	Protamex	44,4620	11,00078	3
		Viscozym	42,6692	9,59290	3
		Total	43,5656	9,28335	6
	S. ilicifolium	Protamex	48,9907	6,81602	3
		Viscozym	25,2466	14,84923	3
		Total	37,1187	16,61076	6
	S. polycystum	Protamex	40,7081	13,48363	3
		Viscozym	36,8840	12,85987	3
		Total	38,7960	11,96917	6
	Total	Protamex	44,7203	10,01106	9
		Viscozym	34,9333	13,36151	9
		Total	39,8268	12,51127	18
Escherichia coli	S. aquifolium	Protamex	9,1169	5,30387	3
		Viscozym	26,3226	10,98586	3
		Total	17,7198	12,17942	6
	S. ilicifolium	Protamex	18,1880	3,14442	3
		Viscozym	19,9577	11,81075	3
		Total	19,0729	7,79051	6
	S. polycystum	Protamex	45,8097	3,80399	3
		Viscozym	14,4961	4,36150	3
		Total	30,1529	17,53736	6
	Total	Protamex	24,3716	16,94320	9
		Viscozym	20,2588	9,80190	9
		Total	22,3152	13,59349	18

Pseudomonas	S. aquifolium	Protamex	16,3049	4,47868	3
aeruginosa		Viscozym	11,6177	2,30003	3
		Total	13,9613	4,09029	6
	S. ilicifolium	Protamex	9,9637	4,75706	3
		Viscozym	9,0844	2,98026	3
		Total	9,5240	3,58282	6
	S. polycystum	Protamex	12,0483	2,65659	3
		Viscozym	11,4821	6,07117	3
		Total	11,7652	4,20272	6
	Total	Protamex	12,7723	4,50231	9
		Viscozym	10,7280	3,77902	9
		Total	11,7502	4,16723	18
Total	S. aquifolium	Protamex	23,2946	17,43621	9
		Viscozym	26,8698	15,34457	9
		Total	25,0822	16,03918	18
	S. ilicifolium	Protamex	25,7141	18,36269	9
		Viscozym	18,0962	11,96439	9
		Total	21,9052	15,53710	18
	S. polycystum	Protamex	32,8554	17,29853	9
		Viscozym	20,9541	14,13361	9
		Total	26,9047	16,50199	18
	Total	Protamex	27,2880	17,50645	27
		Viscozym	21,9734	13,84989	27
		Total	24,6307	15,86321	54

#### Levene's Test of Equality of Error Variances<sup>a</sup>

Dependent Variable: Antibiofilm

F	df1	df2	Sig.
2,118	17	36	,029

Tests the null hypothesis that the error variance of the dependent variable is equal across groups.

a. Design: Intercept + Bacteria + Species + EAE +Bacteria \* Species + Bacteria \* EAE + Species \*

Bactoria Opocios i Bactoria Ente i Opocios

EAE + Bacteria \* Species \* EAE

#### **Tests of Between-Subjects Effects**

Dependent Variable: Antibiofilm

Dependent Variable. Antibioni	Type III Sum of				
Source	Squares	df	Mean Square	F	Sig.
Corrected Model	10817,682 <sup>a</sup>	17	636,334	9,093	,000
Intercept	32760,262	1	32760,262	468,132	,000
Bacteria	7239,420	2	3619,710	51,724	,000
Species	230,461	2	115,231	1,647	,207
EAE	381,315	1	381,315	5,449	,025
Bacteria * Species	521,218	4	130,305	1,862	,138
Bacteria * EAE	144,639	2	72,320	1,033	,366
Species * EAE	574,733	2	287,367	4,106	,025
Bacteria * Species * EAE	1725,894	4	431,474	6,166	,001
Error	2519,309	36	69,981		
Total	46097,253	54			
Corrected Total	13336,991	53			

a. R Squared = ,811 (Adjusted R Squared = ,722)

## **Estimated Marginal Means**

#### 1. Grand Mean

Dependent Variable: Antibiofilm

		95% Confidence Interval		
Mean	Std. Error	Lower Bound	Upper Bound	
24,631	1,138	22,322	26,939	

#### 2. Bacteria

Dependent Variable: Antibiofilm

			95% Confidence Interval	
Bacteria	Mean	Std. Error	Lower Bound	Upper Bound
Bacillus subtilis	39,827	1,972	35,828	43,826
Escherichia coli	22,315	1,972	18,316	26,314
Pseudomonas aeruginosa	11,750	1,972	7,751	15,749

#### 3. Species

Dependent Variable: Antibiofilm

			95% Confidence Interval	
Species	Mean	Std. Error	Lower Bound	Upper Bound

S. aquifolium	25,082	1,972	21,083	29,081
S. ilicifolium	21,905	1,972	17,906	25,904
S. polycystum	26,905	1,972	22,906	30,904

#### 4. EAE

Dependent Variable: Antibiofilm

			95% Confidence Interval		
EAE	Mean	Std. Error	Lower Bound	Upper Bound	
Protamex	27,288	1,610	24,023	30,553	
Viscozym	21,973	1,610	18,708	25,238	

#### 5. Bacteria \* Species

Dependent Variable: Antibiofilm

Dependent variable. Antibio	111111			:	
				95% Confide	ence Interval
Bacteria	Species	Mean	Std. Error	Lower Bound	Upper Bound
Bacillus subtilis	S. aquifolium	43,566	3,415	36,639	50,492
	S. ilicifolium	37,119	3,415	30,192	44,045
	S. polycystum	38,796	3,415	31,870	45,722
Escherichia coli	S. aquifolium	17,720	3,415	10,793	24,646
	S. ilicifolium	19,073	3,415	12,147	25,999
	S. polycystum	30,153	3,415	23,227	37,079
Pseudomonas aeruginosa	S. aquifolium	13,961	3,415	7,035	20,888
	S. ilicifolium	9,524	3,415	2,598	16,450
	S. polycystum	11,765	3,415	4,839	18,692

#### 6. Bacteria \* EAE

Dependent Variable: Antibiofilm

	Ē			95% Confidence Interval	
Bacteria	EAE	Mean	Std. Error	Lower Bound	Upper Bound
Bacillus subtilis	Protamex	44,720	2,788	39,065	50,376
	Viscozym	34,933	2,788	29,278	40,589
Escherichia coli	Protamex	24,372	2,788	18,716	30,027
	Viscozym	20,259	2,788	14,604	25,914
Pseudomonas aeruginosa	Protamex	12,772	2,788	7,117	18,428
	Viscozym	10,728	2,788	5,073	16,383

7. Species \* EAE

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Dependent Variable: Antibiofilm

	-			95% Confidence Interval	
Species	EAE	Mean	Std. Error	Lower Bound	Upper Bound
S. aquifolium	Protamex	23,295	2,788	17,639	28,950
	Viscozym	26,870	2,788	21,215	32,525
S. ilicifolium	Protamex	25,714	2,788	20,059	31,369
	Viscozym	18,096	2,788	12,441	23,752
S. polycystum	Protamex	32,855	2,788	27,200	38,511
	Viscozym	20,954	2,788	15,299	26,609

#### 8. Bacteria \* Species \* EAE

Dependent Variable: Antibiofilm

	_	-			95% Confide	ence Interval
_					Lower	Upper
Bacteria	Species	EAE	Mean	Std. Error	Bound	Bound
Bacillus subtilis	S. aquifolium	Protamex	44,462	4,830	34,667	54,257
		Viscozym	42,669	4,830	32,874	52,464
	S. ilicifolium	Protamex	48,991	4,830	39,195	58,786
		Viscozym	25,247	4,830	15,451	35,042
	S. polycystum	Protamex	40,708	4,830	30,913	50,503
		Viscozym	36,884	4,830	27,089	46,679
Escherichia coli	S. aquifolium	Protamex	9,117	4,830	-,678	18,912
		Viscozym	26,323	4,830	16,527	36,118
	S. ilicifolium	Protamex	18,188	4,830	8,393	27,983
		Viscozym	19,958	4,830	10,162	29,753
	S. polycystum	Protamex	45,810	4,830	36,014	55,605
		Viscozym	14,496	4,830	4,701	24,291
Pseudomonas	S. aquifolium	Protamex	16,305	4,830	6,510	26,100
aerugino		Viscozym	11,618	4,830	1,822	21,413
	S. ilicifolium	Protamex	9,964	4,830	,168	19,759
		Viscozym	9,084	4,830	-,711	18,880
	S. polycystum	Protamex	12,048	4,830	2,253	21,844
		Viscozym	11,482	4,830	1,687	21,277

#### Post Hoc Tests Bacteria

#### **Multiple Comparisons**

Dependent Variable: Antibiofilm

Tukey HSD

		Mean			95% Con Inter	
(I) Bacteria	(J) Bacteria	Difference (I-J)	Std. Error	Sig.	Lower Bound	Upper Bound
Bacillus subtilis	Escherichia coli	17,5116 <sup>*</sup>	2,78848	,000	10,6957	24,3275
	Pseudomonas aeruginosa	28,0766 <sup>*</sup>	2,78848	,000	21,2607	34,8925
Escherichia coli	Bacillus subtilis	-17,5116 <sup>*</sup>	2,78848	,000	-24,3275	-10,6957
	Pseudomonas aeruginosq	10,5650 <sup>*</sup>	2,78848	,002	3,7491	17,3809
Pseudomonas	Bacillus subtilis	-28,0766 <sup>*</sup>	2,78848	,000	-34,8925	-21,2607
aeruginosa	Escherichia coli	-10,5650 <sup>*</sup>	2,78848	,002	-17,3809	-3,7491

Based on observed means.

The error term is Mean Square(Error) = 69,981.

#### **Homogeneous Subsets**

#### Antibiofilm

Tukey HSD<sup>a,b</sup>

		Subset		
Bacteria	N	1	2	3
Pseudomonas aeruginosa	18	11,7502		
Escherichia coli	18		22,3152	
Bacillus subtilis	18			39,8268
Sig.		1,000	1,000	1,000

Means for groups in homogeneous subsets are displayed.

Based on observed means.

The error term is Mean Square(Error) = 69,981.

a. Uses Harmonic Mean Sample Size = 18,000.

b. Alpha = ,05.

<sup>\*.</sup> The mean difference is significant at the ,05 level.

## **Species**

#### **Multiple Comparisons**

Dependent Variable: Antibiofilm

Tukey HSD

	<u>-</u>				95% Confide	ence Interval
		Mean Difference			Lower	Upper
(I) Species	(J) Species	(I-J)	Std. Error	Sig.	Bound	Bound
S. aquifolium	S. ilicifolium	3,1770	2,78848	,497	-3,6389	9,9929
	S. polycystum	-1,8225	2,78848	,792	-8,6384	4,9934
S. ilicifolium	S. aquifolium	-3,1770	2,78848	,497	-9,9929	3,6389
	S. polycystum	-4,9995	2,78848	,186	-11,8154	1,8164
S. polycystum	S. aquifolium	1,8225	2,78848	,792	-4,9934	8,6384
	S. ilicifolium	4,9995	2,78848	,186	-1,8164	11,8154

Based on observed means.

The error term is Mean Square(Error) = 69,981.

#### **Homogeneous Subsets**

#### Antibiofilm

Tukey HSD<sup>a,b</sup>

		Subset
Species	N	1
S. ilicifolium	18	21,9052
S. aquifolium	18	25,0822
S. polycystum	18	26,9047
Sig.		,186

Means for groups in homogeneous subsets are displayed.

Based on observed means.

The error term is Mean Square(Error) = 69,981.

a. Uses Harmonic Mean Sample Size = 18,000.

b. Alpha = ,05.

#### **Submitted Articles**

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H Index : 73

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Impact factor : 2.372

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H Index : 5

Quartiles :-

Impact factor : 0.32

#### 22ND INTERNATIONAL SEAWEED SYMPOSIUM, COPENHAGEN



# Total phenolic content and biological activities of enzymatic extracts from *Sargassum muticum* (Yendo) Fensholt

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**Abstract** Seaweeds are potentially excellent sources of bioactive metabolites that could represent useful leads in the development of new functional ingredients in pharmaceutical and cosmetic industries. In the last decade, new marine bioprocess technologies have allowed the isolation of substances with biological properties. The brown alga Sargassum muticum (Yendo) Fensholt (Ochrophyta) was enzymatically hydrolyzed to prepare water-soluble extracts by using six different commercially available carbohydratedegrading enzymes and two proteases. Evaluation of 2,2diphenyl-1-picrylhydrazyl (DPPH) and ferric reducing antioxidant power (FRAP) antioxidant, tyrosinase, elastase, and biofilm inhibition, antibacterial and antiviral activities as well as evaluation of cytotoxicity were realized for each extract. Total phenolic content was measured for extract characterization, and solid-phase extraction was useful to purify the enzymatic extract. Soluble total phenolic content of S. muticum Viscozyme extract was highest with 6.4% of dry weight. Enzymatic Celluclast and Viscozyme extracts had the lowest value of DPPH IC50 indicating a strong antiradical activity, 0.6 mg mL<sup>-1</sup>, in comparison with other enzymes. The ferric reducing antioxidant power ranged between 48.7 µM Fe<sup>2+</sup> Eq. digested with Viscozyme, and 60.8 µM Fe<sup>2+</sup> Eq, digested with Amyloglucosidase. Tyrosinase inhibition activity of S. muticum Neutrase extract was 41.3% higher compared to other enzymes. Elastase inhibition activity of S. muticum Shearzyme extract had highest activity (32.8%). All enzymatic extracts showed no cytotoxic effect towards the kidney Vero cells. Meanwhile, only S. muticum Neutrase and Alcalase extracts exhibited potential antiviral activity. In addition, S. muticum Viscozyme and Shearzyme extracts showed promising activity in suppressing the biofilm formation against Pseudomonas aeruginosa and Escherichia coli, respectively. Purification of S. muticum Viscozyme extracts by solid-phase extraction managed to concentrate the phenolic content and improve the bioactivity. These results indicate the promising potential of enzyme-assisted followed by solid-phase extraction in recovering phenolic content and in improving its bioactivity.

**Keywords** Bioactivities · Phaeophyceae · Carbohydrase · Enzyme-assisted extraction · Protease · Solid-phase extraction

#### Introduction

Marine algae have attracted a lot of attention in the last few years. Besides their important ecological role in the ocean, marine algae contain significant quantities of vitamins, minerals, dietary fibers, proteins, polysaccharides, and various functional polyphenols while nutrient contents can vary with species and geographical location, season, and temperature. Algae, rich in nutrition elements of interest, are traditionally consumed as food in East Asian countries, especially in Korea, Japan, China, and Indonesia. Algae contain a large amount of phycocolloids such as agar, alginate, and carrageenan, which have been utilized as gelling agents and emulsifiers in the food and pharmaceutical industries (Liu et al. 2012; Andrade et al. 2013).

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Over the last few years, there have been more investigations in the research into new bioactive natural molecules and valuable metabolites from seaweeds having an economic impact. For example, sulfated anionic polysaccharides have been reported to exhibit anti-inflammatory (Mhadhebi et al. 2014), antioxidant (Costa et al. 2011, Chale-Dzul et al. 2015), antiviral (Eom et al. 2015), antimicrobial (Kantachumpoo and Chirapart 2010), and antiproliferative (Lowenthal and Fitton 2015; Ermakova et al. 2016) activities. Proteins also exhibit a potent bioactivity as antioxidant, antityrosinase, antihypertensive, anticoagulant, and antitumor (Suetsuna et al. 2004; Heo et al. 2005; Bondu et al. 2015).

For many years, phenolic compounds from brown algae have been widely studied for their biological activities. Phenolic compounds also called phlorotannins are secondary defense metabolites synthesized during development as components of algal cell walls (Schoenwaelder and Clayton 1999) or as chemical defense in response to abiotic and biotic stress conditions, such as UV radiation, grazing, bacterial infection, and epiphytism, as well as in intra- and interspecific communication (Burtin 2003; Koivikko 2008). Phlorotannins correspond to polymerization of phloroglucinol monomer units containing both phenyl and phenoxy units (Li et al. 2011). Accumulation of these phloroglucinol-based phenolics contributes to the strong antioxidant activity against free radicalmediated oxidation damage (Kang et al. 2012a; Fujii et al. 2013). They also show other potential bioactivity such as a bactericidal, antiviral, and anticancer activity and radio protective and antiallergic effects (Li et al. 2011; Eom et al. 2012; Le Lann et al. 2016).

In the last decades, natural bioactives from plant, fungal, and marine sources are actively investigated as replacements for synthetic antioxidants currently used as food additives. Even if phenolic compounds are sensitive to extraction techniques based on heat or solvent use (Kadam et al. 2013), conventional solid-liquid extraction is frequently used to extract bioactive compounds from algae. There is no standardized protocol for the phenolic compound extraction, as each species produces different chemically structured phenolic compounds. Principal characteristic of solid-liquid extraction method is the utilization of organic toxic solvents and is often time-consuming. As stated by Ibanez et al. (2012), a green chemistry technology concerns and searches for alternatives in order to reduce hazardous chemical exposure to humans and the environment, prevent the waste, and improve the extraction yield. Several methods have appeared, namely supercritical fluid and pressurized liquid extraction and microwave-, ultrasound-, and enzyme-assisted extraction (Michalak and Chojnacka 2014).

Sulfated fucans and alginates encompass the main portion of the wall in brown algae (up to 45% of algal dry weight), while cellulose only accounts for a small fraction (1–8% of algal dry weight). Proteins, phlorotannins, and halide

compounds (i.e., iodide, bromide, and chloride) are additional components in brown algal cell walls (Deniaud-Bouet et al. 2014). The chemical and structural complexity of brown algae cell walls and their heterogeneity have become the main drawbacks during the extraction process (Jeon et al. 2012; Wijesinghe and Jeon 2012). These drawbacks decrease the extraction rate and efficiency. As a consequence, the degradation of algal cell wall must be taken into consideration when choosing an extraction method since it is an important step in the release of active compounds (Wijesinghe and Jeon 2012). Application of enzyme in degrading the cell wall of algae has been considered for the past decades as it has been proven to be effective in improving the yield of active compounds and enhancing the bioactivity (Hardouin et al. 2014; Kulshreshtha et al. 2015; Olivares-Molina and Fernández 2016).

As an invasive species that is widely spread along the European Atlantic coasts (Plouguerné et al. 2006; Tanniou et al. 2014), the brown macroalga *Sargassum muticum* is one of the most readily available Sargassaceae species in Europe and has been remarked as a potential sustainable source of bioactive compounds. The genus *Sargassum* is known to produce interesting phenolic compounds of high molecular weight, 5000 and 14,000 Da (Le Lann et al. 2012), which demonstrated biological activities such as antifouling, antiproliferative, antiangiogenesis, and antioxidant.

The aim of this work is to propose the use of nonselective commercial carbohydrase and protease as a tool to improve the extraction efficiency of phenolic compounds from *S. muticum*. Recovery and characterization of total phenolic content and bioactivities of the water-soluble fractions were determined, and results were discussed in relation to the different enzymes used during extraction. Evaluation of 2,2-diphenyl-1-picrylhydrazyl (DPPH) and ferric reducing antioxidant power (FRAP) antioxidant, tyrosinase, elastase, and biofilm inhibition and antibacterial and antiviral activities as well as evaluation of cytotoxicity were performed in vitro. Solid-phase extraction was conducted to obtain a phlorotannin-enriched fraction.

#### Materials and methods

#### Algal biomass collection

The brown alga *Sargassum muticum* (Yendo) Fensholt was collected in June 2014 from the rocky shore of St. Gildas de Rhuys peninsula, South Brittany, France (47° 30′ 03″ N, 2° 50′ 12″ W). Apical parts of *S. muticum* thalli were cut and placed in closed boxes to avoid direct contact with sunlight. The collected algal materials were rinsed with tap water to remove the epiphytes and other residual sand. They were ground with a Jupiter's Universal grinder (Wernau, Germany) using a 3-mm perforated grill, lyophilized, and the dry algal materials were stored in the dark for further analysis.



#### Enzyme-assisted extraction of S. muticum

Approximately, 6.5% of dry algal material (19.5 g of dry material) was diluted in 300 mL of distilled water and 2.5% (of dry algal material) commercial enzyme was added. The eight enzymes used were two proteases Neutrase and Alcalase and six carbohydrases Ultraflo, Amyloglucosidase (AMG), Shearzyme, Termamyl, Viscozyme, and Celluclast. Extraction without enzyme or aqueous extract served as negative control. Figure 1 illustrates the enzyme-assisted extraction (EAE) of *S. muticum*.

Sample solutions were incubated for 3 h at 50 °C. Prior to filtration, all samples were denatured for 15 min in 90 °C. Soluble enzymatic extracts were frozen and lyophilized for further analysis. Insoluble enzymatic extracts (residual materials) were stored in -20 °C for total phenolic analysis to determine the insoluble phenolic compounds that may not be digested by the enzymes. All *S. muticum* enzymatic extracts were prepared in triplicates.

#### Total phenolic content analysis

Folin-Ciocalteu method is commonly used for measuring the total phenolic content (TPC). This assay is based on the reduction-oxidation (redox) reactions, which are usually considered to be relatively stoichiometric and on the redox potential of the phenolic hydroxyl group (Hagerman and Butler 1989).

Briefly, 0.5 mL of samples or standard (Phloroglucinol) was added to 0.5 mL 95% ethanol. And 2.5 mL  $_{2}$ O was then added, followed by 0.25 mL of Folin-Ciocalteu reagent (50%,  $_{2}$ V/ $_{2}$ ). All samples were agitated and left to stand for 5 min. Finally, 0.5 mL of  $_{2}$ CO $_{3}$  solution (5%) was added. Optical density was read against the blank at 725 nm. Phloroglucinol was used as standard prepared in different concentrations ranging from 0 to 100  $_{1}$ Mg mL $_{2}$ 1 (Chandler and Dodds 1983; Heo et al. 2005; Wang et al. 2010).

#### Free radical scavenging activity

DPPH is a stable free radical. DPPH assay is based on the theory that a hydrogen donor is an antioxidant. It measures compounds that are radical scavengers. DPPH shows a strong absorption maximum at 517 nm. This analysis is based on modified method of Yen and Chen (1995) and Chen et al. (2008).

A series of butylated hydroxytoluene (BHA), butylated hydroxyanisole (BHT), and ascorbic acid solutions—serves as standard—in different concentrations was prepared (2–50 μg mL<sup>-1</sup>). 0.25 mM of DPPH solution was introduced to 100 μL sample solution. Sample solutions of *S. muticum* enzymatic extracts were made in different concentrations by diluting the stock solution in

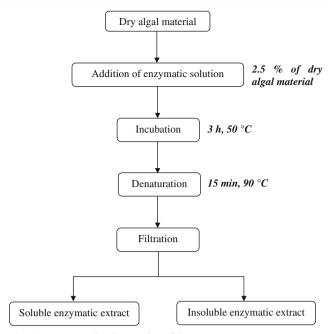


Fig. 1 Enzyme-assisted extraction of S. muticum

methanol (0–1000  $\mu g$  mL<sup>-1</sup>). Before reading the optical density at 517 nm, all samples were incubated at 40 °C for 30 min. Percentage of inhibition was calculated using the following formula:

$$I\left(\%\right) = \left\lceil {^{\left(A_{\rm C} - A_{\rm S}\right)} / _{A_{\rm C}}} \right\rceil \times 100$$

where I (%) = inhibition (expressed in %),  $A_{\rm C}$  = absorbance of control, and  $A_{\rm S}$  = absorbance of samples.

 $IC_{50}$ , corresponding to the concentration sufficient to obtain 50% of a maximum scavenging capacity, of samples was determined based on the regression obtained from dose–response curve.

#### Ferric reducing antioxidant power

This method is based on the reduction of a ferric ion analogue, the complex of tripyridyltriazine Fe(TPTZ)<sup>3+</sup>, to the intensely bluish color complex of Fe(TPTZ)<sup>2+</sup> by the presence of antioxidants in acidic medium. Then, the results were obtained from the absorbance at 593 nm and expressed as micromolar Fe<sup>2+</sup> equivalents (Benzie and Strain 1996, 1999). The TPTZ reagent consisted of 300 mM of acetate buffer with pH 3.6, 10 mM 2,4,6-tripyridyl-s-triazine (TPTZ), and 20 mM FeCl<sub>3</sub>· 6H<sub>2</sub>O solution. Fifty microliter of *S. muticum* enzymatic extract was introduced in a microplate mixed with 150 μL reagent. After 15 min incubation, the absorbance was read at 593 nm. The results were expressed in micromolar equivalent to FeSO<sub>4</sub>·7H<sub>2</sub>O (Shahwar et al. 2012).



#### Tyrosinase inhibition activity

Inhibition of tyrosinase activity was performed based on Chan et al. (2008) and Lim et al. (2009). Phosphate buffer (80  $\mu L$ ) was introduced to 20  $\mu L$  of S. muticum enzymatic extract (100  $\mu g$  mL $^{-1}$ ) added to 20  $\mu L$  of 50% (v/v) dimethyl sulfoxide (DMSO). Later, 40  $\mu L$  of L-3,4-dihydroxyphenylalanine (L-DOPA) and 40  $\mu L$  of mushroom tyrosinase enzyme were added to complete the reaction. Before reading the absorbance at 475 nm, all microplates were incubated at 37 °C for 30 min. Quercetin and kojic acid served as standards prepared in different concentrations (0.02–0.4%). All reactions were made in triplicate. Percentage of inhibition was calculated using the following formula:

$$I\left(\%\right) = \left\lceil {^{\left(A_{\rm C} - A_{\rm S}\right)}} \middle/_{A_{\rm C}} \right\rceil \times 100$$

where I(%) = inhibition (expressed in %),  $A_C$  = absorbance of control, and  $A_S$  = absorbance of samples.

#### Elastase inhibition activity

As an inhibitor, trypsin inhibitor type II-O was used, prepared in phosphate buffer in final concentration 1 g L<sup>-1</sup>. All tested samples were dissolved in 10% (v/v) DMSO with final concentration same as standard (1 mg mL<sup>-1</sup>). The activity of porcine pancreatic elastase (type I; Sigma Chem. Co) was examined using substrate (N-methoxysuccinyl-ala-ala-proval-p-nitroanilide/N-Suc-(Ala)<sub>3</sub>-nitroanilide), and the release of  $\rho$ -nitroaniline was measured at 405 nm.

The reaction of 20  $\mu$ L samples (1  $\mu$ g mL<sup>-1</sup>) was carried out in 80  $\mu$ L of 100 mM Tris and 500 mM NaCl pH 8, 50  $\mu$ L of 0.4 U mL<sup>-1</sup> elastase solution, and 50  $\mu$ L of 0.64 mM substrate. Elastase inhibition was assessed every 10 min at 25 °C for 40 min. The same reactions were applied for the standard, but the volume of standard introduced in the well ranged from 0 to 100  $\mu$ L ( $C_{\rm f} = 0$ –0.5  $\mu$ g mL<sup>-1</sup>) in order to obtain the calibration curve. All reactions were made in triplicate (Sallenave et al. 1998).

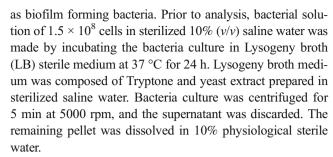
The percentage of inhibition of elastase was calculated as follows:

$$I\left(\%\right) = \left\lceil {^{\left(A_{\rm C} - A_{\rm S}\right)}} \middle/_{A_{\rm C}} \right\rceil \times 100$$

where I(%) = inhibition (expressed in %),  $A_C$  = absorbance of control, and  $A_S$  = absorbance of samples

#### Biofilm inhibition activity

This test is based on the crystal violet assay (Adetunji and Isola 2011; Welch et al. 2012). *Pseudomonas aeruginosa* and *Escherichia coli* (Gram-negative bacteria) were chosen



Bacterial solution (200 µL) was introduced into a sterile 96-well microplate and left for 24 h at 37 °C in order to allow the bacteria to form a film layer in the microplate. After a 24-h incubation, the plate was washed two times with the 10% physiological sterile water. Then, 200 µL of S. muticum enzymatic extracts (1 mg mL<sup>-1</sup>) dissolved in LB was added to the microplate and incubated for 24 h at 37 °C. In the next 24 h, the microplate was rinsed two times with physiological sterile water and 200  $\mu$ L of 10% crystal violet ( $\nu/\nu$ )—prepared in ultrapure water—was added. Fifteen minutes after the crystal violet was added, the microplate was rinsed and the biofilm formed in the microplate was suspended in 200 µL of ethanol. The microplate was read in 600 nm. Negative and positive controls consisted of sterilized saline water and bacteria were applied. Both the control and the samples were made in triplicates.

#### Antibacterial activity

The agar diffusion method (Bauer et al. 1966) was chosen for evaluating the antibacterial activity of *S. muticum* enzymatic extracts. Two Gram-positive bacteria—*Staphylococcus hominis* and *Bacillus subtilis*—as well as two Gram-negative bacteria—*P. aeruginosa* and *E. coli*—were chosen for this test. These bacteria were prepared in LB medium with bacterial density  $1.5 \times 10^8$  cfu mL<sup>-1</sup>. Bacterial suspension was poured on solidified LB agar media and incubated for 1 h at 37 °C.

For screening, *S. muticum* enzymatic extract of 5 mg mL<sup>-1</sup> was prepared in sterile physiological water. Twenty microliters of sample solution—extract solution or positive control (phosphomycin, ampicillin, and streptomycin at 1 mg mL<sup>-1</sup>)—was added to sterile paper discs. All samples were in triplicate. Petri dishes were then incubated at 37 °C for 48 h. The zone of inhibition was established on the agar media and was then measured and expressed in centimeters.

#### Antiviral and cytotoxicity activity

Antiviral activity of *S. muticum* enzymatic extracts and their cytotoxicity were analyzed. For cytotoxicity, the kidney cells of African green monkey (Vero cells) were used. Herpes simplex virus type 1 (HSV-1) model was used to determine the antiviral effect.



One hundred microliter of cell suspension  $(3.5 \times 10^5 \text{ cells mL}^{-1})$  in Eagle's MEM containing 8% FCS was incubated with 50  $\mu$ L of diluted enzymatic extract (concentration from 10 to 500  $\mu$ g mL<sup>-1</sup>) in 96-well plates (72 h, 37 °C, 5% CO<sub>2</sub>). Three replicates were infected using 50  $\mu$ L of medium and a virus suspension at a multiplicity of infection (MOI) of 0.001 ID<sub>50</sub> cells<sup>-1</sup>. After incubation, antiviral activity was evaluated by the neutral red dye method. Zovirax, a commercial antiherpetic acyclovir (9-2(2-hydroxyethoxymethyl) guanine), was used as reference inhibitor (positive control).

For antiviral, 50% effective antiviral concentration (EC<sub>50</sub>) was expressed as the concentration that achieved 50% protection of virus-infected cells from virus-induced destruction. The optical density (OD) was related directly to the percentage of viable cells, which was inversely related to the cytopathic effect (CPE). The linear regression was determined for each assay on the basis of cell controls (0% CPE) and virus controls (100% CPE). Data were expressed as a percentage of protection (%P):

$$\left| \frac{\left( (\mathrm{OD_t})_{\mathrm{virus}} - (\mathrm{OD_c})_{\mathrm{virus}} \right)}{\left( (\mathrm{OD_c})_{\mathrm{MOCK}} - (\mathrm{OD_c})_{\mathrm{virus}} \right)} \right| \times 100$$

where  $(OD_t)_{virus}$  is the OD of the test sample,  $(OD_c)_{virus}$  is the OD of the virus control, and  $(OD_c)_{MOCK}$  is the OD of the mock-infected control (McLaren et al. 1983; Langlois et al. 1986).

For cytotoxicity, 50% cytotoxic concentration (CC<sub>50</sub>) was defined as the concentration that reduced the OD of treated cells to 50% of that of untreated cells. Data were expressed as a percentage of protection (%D):

$$\left\lfloor \frac{\left( (\mathrm{OD_c})_{\mathrm{C}} - (\mathrm{OD_c})_{\mathrm{MOCK}} \right)}{\left( \mathrm{OD_c} \right)_{\mathrm{C}}} \right\rfloor \times 100$$

where (OD<sub>c</sub>)<sub>C</sub> and (OD<sub>c</sub>)<sub>MOCK</sub> are the OD values of untreated cells and treated cells, respectively (Langlois et al. 1986).

#### Solid-phase extraction of S. muticum enzymatic extracts

Solid-phase extraction (SPE) was performed by using C-18 cartridges based on the method of Tanniou et al. (2013), with slight modification. Prior to the fractionation process, the cartridges were conditioned with the following elution: 20 mL methanol and 20 mL ultrapure  $\rm H_2O$  successively. Raw enzymatic extract (4 mL) was loaded into the cartridge and left to absorb the column. The phlorotannins were then eluted with 40 mL methanol 50% ( $\nu/\nu$ ). Prior to elution, the cartridge was washed with 1% acidic water to remove sugars, organic acids, and other water-soluble constituents (Dai and Mumper 2010). All processes were repeated four times in order to accumulate sufficient amount of fractions. The same fractions were pooled and evaporated under reduced pressure at 40 °C on a rotary evaporator. Each fraction was redissolved using acetonitrile-

methanol (2:1; v/v). The phenolic content of methanolic fraction was determined with the Folin-Ciocalteu method, and its antioxidant activity was tested by means of DPPH and FRAP.

#### Fourier transform infrared spectroscopy (FT-IR)

Spectra of *S. muticum* enzymatic extracts were determined using a Nicolet iS 5 FT-IR spectrometer (Thermo Scientific) with diamond crystal plates. The spectra were collected with subsequent scanning of samples in triplicate ranging from 4000 to 400 cm<sup>-1</sup>. Dry enzymatic extracts (2 mg) were introduced onto the crystal plate. The spectra were acquired as the percentage of absorbance. The detected peaks were then compared with a standard of phloroglucinol.

#### High performance liquid chromatography analysis

The chromatographic analysis was performed by UltiMate 3000 Rapid Separation LC (RSLC) System consisting of pump, auto-samplers, and diode array detector (DAD-3000RS) with UV–vis detection (Thermo Scientific). The column was a Nucleodur RP  $C_{18}$  ec (250 × 4.6 mm i.d., 5  $\mu$ m) of Macherey-Nagel. The reversed-phase high performance liquid chromatography (HPLC) analysis was performed based on the modified method of Koivikko et al. (2007) and Sanchez-Camargo et al. (2015). Sargassum muticum enzymatic extracts were eluted by a multistep gradient of solvents A (1 % acid formic in water) and B (acetonitrile). The elution was as follows: 0–20 min, 5 % B; 20–35 min, 20 % B; 35–45 min, 80 % B; and 45–55 min, 0 % B.

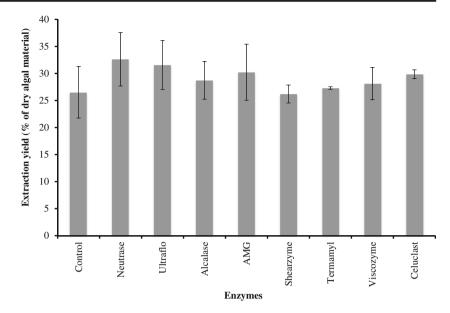
Phenolic compounds were detected at 280 nm, while the wavelength was set ranging from 240 to 300 nm. Injection volume was 100  $\mu$ L with 1 mL min<sup>-1</sup> flow rate. Internal standard range of phloroglucinol was prepared with concentration ranging from 100 to 800  $\mu$ g mL<sup>-1</sup>.

#### Statistical analysis

All results are presented as mean  $\pm$  standard deviation (SD) with n=3. Analysis of variance (ANOVA) was performed using IBM SPSS Statistics 20. Normality and homogeneity tests were performed on all data. Furthermore, the statistical differences between samples were determined using the Tukey test with significance level at 5 % (p < 0.05). Significant differences showed by each sample tested were labeled as different superscript letters. Principal component analysis (PCA) from the FT-IR spectra of *S. muticum* enzymatic extract was performed to transform a number of potentially correlated variables into a number of relatively independent variables that could be ranked based upon their contribution for explaining the variation of the whole data set (Xu and Hagler 2002).



Fig. 2 Yield of enzyme-assisted extraction of *S. muticum* (there is no significant difference between tested samples, p > 0.05). Aqueous extraction was used as control



#### **Results**

#### Yield of enzyme-assisted extraction

After the extraction, the insoluble phase was separated from the soluble aqueous phase. Only the soluble aqueous phase was studied. Extraction yields are presented in Fig. 2. There was no significant difference in the yield of enzyme-assisted extraction (F(8,18)=1.127, p=0.392). In comparison to the aqueous extract ( $26.5\pm4.7\%$  of dry algal material), the enzymatic extracts produced higher yield, except for Shearzyme extract ( $26.2\pm1.7\%$  dry algal material). Neutrase extract yielded  $32.6\pm4.9\%$  of dry algal material, while Alcalase extract only yielded  $28.7\pm3.5\%$  dry algal material. Second

highest yield was obtained from Ultraflo extract,  $31.6 \pm 4.5\%$  dry algal material. The yield of enzyme-assisted extraction from other enzymes, i.e., AMG, Termamyl, Viscozyme, and Celluclast extracts, ranged between  $27.3 \pm 0.2$  and  $30.2 \pm 5.2\%$  of dry algal material.

#### Total phenolic content of S. muticum enzymatic extracts

Insoluble and soluble TPC of *S. muticum* enzymatic extracts were determined, and results were expressed as percent of dry algal material and are presented in Table 1. For insoluble TPC, all enzymatic extracts showed the same content of phenolic content, i.e., 0.9% dry algal material. As for control, there was  $1.0 \pm 0.0\%$  dry algal material (data not shown). Despite the

Table 1 Soluble total phenolic content, DPPH IC<sub>50</sub>, FRAP, tyrosinase, and elastase inhibition activity of S. muticum enzymatic extracts

Samples	Soluble TPC ( % of dry algal material)	DPPH IC <sub>50</sub> (mg mL <sup>-1</sup> )	FRAP (μM Fe <sup>2+</sup> Eq)	Tyrosinase in (% of inhibit	hibition activity	Elastase inhibition	n activity
			5 Vitamin C $58.1 \pm 0.7$	Kojic acid	$67.6 \pm 6.5$	Trypsin inhibitor	$50.0 \pm 0.0$
		BHT $0.012 \pm 4.8$	3	Quecertin	$42.7 \pm 9.4$		
Control	$2.6\pm0.0^a$	$0.7\pm0.2^a$	$65.7 \pm 1.0^{a}$		$14.7\pm3.1^a$		$23.1\pm4.8^{ac}$
Neutrase	$3.8\pm0.0^{ac}$	$0.9\pm0.1^a$	$58.8 \pm 0.6^{bd}$		$6.6\pm1.5^{b}$		$23.4\pm3.3^{ac}$
Ultraflo	$4.2\pm0.0^{ac}$	$2.2\pm0.9^{c}$	$58.4 \pm 0.8^b$		$24.5\pm2.6^c$		$24.0\pm4.2^{ac}$
Alcalase	$3.8\pm0.0^{ac}$	$1.9 \pm 0.3^{bc}$	$56.1 \pm 3.0^{\circ}$		$10.7\pm3.5^a$		$30.9\pm5.9^{ac}$
Celluclast	$4.5\pm0.0^{bc}$	$0.6\pm0.1^a$	$59.3 \pm 1.7^{\text{bd}}$		$19.9 \pm 5.1^a$		$29.9\pm0.5^{ac}$
AMG	$4.8\pm0.0^{bc}$	$0.7\pm0.3^{a}$	$60.8 \pm 1.8^{ad}$		$15.4\pm2.3^a$		$21.6\pm0.8^a$
Shearzyme	$4.7\pm0.0^{bc}$	$1.0\pm0.3^{ab}$	$58.4 \pm 0.7^{bd}$		$13.0\pm0.7^{\rm a}$		$32.8\pm0.9^{bc}$
Viscozyme	$6.4\pm0.5^{bc}$	$0.6\pm0.1^a$	$48.7 \pm 0.9^{c}$		$23.5\pm2.6^{c}$		$29.7\pm1.9^{ac}$
Termamyl	$5.0\pm0.1^{bc}$	$0.8 \pm 0.2^a$	$57.4 \pm 1.0^b$		$14.2\pm2.5^a$		$28.7\pm3.8^{ac}$

Each value represented is mean  $\pm$  standard deviation (n = 3), distinguished by different superscript letters in order to indicate a significant difference (p < 0.05)

 $IC_{50}$  inhibitory concentration 50 %—concentration needed to inhibit 50 % of the reaction, Eq equivalent



low value, this result indicated that there were still some traces of phenolic compound in the residues of *S. muticum* enzymatic extracts.

Statistical analysis showed that there was a significant difference in total phenolic content of S. muticum enzymatic extract (F(8, 18) = 4.097, p = 0.002). Aqueous extracts of S. muticum was significantly different from Celluclast, Shearzyme, AMG, Termamyl, and Viscozyme extracts (p < 0.05). Nevertheless, there was no significant difference exhibited in total phenolic content of aqueous, Alcalase, Neutrase, and Ultraflo extracts of S. muticum (p > 0.05). Neutrase extract (Fig. 2) had the highest yield of dry algal material. However, the highest soluble TPC was shown by Viscozyme extract,  $6.4 \pm 0.5\%$  of dry algal material, followed by Termamyl extract,  $5.0 \pm 0.1$  % of dry algal material. The proteases, i.e., Neutrase and Alcalase, only extracted  $3.8 \pm 0.0$  % of dry algal material. Nevertheless, in comparison with the aqueous extract ( $2.6 \pm 0.0 \%$  of dry algal material), all eight enzymes enhanced the extraction of phenolic compounds in S. muticum.

#### Bioactivities of S. muticum enzymatic extracts

Results for *S. muticum* enzymatic extract bioactivities are presented in Tables 1 and 2. Antioxidant activities of *S. muticum* were determined through the free radical scavenging ability and the reducing power by means of DPPH and FRAP analysis. As summarized in Table 1, certain samples showed a significant difference of DPPH  $IC_{50}$  compared to aqueous extract (F (8, 18) = 8.640, p = 0.000). Ultraflo extract of *S. muticum* was significantly different from Shearzyme, Neutrase, Termamyl, AMG, Celluclast, Viscozyme, and

aqueous extracts (p < 0.05). Alcalase extract also showed a significant difference towards other extracts (p > 0.05) except for Shearzyme and Ultraflo extracts (p > 0.05).

Having low value of  $IC_{50}$  indicates that the sample has a strong antiradical DPPH activity. In this case, Viscozyme and Celluclast extracts  $(0.6 \pm 0.1 \text{ mg mL}^{-1})$  would be mostly effective against the free radical compared to the other. Ultraflo  $(2.2 \pm 0.9 \text{ mg mL}^{-1})$  and Alcalase extracts  $(1.9 \pm 0.3 \text{ mg mL}^{-1})$  were extracts with less effective activity against the free radical as shown by their high value of DPPH  $IC_{50}$ . Nevertheless, free radical scavenging activity of *S. muticum* enzymatic extracts was inferior to the commercial antioxidant BHA and BHT ( $IC_{50} = 0.008 \pm 0.5$  and  $0.012 \pm 4.8 \text{ mg mL}^{-1}$ ).

All samples had significantly different reducing capacity from the aqueous extract (65.7  $\pm$  1.0  $\mu$ M Fe<sup>2+</sup> Eq; F (8, 18) = 35.288, p = 0.000). The most effective reducing capacity was shown by AMG extract in 60.8  $\pm$  1.8  $\mu$ M Fe<sup>2+</sup> Eq. On the contrary, Viscozyme extract did not show a strong reducing capacity with only 48.7  $\pm$  0.9  $\mu$ M Fe<sup>2+</sup> Eq. The reducing power activity of S. *muticum* enzymatic extracts was still inferior to commercial vitamin C, 58.1  $\pm$  0.7  $\mu$ M Fe<sup>2+</sup> Eq. Results are presented in Table 1.

For the tyrosinase inhibition activity of *S. muticum* enzymatic extracts, their activity showed significant difference (F (8, 18) = 16.895, p = 0.000). Neutrase, Ultraflo, and Viscozyme extracts were significantly different from all extracts (p < 0.05). There was no significant difference between aqueous, Alcalase, Celluclast, AMG, Shearzyme, and Termamyl extracts of *S. muticum* (p > 0.05). *S. muticum* enzymatic extract effects on tyrosinase enzyme were still low compared to kojic acid (67.6  $\pm$  6.5 % of inhibition) and

**Table 2** Antiviral and biofilm inhibition activity of *S. muticum* enzymatic extract

Samples	Biofilm inhibition	activity (% of inhibition)	Antiviral activity		
	P. aeruginosa	E. coli		$CC_{50} (\mu g mL^{-1})$	EC <sub>50</sub> (μg mL <sup>-1</sup> )
			Zovirax	>500	$0.7 \pm 0.3$
Control	n.a.	$14.7\pm0.9^a$		>500	>500
Neutrase	$36.1\pm7.8^a$	$29.1\pm10.0^{ac}$		>500	$430.1 \pm 16.3$
Ultraflo	$3.7\pm2.0^{b}$	$32.8\pm3.7^{ac}$		>500	>500
Alcalase	n.a.	$38.0\pm7.9^b$		>500	$225.1 \pm 23.3$
Celluclast	n.a.	$33.8\pm7.7^a$		>500	>500
AMG	n.a.	$43.2\pm5.9^{be}$		>500	>500
Shearzyme	$26.9\pm2.1^b$	$64.3 \pm 10.3^{de}$		>500	>500
Viscozyme	$43.7\pm4.2^a$	$28.8 \pm 9.4^a$		>500	>500
Termamyl	$12.4\pm3.9^b$	$49.2\pm13.3^{be}$		>500	>500

Each value represented is mean  $\pm$  standard deviation (n = 3), distinguished by different superscript letters in order to indicate a significant difference (p < 0.05)

 $CC_{50}$  cytotoxic concentration 50 %—a concentration of product generating 50 % of cell viability compared to control,  $EC_{50}$  antiviral effective concentration—a concentration of product causing 50 % of inhibition of HSV-induced destruction towards the cells, n.a. no activity detected



quercetin (42.7  $\pm$  9.4 % of inhibition)—serving as positive controls. The inhibition activity of *S. muticum* enzymatic extracts ranged from 6.6  $\pm$  1.5 % (Neutrase extract) to 24.5  $\pm$  2.6 % of inhibition (Ultraflo extract).

In elastase inhibition analysis, a significant difference was only showed between the AMG ( $21.6 \pm 0.8$  % of inhibition) and Shearzyme extracts ( $32.8 \pm 0.9$  % of inhibition) (F (8, 18) = 4.231, p = 0.05). It seems that AMG extract has the lowest activity while Shearzyme extract had the highest. All S. muticum enzymatic extracts showed no significant difference with the control ( $23.1 \pm 4.8$  % of inhibition) (p > 0.05). Furthermore, the elastase inhibition activity of S. muticum enzymatic extracts was weaker than that of the trypsin inhibitor ( $50.0 \pm 0.0$  % of inhibition).

For the biofilm generated from *P. aeruginosa*, there was no sign of inhibition from the control, S. muticum enzymatic extracts of Alcalase, Celluclast, and AMG. Ultraflo and Termamyl extract showed weak inhibition with  $3.7 \pm 2.0$ and  $12.4 \pm 3.9$  % of inhibition, respectively. Highest activity was shown by Viscozyme extract with  $43.7 \pm 4.2 \%$  of inhibition. Shearzyme and Termamyl extract (64.3  $\pm$  10.3 and  $49.2 \pm 13.3$  % of inhibition, respectively) performed strong biofilm inhibition activity formed by E. coli. In addition, their activity was significantly better than the aqueous extract,  $14.7 \pm 0.9$  % inhibition. All S. muticum enzymatic extracts were less effective in inhibiting the formation of biofilm formed by P. aeruginosa compared to E. coli, as can be seen in Table 2. Statistical analysis exhibited a significant difference in biofilm inhibition activity of S. muticum against P. aeruginosa and E. coli (F (8, 18) = 5.870, p = 0.01; F (8, 18) = 6.621, p = 0.000).

Based on the antibacterial activity of *S. muticum* enzymatic extracts against four pathogenic bacteria, i.e., S. hominis, B. subtilis, P. aeruginosa, and E. coli, all samples showed no activity (data not shown). It is suggested that the presence of polysaccharides in raw enzymatic extracts provoked the growth of the bacteria instead of suppressing it. On the contrary, phosphomycin, ampicillin, and streptomycin—antibiotics served as positive control—showed substantial antibacterial activity. Inhibition zone formed by phosphomycin in S. hominis, B. subtilis, P. aeruginosa, and E. coli was  $1.9 \pm 0.1$ ,  $1.7 \pm 0.1$ ,  $0.9 \pm 0.1$ , and  $2.2 \pm 0.1$  cm, respectively. Ampicillin showed inhibition zone of 3.6  $\pm$  0.1, 1.6  $\pm$  0.1,  $0.5 \pm 0.0$ , and  $1.7 \pm 0.1$  cm in S. hominis, B. subtilis, P. aeruginosa, and E. coli. As for streptomycin, the inhibition zone formed was  $3.4 \pm 0.1$ ,  $1.1 \pm 0.2$ ,  $2.7 \pm 0.1$ , and  $1.8 \pm 0.2$  cm in S. hominis, B. subtilis, P. aeruginosa, and E. coli, respectively.

No cytotoxicity effect of the extracts on the Vero cells was observed in the range of the concentrations assayed. After 3 days of treatment, no microscopically visible alteration of normal cell morphology was observed even at 200.0  $\mu g \, mL^{-1}$ . As for their potentials as antiviral agent against the HSV-1,

only Neutrase and Alcalase extracts showed a potential antiviral activity with EC  $_{50}$  at 430.1  $\pm$  16.3 and 225.1  $\pm$  23.3  $\mu g$  mL $^{-1}$ .

# Phlorotannin-rich fraction of *S. muticum* enzymatic extract

Semi-purification of *S. muticum* digested with Viscozyme was performed by means of SPE method. The methanolic fraction of Viscozyme extract reached 51.5  $\pm$  0.6 % of dry weight of phenolic content. It showed that there were tenfold phenolic content compared to the raw extract (6.4  $\pm$  0.5 % dry algal material). Antioxidant activity of this fraction was also improved. The value of DPPH IC<sub>50</sub> was 0.012  $\pm$  0.6 from 0.6  $\pm$  0.1 mg mL<sup>-1</sup>. Reducing power capacity of this fraction also improved to 62.4  $\pm$  2.0 from 48.7  $\pm$  0.9  $\mu$ M Fe<sup>2+</sup> Eq.

# FT-IR and HPLC analysis of *S. muticum* Viscozyme extracts

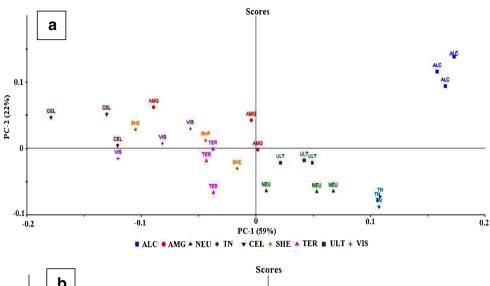
A preliminary analysis to detect the presence of phenolic compound in S. muticum enzymatic extracts was performed via FT-IR analysis. Figure 3 presents the score plot of S. muticum enzymatic extract from the FT-IR spectra: (A) the zone at 1800–600 cm<sup>-1</sup> and (B) wave numbers that are particular to phenolic compounds (3200-3300, 2930-2945, 1605-1620, 1445–1460, 1075–1090, 920–935, and 865–880 cm<sup>-1</sup>). Figure 2b shows a slight of grouping between Neutrase, Alcalase, Ultraflo, and aqueous extracts while the others were dispersed. It was assumed that they probably had similar characteristic of phenolic compounds that might be related to the chemical structure or numbers of polymers. Samples that are close to each other are similar in terms of what the components represent, in this case, the phenolic compounds based on the selected wave numbers. Unfortunately, as there were no additional measurable variables, it would be delicate to make further perception concerning the components influencing this similarity.

Viscozyme extract was chosen between the eight enzymes as this enzyme had the highest phenolic content based on the Folin-Ciocalteu method. FT-IR managed to detect the presence of phenolic compound based on the identified wavelengths as can be seen in Table 3. The liaison of phenolic compound detected from the raw and purified fraction of Viscozyme extract was at 873.72, 930.47, 1081.72, 1450.96, 2934.75, and 3239.54 cm<sup>-1</sup>. These wave numbers were in accordance to those detected in phloroglucinol (standard) and from previous studies. Additional bonds which were not present in the purified fraction were detected at 1325 and 899.15 cm<sup>-1</sup> in raw Viscozyme extract, related to the C=CH<sub>2</sub> and C-H deformation bonds, respectively.

Chromatogram of HPLC from purified fraction of Viscozyme extract is presented in Fig. 4. Based on the



Fig. 3 Principal component analysis of *S. muticum*. **a** From all FT-IR spectra (600–1800 cm<sup>-1</sup>). **b** Phenolic compound of enzymatic extraction from the selected FT-IR spectra (865–880, 920–935, 1075–1090, 1445–1460, 1605–1620, 2930–2945, 3200–3300 cm<sup>-1</sup>). *ALC* Alcalase; *AMG* Amyglucosidase, *NEU* Neutrase; *TN* control/aqueous extract, *CEL* Celuclast, *SHE* Shearzyme, *TER* Termamyl, *ULT* Ultraflo, *VIS* Viscozyme



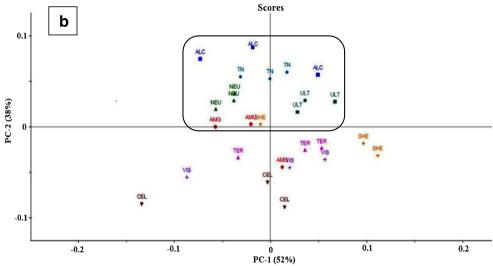


Table 3 Identified wave numbers of Viscozyme extracts (raw and purified sample) correspond to the phenolic compounds

Wave number (cm <sup>-1</sup> )	Bonds	References
873.72 <sup>a</sup>	C–H aromatics	Haeuser et al. (1961), standard (phloroglucinol)
887.07 <sup>b</sup>		
899.15 <sup>a</sup>	C=CH <sub>2</sub>	Karunakaran et al. (2015), Haeuser et al. (1961)
930.47 <sup>a</sup>	Vibration C-O-C	Sekkal et al. (1993)
932.22 <sup>b</sup>		
1081.72 <sup>a</sup>	C-O stretch	Gómez-Ordóñez and Rupérez (2011)
1075.70 <sup>b</sup>		
1325 <sup>a</sup>	C-H deformation	Gómez-Ordóñez and Rupérez (2011), Sartori (1997)
1450.96 <sup>a</sup>	δ CH <sub>2</sub> –CH <sub>3</sub> bend	Sigee et al. (2007)
1493.98 <sup>b</sup>		
1618.11 <sup>a</sup>	C=C cyclic	Kusumaningsih et al. (2016)
1617.51 <sup>b</sup>		
2934.75 <sup>a</sup>	C-H stretch	Ahn et al. (2006), Girija et al. (2013)
2935.76 <sup>b</sup>		
3239.54 <sup>a</sup>	O-H stretch	Standard (phloroglucinol)
3227.27 <sup>b</sup>		

<sup>&</sup>lt;sup>a</sup> Raw hydrolysate



<sup>&</sup>lt;sup>b</sup> Purified hydrolysate

HPLC analysis, Viscozyme crude extract of *S. muticum* contained a compound with the same characteristic as phloroglucinol. This was due to the peak that was detected at the same retention time as phloroglucinols as shown in Fig. 4a Peak of phloroglucinols was detected at 280 nm. Figure 4b presents the detected peak of phloroglucinol-like compound in purified Viscozyme extract. As can be seen from this figure, the peaks were well separated one to each other, indicating that the purification method applied, i.e., SPE, managed to purify the phenolic compounds. In addition, the chromatogram method successfully elucidated the compound.

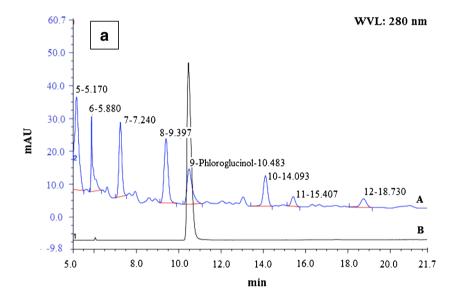
#### **Discussion**

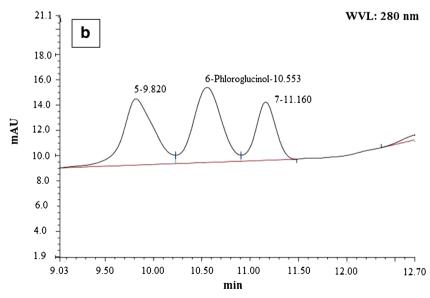
The focus of this study was to evaluate the bioactivity of *S. muticum* phenolic compound obtained after enzymatic-

Fig. 4 Chromatogram of S. muticum from HPLC analysis. a S. muticum Viscozyme raw extract overlaid with phloroglucinol detected at 280 nm. b Purified fraction of S. muticum Viscozyme extract shows a well-separated fraction that is previously assumed to have aromatic rings same as phloroglucinol

assisted hydrolysis. This process has been relatively well-known for the past few years as an alternative method to extract the bioactive compounds (Athukorala et al. 2006; Charoensiddhi et al. 2014; Sanchez-Camargo et al. 2015) from algae. EAE is an environmentally friendly extraction method known by its high efficiency and reduced solvent consumption and time. It has been reported to be effective in releasing the bounded compound, increasing the overall yield (Rosenthal et al. 1996), and removing the barriers of water solubility and insolubility for bioactive compounds (Wijesinghe and Jeon 2012; Kadam et al. 2013). Efficiency of EAE particularly depends on the maintenance of its optimal conditions including time and temperature during the process to maximize the extraction yield (Jeon et al. 2012; Wijesinghe and Jeon 2012).

Eight commercial enzymes consisted of two proteases (Neutrase and Alcalase) and six carbohydrases (Ultraflo,







Celluclast, AMG, Termamyl, Shearzyme, and Viscozyme) were used to release the phenolic compounds from S. muticum. Enzymes used in this study were proven to be effective in increasing the yield of extraction in comparison with water extraction, except for Shearzyme. Furthermore, they have been mostly used in brown algae (Heo et al. 2003; Ahn et al. 2004a; Heo et al. 2005; Athukorala et al. 2006; Kim et al. 2006; Je et al. 2009; Sanchez-Camargo et al. 2015). Highest yield was obtained from S. muticum digested with Neutrase, one of proteolytic enzymes, though the value was not significantly different from the other enzymes and water extraction. Enzymes from the protease group have been reported from several studies as an effective enzyme in enhancing the extraction yield in Ishige okamurae, Sargassum coreanum, Sargassum fulvellum, Sargassum thunbergii, and Scytosiphon lomentaria (Heo et al. 2003; Wang et al. 2010; Hardouin et al. 2014). Nevertheless, the proteolytic enzymes yielding the highest as reported by previous studies were different from the present one; they were Flavourzyme and Alcalase (Heo et al. 2003). Protamex yielded 19% of dry material for S. fulvellum, meanwhile for S. horneri, S. coreanum, and S. thunbergii extracted with Alcalase had 30-40% of extraction yield. In addition, S. coreanum extracted with Protamex contained 9.9 mg g<sup>-1</sup> algal material of phenolic content, while Viscozyme extract had 11 mg g<sup>-1</sup> of algal material. The highest phenolic content in S. horneri, S. fulvellum, and S. thunbergii came from the Alcalase extract with 5.3, 3.6, and 4.16 mg  $g^{-1}$  (Heo et al. 2003). In the present study, the highest phenolic content was also shown by Viscozyme extract of *S. muticum* with almost 6% of phenolic (6 mg g<sup>-1</sup>). Variety in phenolic content obtained from this study and previous one might be due to the different condition of extraction that was applied, i.e., temperature and concentration of enzyme, and also the type of species. It leads to an assumption that the structural heterogeneity and various amounts of algal polysaccharides—alginates, fucoidans, and laminarian in the case of brown algae (Rhein-Knudsen et al. 2015), may influence the efficiency of enzymatic degradation. It is possible that the relative amount and sequence distribution of mannuronic (M) and guluronic (G) residues are two main factors affecting the extraction process since these are the main components in brown algal cell wall constituted as alginates. Mannuronic acids form  $\beta$ -1,4 linkages giving MM blocks a linear and flexible conformation. Meanwhile, guluronic acids generate  $\alpha$ -1,4 linkages and introduce a steric hindrance around the carboxyl groups, thereby providing a folded and rigid structure, ensuring the stiffness in the polymer chain (Draget et al. 2005). M/G ratio of Sargassum alginates varies from 0.8 to 1.5 (Andriamanantoanina and Rinaudo 2010).

Most of the temperatures suggested for the extraction of marine plant phenolic compounds are below 60 °C (Tanniou et al. 2013; de Andrade et al. 2015; Ismail and Abdullah

2015). This is due to phenolic compounds being easily degraded because of high temperature. Nevertheless, there are several studies that tested the extraction of phenolic compound using temperature above 60 °C depending on the method and type of solvents that were applied for the extraction (Siriwardhana et al. 2008; Park et al. 2012; Shitanda et al. 2014). For marine algae, their maceration into a high level of heat might accelerate the extraction process as the heating softens the tissues and facilitate the depolymerization of polysaccharides (Siriwardhana et al. 2008), phenol-protein, and phenol-polysaccharide linkages (Tan et al. 2013). As shown by Siriwardhana et al. (2008), the brown alga Hizikia fusiformis extracted in water at 100 °C for 60 min yielded 4.65 % phenolic content, while at 25, 50, and 75 °C, it was only 1.40, 2.02, and 2.74 % of dry algal material, respectively. An elevated temperature may also help the solvents to reach the sample matrices as the viscosity and the surface tension of the solvents are decreased. In addition, it facilitates the solubility of analytes and promotes higher mass transfer rate (Dai and Mumper 2010). Using high temperature will be mostly compatible with water as solvent, since water becomes a strong source of hydronium (H<sub>3</sub>O<sup>+</sup>) and hydroxide (OH<sup>-</sup>) ions, which can catalyze reactions, including hydrolysis of polysaccharides and proteins into smaller molecules. These molecules are more susceptible to react (Sereewatthanawut et al. 2008). Organic solvents such as methanol, acetone, and many others tend to evaporate at high temperatures (approximately >50 °C); thus, their efficiency decreases due to the shifting of solvent-to-water ratio leading to the change of sample-to-solvent ratio (Tan et al. 2013). As reported by Dai and Mumper (2010), besides temperature, the sample-tosolvent ratio is another one of the crucial factors that influence the yield of chemical extraction.

Antioxidative capacity of phlorotannins, phenolic compounds of brown algae, might also be degraded due to their heat and light sensitivity. The chemistry behind total phenolic assay like Folin-Ciocalteu assay relies on the transfer of electrons from phenolic compounds and other reducing species to molybdenum, forming blue complexes that can be detected spectrophotometrically. It is obvious that other reducing species besides phenolic compounds can also be involved in transferring the required electron (Plaza and Turner 2015). Higher phenolic content from the enzymatic extracts of H. fusiformis result from the study by Siriwardhana et al. (2008) is likely due to the reaction of other molecules, such as polysaccharides and proteins. In addition, these molecules are suggested to contribute in the antioxidant activity of H. fusiformis. Hizikia fusiformis is regarded as a taxonomic synonym of S. fusiforme (Harvey) Hetchell (Stiger et al. 2003). Therefore, it implies that the antioxidant activity in S. muticum enzymatic extract might also be due to the presence of polysaccharides and proteins.



In relation to algae, the introduction of the enzyme followed by heat treatment can further hydrolyze proteins and laminarin (Siriwardhana et al. 2008). Soluble phlorotanninsphenolics of brown algae—are stored in cell organelles, called physodes observed in the cytoplasm of brown algae (Schoenwaelder 2002). It is suggested that phlorotannins transformed into components of algal cell wall when physodes fuse with the cell membrane and the phlorotannins get secreted into the cell wall, complexing with alginic acid (Arnold and Targett 2003). Phlorotannins are also known to form strong complexes with proteins (Stern et al. 1996). As a consequence, enzyme facilitates the breaking of such complex linkages and this process results in liberating more phenolic compound to the solvents. However, it is important to note that high temperature had an adverse impact on the enzymes but the same case applies for the phenolic extraction (Wijesinghe and Jeon 2012). This is due to the instability of phenolic compounds as the elevated temperature stimulates a degradative reaction which is the oxidation of compound resulting in a decrease of phenolic yields (Dai and Mumper 2010). In summary, the temperature of EAE applied in this study was more favorable and suitable for the extraction of phenolic compound. In addition, this method offers a softer condition compared to other methods that apply higher temperatures as previously described.

This study reveals that the phenolic content of S. muticum extracted with carbohydrases was higher compared to the proteases. It leads to an assumption that the carbohydrases disrupt the algal cell wall and release the metabolites linked to the polysaccharides, in this case the phenolic compounds. It is also possible that the phenolic compounds of brown algae are mostly attached to the alginates instead of proteins. As mentioned by previous studies, the phenolic compounds are not only covalently attached to the proteins but also to the alginates (Deniaud-Bouet et al. 2014). The enzymatic degradation of alginate fractions always releases a significant amount of phenolics since phenolic substances form high molecular weight complexes with alginates due to oxidation with haloperoxidases (Bitton et al. 2006). This type of linkage serves as structural component constituting the brown algal cell wall. Phenolic cross-linking of alginate may be an integral part of initial wall formation as shown in the Fucus zygote (Arnold and Targett 2003).

From eight commercial enzymes used, Viscozyme—one of the carbohydrases—yielded the highest phenolic content. Viscozyme was also reported effective in extracting the phenolic content of *Ecklonia cava* (Laminariales, Lessoniaceae) and *S. coreanum* (Heo et al. 2005). The result in this study was also in accordance with previous study conducted by Sanchez-Camargo et al. (2015). They reported that *S. muticum* digested with Viscozyme had a higher content of phenolic compounds compared to Alcalase. However, the efficiency of Viscozyme in extracting the phenolic content from

the brown algae works differently in every species. Siriwardhana et al. (2004) studied the effect of different commercial enzymes in antioxidant activity of *Hizkia fusiformis*. Based on their result, combination of Alcalase and Ultraflo showed the highest content of phenolic compound. Alcalase also enhanced the phenolic content of *S. thunbergii*, *S. horneri*, *S. fullvelum*, and *I. okamurae* (Heo et al. 2005). The difference in the response obtained in this study compared with others might be due to the optimum conditions that were applied (pH, temperature and time, and/or the carbohydrate compositions in different species leading to different mechanism of breakdown).

From the results of the bioactivities assay presented in Tables 1 and 2, it is apparent that the eight enzymatic extracts of *S. muticum* showed different levels of potential. It indicates that the extracted phenolic compound might function differently in responding to the test applied. In addition, it is also important to consider the possibility of an intervention from other compound as in this study the enzymatic extracts tested were all crude material.

The phenolic compounds are compounds with monomeric, oligomeric, or polymeric rings with an aromatic ring having one or more hydroxyl substituents (Balboa et al. 2013). Phlorotannins belonging to the phenolic compounds are exclusively found in brown algae. They have been known as a promising and strong antioxidant agent since they have up to eight interconnected rings of hydroxyl groups (Wang et al. 2009). The phenolic group is known as an excellent hydrogen donor forming hydrogen bonds with the carboxyl group of protein (Ozdal et al. 2013). Later, the existence of these hydroxyl groups provides the molecule with the ability of donating a proton to radical leading to its role as a possible chain breaking molecule or antioxidant upon secondary oxidation (Franco et al. 2008).

Phenolic compounds are one of the compounds that can upregulate the expression of antioxidant enzymes for examples the superoxide dismutase (SOD), catalase (CAT), glutathione peroxidase (GPx), and reductase (GR). The different types of bioactivity chosen in this study were intended for the possibilities of phenolic compounds to be further applied in the nutraceuticals, cosmeceuticals, and painting industries. Therefore, the antioxidant activity of brown algae is strongly related to their phenolic content. As in this study, *S. muticum* digested with Viscozyme had strongest antiradical activity indicated by the low value of IC<sub>50</sub>. Reducing the particle size can increase the extraction rate and the yield of phenolic compounds that can be achieved by grinding and enzyme-assisted extraction (Franco et al. 2008).

S. horneri digested with AMG, Celluclast, Termamyl, Ultraflo, and Viscozyme was reported to show DPPH radical scavenging activities at 25 μg mL<sup>-1</sup> (Park et al. 2004, 2005). Heo et al. (2005) also studied the antioxidant activity via DPPH assay of S. horneri enzymatic extracts.



However, their results showed that the Viscozyme, Celluclast, AMG, and Ultraflo extracts with final concentration 2 mg mL<sup>-1</sup> did not show any inhibition. Instead, the Termamyl extract of *S. horneri* showed 11.2 % inhibition. This activity was poor compared to the commercial standard BHA and BHT with 87 and 56 % inhibition, respectively. The same authors also reported that the antiradical activity of *S. fullvelum*, *S. coreanum*, and *S. thunbergii* extracted with carbohydrate enzymes also showed weak activity ranging from 4 to 38 % compared to the standards. This suggests that there is an interindividual and interspecies variation affecting the performance of antioxidant activity.

The antioxidant effects of S. muticum enzymatic extract might also be due to the presence of polysaccharides. The heterofucans of Sargassum filipendula have been reported to show antioxidant activity such as hydroxyl and superoxide radical scavenging, chelating effects on ferrous ions, and reducing powers in active concentration at 2 mg mL<sup>-1</sup> (Costa et al. 2011). Lim et al. (2014) in their study concerning the antioxidant effects of Sargassum binderi also reported that the antiradical effect of this sample was at 2 mg mL<sup>-1</sup>. It indicated that their antioxidant activity was weaker in comparison to the commercial standard, BHA, BHT, and ascorbic acid with  $6.0 \times 10^{-4}$ ,  $9.6 \times 10^{-4}$ , and  $3.5 \times 10^{-4}$  mg mL<sup>-1</sup>. However, it had performed high superoxide (compared to commercial food-grade fucoidan, BHA, and BHT) and hydroxyl radical scavenging activities (compared to BHT) (Lim et al. 2014). The ability of Sargassum graminifolium polysaccharides to inhibit 50% of superoxide anions scavenging and antiradical activities was at 1.9 and 0.6 mg mL<sup>-1</sup>, respectively (Zhang et al. 2012).

Tyrosinase is a metalloenzyme oxidase catalyzing two distinct reactions of melanin synthesis. Melanin helps to protect skin from the damaging ultraviolet radiation of the sun. However, elevated concentrations of melanin in the skin can result in hyperpigmentation such as freckles and moles (Kang et al. 2012b). While tyrosinase is responsible for skin pigmentation, elastase is a proteinase enzyme capable of degrading elastin. Elastin is an extracellular matrix protein providing elasticity to the connective tissues. It forms elastic fiber in skin dermis having an influence on skin elasticity (Kim et al. 2009; Moon et al. 2010). Therefore, inhibition of tyrosinase and elastase activities become a method to protect against skin aging (Kim et al. 2009; Chang 2012). There is interest in the potential of marine algal bioactive compounds in cosmeceutical industries related to their tyrosinase and elastase inhibition activities. In the case of phlorotannins, many studies have reported their potential in inhibiting the tyrosinase activity. Dieckol, isolated phlorotannins from the brown alga E. cava, has been examined for its inhibitory effects on mushroom tyrosinase. Dieckol was capable to inhibit the mushroom tyrosinase with IC50 20 µM and it showed no cytotoxicity (Kang et al. 2012b). 7-Phloroekol of E. cava has been observed to have an inhibitory effect on mushroom tyrosinase at 50 µg mL<sup>-1</sup>. This compound was isolated from the ethanol extract of *E. cava* (Yoon et al. 2009). These examples provide knowledge concerning the tyrosinase inhibition activity from marine brown algal phlorotannins. Some results presented in the present study are in accordance with previous studies as previously mentioned. Polyphenol extracts of *S. muticum* extracted in different commercial enzymes showed interesting results regarding the tyrosinase and elastase inhibition activities. Attention was given to the tyrosinase inhibition activity; *S. muticum* Neutrase extract demonstrated the highest activity at 41 % compared to other enzymes. Accordingly, this result indicates another promising potential of enzymatic extracts of *S. muticum* polyphenol in the point of view of cosmeceutical industry.

Enzymatic extracts of S. muticum polyphenol did not show expected result against bacteria, as they did not demonstrate any antibacterial activity. It is suspected that the presence of sugar could contribute to the growth of bacteria in the media. However, this assumption will need further verification by evaluating the purified enzymatic extract of Sargassum polyphenol. It has been demonstrated that purified phlorotannin showed potent inhibition of cell membrane integrity as well as cell metabolism against Trichophyton rubrum (Eom et al. 2012). Phlorotannins from Ecklonia kurome were active against food-borne pathogen bacteria. Interaction between protein and brown algal phlorotannin has been considered to play important role in bactericidal action of purified phlorotannins (Ahn et al. 2004b). Even though the enzymatic extract of Sargassum polyphenol did not show antibacterial activity, biofilm inhibition activity demonstrated the opposite result. It seems that polyphenol of Sargassum enzymatic extracts was more active against biofilm formation. This contradictive result is caused by different mechanism of inhibition. In antibacterial activity, the mechanisms of inhibition are most likely thought to rely on inhibition of extracellular microbial enzymes, deprivation of substrates required for microbial growth, or direct inhibition of oxidative phosphorylation (Scalbert 1991). Regarding biofilm inhibition activity, two main mechanisms have been proposed, i.e., direct inhibition of bacterial settlement or survival of the macrofouler larvae and regulation of macrofouler growth, which in turn affects larval settlement of the macrofoulers (Avelin Mary et al. 1993). Some studies have reported the promising potential of brown algal polyphenol compound as antifouling agent. Water extract of S. vulgare polyphenol from its pneumatocysts presented 75 to 99 % inhibition of mussel Perna perna attachment. Polyphenol from leaflets of S. vulgare had 67 to 80 % inhibition of byssal adhesion (Plouguerné et al. 2012). Phlorotannin extract of S. tenerrimum had EC<sub>50</sub> of larval settlement inhibition of Hydroides elegans at 0.526 ppm. It indicated that this



extract was effective in inhibiting the settlement of *H. elegans* larval attachment (Lau and Qian 1997).

Antiviral effects of brown algae are usually shown by their polysaccharides as explained by previous studies (Ahn et al. 2006; Lee et al. 2011; Elizondo-Gonzalez et al. 2012). However, for the past few years, several studies concerning the antiviral effects of brown algae phlorotannins have appeared. Ahn et al. (2004b) reported an inhibitory effect of 8,8'-bieckol and 8-4"-dieckol isolated from E. cava on HIV-1 reverse transcriptase and protease in vitro. Phloroglucinol, eckol, 7-phloroeckol, phlorofucoroeckol, and dieckol isolated from E. cava were effective against porcine epidemic diarrhea coronavirus at concentration ranging from 10.8 to 14.6 µM (Wang et al. 2008). Ethyl acetate fractions of phlorotannins from brown algae Eisenia bicyclis (Lessoniaceae) have been reported to show antiviral effects against genital human papilloma virus (HPV) at 50 µg mL<sup>-1</sup> (Kim and Kwak 2015). Dichloromethane/methanol extract of S. cymosum, S. polyceratium, S. vulgare, Padina gymnospora, Lobophora variegata, Dictyopteris delicatula, Sictyota menstrualis, and Stypodium zonale demonstrated herpes simplex virus-1 (HSV-1) inhibition activity with percentage of inhibition ranging from 20 to 96 % at 400 μg mL<sup>-1</sup>. Spectroscopic analysis of these crude extracts detected the presence of polyphenol compounds along with other active compounds like the halogenated sesquiterpenes obtusol, elatol, meroditerpenoid atomaric acid, triacylglycerols, and fatty acids (Soares et al. 2012). Out of eight, only two enzymatic extracts were found to be effective against the HSV-1; they were S. muticum digested with Neutrase and Alcalase. Result in this study regarding the antiviral effect of S. muticum enzymatic extract seems to be different from the previous studies. Out of eight, only two enzymatic extracts were found to be effective against the HSV-1; they were S. muticum digested with Neutrase and Alcalase. However, the antiviral effects of these two enzymatic extracts might not be provoked by the phlorotannins since the Viscozyme extract of S. muticum with high content of phenolic compound did not show any antiviral activity. It is possible that the presence of other compounds such as protein and polysaccharides that are in the enzymatic extracts is involved in disrupting the viral cells. As part of brown algal cell wall, phlorotannins are known to form a complex linkage with alginates and proteins. Early characterization by FT-IR has confirmed the presence of carbohydrates and proteins in S. muticum raw extracts. Zone 1800-600 cm<sup>-1</sup> that was selected for PCA is the type that is usually used for polysaccharides and protein studies in algae (Rodrigues et al. 2015).

In tyrosinase, elastase, and biofilm inhibition activity as well as antiviral effects of *S. muticum* enzymatic extract, other compounds aside from polyphenol might also contribute to these bioactivities such as polysaccharides and proteins. To our best knowledge, there have not been any reported studies concerning the tyrosinase,

elastase, biofilm inhibition, and antiviral activity of polyphenol from enzymatic extracts.

The semi-purification fraction of *S. muticum* Viscozyme extract yielded an important percentage of phenolic compounds assumed to be phlorotannins. The antiradical and the reducing capacity of the semi-purified fraction was also improved compared to the raw enzymatic extract. This result suggests that the antioxidant activity shown by *S. muticum* enzymatic extract is generated by the presence of phenolic compounds. This suggestion is supported by FT-IR (Table 3) and HPLC analysis (Fig. 3) that detects the presence of peaks corresponding to phloroglucinol.

FT-IR detects other peaks that appear to be correlated with phenolic bonds characterized by its aromatic and hydroxyl groups with the detection of C–H aromatics, C–O–C vibration, C–O stretch, and C=C. The characteristic of phenolic compounds lies on its aromatic ring (1260, 1620, 2970, and 2850–2970 cm<sup>-1</sup>) and hydroxyl group (3415 cm<sup>-1</sup>) (Ganapathi et al. 2013). However, a protein bond is also exposed through this analysis by the presence of amide bond either in the raw or semi-purified enzymatic extracts. On the contrary, the purification method has successfully removed the cellulose and mannuronic acid. Due to the lack of semi-purified extracts, it was impossible to characterize them using mass spectrometry. This will be performed in future work to help to identify the group of phlorotannins that exists in *S. muticum* extracted with enzyme.

This study has demonstrated that the application of enzymes in extracting the phenolic compound in Sargassum is quite effective. Based on the results presented in this study, different enzymes used give various results concerning the total phenolic content and their bioactivity. Among the eight enzymes, Viscozyme releases higher content of phenolic compounds and are correlated with strong antiradical activity. This might be supported by the optimum conditions provided, i.e., 50 °C for 3 h, during the extraction process, which is favorable for the Viscozyme, extract activity. Since phenolic compounds are sensitive to heat and easily oxidized by the exposure of light, it is strongly recommended to apply a mild temperature between 40 to 60 °C. Besides the extraction condition, in the case of phenolic compound, it is important to take the period of collection into further consideration. As commonly known, the production of phenolic compound is mostly stimulated by their environmental condition, such as the presence of grazer and direct exposition to sunlight. As a result, the content will vary according to the season.

Another important note to take is that by using enzymes, the active molecules released might not exclusively come from the phenolic groups. It is possible that it also extracts other molecules since the objective of this method is to degrade algae cell walls. Therefore, it will not be surprising that the fucoidan and protein might be responsible for the



bioactivity of *S. muticum*. As a consequence, it will be interesting to further work on the purified samples pretreated by the enzyme-assisted extraction in the future. In addition, the characterization of *S. muticum* phenolic compound might help in understanding more about the mechanisms of their bioactivity.

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# Antioxidant and antibacterial activity of solid-liquid and enzyme-assisted extraction of phenolic compound from three species of tropical *Sargassum*

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**Abstract.** Sargassum has been well acknowledged for the potential natural product of its phlorotannins. Solid-liquid extraction (SLE) is the most common method used to extract them. However, this method has some drawbacks such as low yield and toxic. An alternative ecofriendly method has been proposed, i.e. enzyme-assisted extraction (EAE), proven to be more efficient. The aim is to compare the efficiency of SLE and EAE concerning their extraction yield, total phenolic content and antioxidant activity. S. aquifolium, S. ilicifolium and S. polycystum were extracted using water, methanol, methanol 50%, and ethanol 75% and enzymes (Viscozyme and Protamex). Total phenolic content (TPC) was analyzed by Folin-Ciocalteu and antioxidant activity via DPPH and FRAP analysis. This study implied that bioactivity of Sargassum extracted with enzymes is better compared to the one using organic solvents.

Keywords: Conventional extraction, alternative extraction, brown algae, bioactivities, polyphenols

#### 1. Introduction

As an archipelagic country located in the tropical zone and the coral triangle, Indonesia has been extensively known for its remarkable diversity of marine organisms. One of them is marine seaweed. There are three groups of marine seaweed that can be found throughout Indonesian water, namely, green, red and brown seaweed. For decades, the green and red seaweed, particularly from the genera *Kappaphycus* and *Gracillaria*, have been cultivated for their carrageenan and agar [1–3].

Marine seaweed is also distinguished by its polysaccharides [4,5] but also by its biologically active compounds that have been studied and proven to have promising potential for the nutraceutical, pharmaceutical and cosmeceutical industries [6–10]. Hence, these marine plants have attracted a lot attention during the exploration of alternative natural products. The natural products derived from seaweed offer a sustainable resource with an infinite application.

Among the other three, brown seaweed from the class of Phaeophyceae is one of marine plants that has not been fully explored and exploited in Indonesia despite of their abundance and their remarkable potential [11–17]. A number of previous studies have reported that polysaccharides and bioactive compounds from the brown seaweed showed an encouraging potency as source of natural product with

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compounds from the brown seaweed showed an encouraging potency as source of natural product with their antimicrobial [18], antioxidant[11,19], antifouling [20,21], antiviral [22,23], anti-proliferative [24–26], anti-inflammatory [27], tyrosinase inhibition [28] activities and many other. This potency can be further applied in foods, medicines and cosmetics in order to reduce the application of chemical products or even to fully substitute them in the future [29–33].

There are five genera of brown seaweed widely distributed in Indonesian waters which four of them are the most species-rich, namely *Dictyota*, *Padina*, *Turbinaria* and *Sargassum* with 111 species in total [34]. These four genera have not been optimally exploited; however, *Sargassum* is the most extensively studied among them concerning its alginates and bioactive compound especially the phenolic compounds also known as phlorotannins. The phlorotannin in which are mostly present in the brown seaweed [35,36] are known for its strong antioxidant activity [37–43]. Nevertheless, it also shows promising effects against cancer, allergy, diabetes, inflammation and viral and microbial infection [36]. Basically, this compound is produced as one of the components composing the cell wall of brown seaweed [35]. However, it also serves as an induced defense system as phlorotannin has plastic responses to the environmental factors for examples nutrient availability, light, ultraviolet radiation, temperature, salinity and the intensity of herbivory [44–47].

Due to the various health beneficial performed by phlorotannin, various extractants are used in order to release this compound from the algal matrix. Solid-liquid extraction (SLE) is one of the most used methods in extracting the phlorotannin of brown seaweed [48–53]. This method relies on the organic solvents during the process such as hexane, ethyl acetate, ethanol, methanol, etc. [52]. The polysaccharides' complexity of marine seaweed often becomes the hindrance during the application of SLE. Thus, it reduces the extraction efficiency causing to the low yield of extraction with only 8% to 30% of the dry material [54]. It is crucial to apply an extraction method that could enhance the target compound, improve the bioactivity, time-saving, and favorable for human and environmental. These are the necessary requirements in order to generate a 'Green concepts' as in recent trend [55].

Enzyme-assisted extraction (EAE) has been widely mentioned as a potential alternative method since it offers an eco-friendly approach, an improvement in the quantity of target compound and an enhancement in the bioactivity [56]. The advance point of EAE compared to SLE method lies on its ability to degrade or disrupt the cell walls and membranes facilitating the release of target compound [57]. The EAE method has shown to improve the extraction yields and to enhance the recovery of bioactive compound in marine seaweed [56,58–62]. Nevertheless, this method has not been extensively applied in extracting the bioactive compound from Indonesian *Sargassum*. Consequently, this study will evaluate the efficiency of EAE in extracting bioactive compound of *Sargassum*. Furthermore, SLE method will also be used as a comparison with the EAE method. Later, the phenolic content and the antioxidant activity from both methods will be determined. In additional, tyrosinase and biofilm inhibition activity of the EAE extracts will also be analyzed in order to acknowledge another possible potential from this extracts. Finally, these extracts will be characterized using the Fourier-Transform Infra-Red (FTIR) spectroscopy. This characterization will serve as a preliminary analysis to detect the presence of the phenolic compound, phlorotannins.

#### 2. Material and Methods

#### 2.1 Seaweed Collection

There were four different species of *Sargassum* collected for this study. *S. polycystum* was collected from Panjang Island, Jepara in October 2013 then *S. aquifolium* and *S.ilicifollium* were collected from Telukawur, Jepara in April 2014. Both of the locations are located in the northern region of Central Java, Indonesia.

The apical and median parts of *Sargassum* thalli were cut leaving the basal part still attached to the substrate for further regeneration, rinsed with seawater and placed in closed boxes. Samples were then transferred to the Central Laboratory of Research and Services Diponegoro University (CORES-DU).

They were washed with tap water to remove the remaining epiphytes and other residual sands and naturally dried away from the sunlight for seven days. After all samples had dried, they were all grinded and stored in a sealed plastic bag that was covered to avoid direct contact with the sunlight.

# 2.2 Solid-liquid extraction (SLE) of Sargassum

25 gr of algal dry material was diluted in 300 ml of four different solvents, i.e. water, methanol 50 % (v/v), pure methanol and ethanol 75 % (v/v). The filtered samples were then evaporated until dryness. The dried extract obtained was then added by 20 ml of  $H_2O$  in order to get the crude extract. Finally, the crude extracts were frozen and lyophilized for further analysis.

# 2.3 Enzyme-assisted extraction (EAE) of Sargassum

19.5 gram of dry material was diluted in 300 ml of water, then 5 % (of dry material) commercial enzymes were added. Viscozyme<sup>®</sup> – a carbohydrase, and Protamex<sup>®</sup> – a protease, were chosen. These enzymes were provided by Novozymes<sup>®</sup>, manufactured in Denmark. Extraction without enzyme served as contro, also called the aqueous extraction.

Sample solutions were incubated for 3 hours in 40 °C. After the filtration, the soluble samples were then frozen and lyophilized for further analysis. The residual materials were stored in -20 °C for TPC analysis to determine the insoluble phenolic compound that might not be digested by the enzymes.

#### 2.4 Total Phenolic Content Analysis

Folin-Ciocalteu is a commonly used method for measuring the phenolic content. This assay is based on the reduction-oxidation (redox) reactions, which are usually considered to be relatively stoichiometric and on the redox potential of the phenolic hydroxyl group [63].

0.5 ml of samples or standard (Phloroglucinol) was added with 0.5 ml ethanol 95 %. 2.5 ml  $\rm H_2O$  was then introduced to the samples or standard solution followed by 0.25 ml of Folin-Ciocalteu 50 %. All samples were agitated and left to stand for 5 minutes. In the end, 0.5 ml of 5 % of  $\rm Na_2CO_3$  (in 100 ml  $\rm H_2O$ ) was added. Then the optical density was read against the blank prepared at 725 nm. Phloroglucinol was a standard for the calibration curve prepared in different concentrations ranged from 0 – 100  $\rm \mu g/ml$  [64–66].

# 2.5 Antioxidant Activity

2,2-diphenyl-1-picrylhydrazyl or DPPH is a stable free radical. DPPH assay is based on the theory that a hydrogen donor is an antioxidant. It measures compounds that are radical scavengers. DPPH show a strong absorption maximum at 517 nm. This analysis is based on modified method of Yen and Chen (1995) and Chen et al. (2008).

A series of BHA, BHT and Ascorbic Acid solution – serves as standard – in different concentration is prepared (2 – 50 µg/ml). 0.25 mM of DPPH solution is introduced to 100 µL samples solution. Samples solution of *S. muticum* hydrolysates was made in different concentration by diluting the solution stock in methanol (0 – 1000 µg/mL). Before reading the optical density at 517 nm, all samples are incubated in 40 °C for 30 minutes. Then, the percentage of inhibition is calculated by following formula:

$$I(\%) = \left[ \frac{(A_C - A_S)}{A_C} \right] \times 100$$

I(%) = Inhibition (express in %)

 $A_{C}$  = Absorbance of control

 $A_S$  = Absorbance of samples

IC<sub>50</sub> of samples is determined based on the regression obtained from dose response curve. IC is defined at the concentration sufficient to obtain 50 % of a maximum scavenging capacity.

# 2.6 FRAP (Ferric Reducing Antioxidant Power)

This method is based on the reduction of a ferroin analogue, the complex of tripyridyltriazine  $Fe(TPTZ)^{3+}$ , to the intensely bluish colour complex of  $Fe(TPTZ)^{2+}$  by the presence of antioxidants in acidic medium. Then, the results are obtained from the absorbance at 593 nm and expressed as  $\mu m$   $Fe^{2+}$  equivalents (Benzie & Strain 1996; 1999)

The reagent of TPTZ consisted of 300 mM of acetate buffer with pH 3.6, 10 mM TPTZ (2, 4, 6-tripyridyl-s-triazine) and 20 mM FeCl<sub>3</sub>  $_6$ H<sub>2</sub>O solution. 50  $\mu$ L of *S. muticum* hydrolysates was introduced in microplate mixed with 150  $\mu$ L reagent. After 15 minutes of incubation, the absorbance was read at 593 nm. The results were expressed in  $\mu$ M equivalent to FeSO<sub>4.7</sub>H<sub>2</sub>O calculated from the calibration curve [69].

# 2.7 Antibacterial Activity

Agar diffusion method [70] was chosen as method for evaluating the antibacterial activity of *Sargassum*. One gram positive bacteria – *Bacillus subtilis* – and two gram negative bacteria – *Pseudomonas aeruginosa* and *Escherichia coli* – were chosen for this test. These bacteria were prepared in liquid nutrient broth media with bacterial density 1.5 x 10<sup>8</sup> cfu/ml. Bacterial suspension was poured on to solidified agar media and incubated for 1 h in 37 °C.

5 mg/ml of SL extracts and hydrolysates were prepared in sterile physiological water. 20  $\mu$ L of tested samples solution – hydrolysates solution or positive control (Phosphomycin, Ampicillin and Streptomycin) – was introduced to sterile paper disc. All samples were made in triplicate. Later, the impregnated sterile paper discs were placed on the agar media in accordance with samples coding made in advance. The petri dishes were then incubated at 37 °C for 48 h. Zone of inhibition established on the agar media was then measured and expressed in cm.

# 2.8 Statistical Analysis

All results are expressed as mean  $\pm$  standard deviation (SD) with n = 3. As the design of this research was a factorial design with more than one independent variable than two ways analysis of variance (ANOVA) was chosen as statistical analysis using IBM SPSS Statistics 20. Further, the statistical difference between samples were determined via Tukey Post Hoc Test with significance level at 5% (p<0.05).

### 3. Results

# 3.1 Comparison of SLE and EAE Extraction Yield, Total Phenolic Content, Antioxidant and Antibacterial activity of Sargassum

Two-ways analysis of variance was applied to determine whether there was a significant effect or not from several independent factors towards the dependant variable. In the case of *Sargassum* yield of extraction, total phenolic content and antioxidant activity – all were the dependant variables, different species and different type of extractants served as independent factors that would influence the dependant variables.

Statistical analysis of *Sargassum* extraction yield showed that there was a significant different between the three species extracted, i.e *S. aquifolium*, *S. ilicifolium* and *S. polycystum*. Different type of extractants used also gave a significant effect to the extraction yield. This result implied that the extraction yield of *Sargassum* showed an inter-individually significant effect.

As seen in Table 1, the EAE yielded higher dry material compared to the SLE. The order of samples with highest yield of enzymatic extraction was *S. polycystum* > *S. aquifolium* > *S. ilicifolium*. The highest extraction yield was shown by *S. polycystum* extracted with Protamex<sup>®</sup>,  $38.1 \pm 6.8 \%$  of dry material.

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**Table 1.** Extraction yield of Sargassum (% of dry material) in different method of extraction (mean  $\pm$  standard deviation, n=3).

Samples	Water	MeOH	MeOH 50%	EtOH 75%	Viscozyme®	Protamex <sup>®</sup>
S. aquifolium	$17.3 \pm 2.2^*$	$5.7 \pm 1.8^*$	$15.8 \pm 1.5^*$	$12.3 \pm 0.4^*$	$29.5 \pm 5.4^*$	$26.2 \pm 5.8^*$
S. ilicifolium	$14.7 \pm 0.8^*$	$3.1 \pm 0.7^*$	$7.3 \pm 1.0^*$	$7.5 \pm 1.3^*$	$27.9 \pm 1.4^*$	$21.4 \pm 2.0^{*}$
S. polycystum	$22.8 \pm 1.1^*$	$8.3 \pm 3.2^*$	$13.4 \pm 1.9^*$	$12.5 \pm 1.9^*$	$35.7 \pm 2.6^*$	$38.1 \pm 6.8^*$

<sup>\*</sup>Significantly different at p<0.05.

The total phenolic content of Sargassum is presented in Table 2 expressed in % of dry material. Viscozyme® extracted more phenolic content in S. aquifolium and S. polycstum with  $7.0 \pm 1.3$  and  $8.1 \pm 2.6$ % of dry material, respectively, compared to the organic solvents. In the contrary, the highest phenolic content of S. ilicifolium was obtained from the methanol  $7.5 \pm 2.4$ % of dry material. Apparently, the efficiency between Protamex and methanol 50% (v/v) in extracting the phenolic compound were the same in all Sargassum. As can be seen on Table 2, they only extracted between 3 up to 5% of phenolic content.

The antiradical activity expressed in the value of IC<sub>50</sub> of Sargassum is presented in Table 3. S. polycystum only showed its antiradical activity when they were extracted with ethanol 75% and Protamex<sup>®</sup> and the best activity was shown by Protamex<sup>®</sup> with IC<sub>50</sub> 1.9  $\pm$  1.1 mg/mL. The weakest antiradical activity came from the methanolic extract of S. aquifolium with IC<sub>50</sub> 23.8  $\pm$  9.0 mg/mL. The three *Sargassum* extracted with Protamex<sup>®</sup> gave a better antiradical activity compared to the other extractants, it can be seen by their IC<sub>50</sub> that were lower as presented on Table 3. The 75% (v/v) ethanolic extracts of S. aquifolium and S. ilicifolium showed no antiradical activity. Meanwhile, the antiradical activity of all samples were still weaker compared to the standards, i.e BHA and BHT with  $IC_{50} 8x10^{-3} \pm 0.5$  and  $12 \times 10^{-3} \pm 4.8$  mg/ml, respectively. The statistical analysis of two-ways ANOVA showed that there was a significant difference between the three species of Sargassum and between the types of extractant that affected the antiradical activity. The statistical analysis of two-ways ANOVA showed that there was a significant difference between the three species of Sargassum and between the types of extractant used. Based on its phenolic content, the order of samples with highest content was S. polycystum > S. aquifolium > S. ilicifolium. In addition, the efficiency order of extractants in extracting the phenolic compounds of S. polycystum was Viscozyme<sup>®</sup> > ethanol 75% (v/v) > Protamex<sup>®</sup> > methanol > methanol 50% (v/v) > water. As for S. aquifolium, the efficiency order was Viscozyme<sup>®</sup> > methanol > ethanol 75% (v/v) > methanol 50% (v/v) > Protamex<sup>®</sup> > water. Then, for S. ilicifolium, the efficiency order was methanol > ethanol 75% (v/v) > methanol 50% (v/v) > Protamex<sup>®</sup> > Viscozyme<sup>®</sup> > water.

Table 4 presents the reducing power activity also known as FRAP of *Sargassum* expressed by a concentration in  $\mu$ M equivalent with Fe<sup>2+</sup> ( $\mu$ M Eq Fe<sup>2+</sup>). Vitamin C was used as standard with reducing power 58.1  $\pm$  0.7  $\mu$ M Eq Fe<sup>2+</sup>. The reducing power activity that was almost near as the standard was the aqueous phase of *S. polycystum* with 43.5  $\pm$  0.7  $\mu$ M Eq Fe<sup>2+</sup>. In other word, this sample had the highest reducing power activity. Meanwhile, the lowest reducing power activity was shown by *S. ilicifolium* extracted with ethanol 75%, 7.9  $\pm$  1.2  $\mu$ M Eq Fe<sup>2+</sup>. In addition, it appears that Viscozyme<sup>®</sup> extracts of *S. aquifolium* and *S. polycystum* had better reducing power activity compared to methanol, methanol 50%, ethanol 75% (v/v) and even Protamex<sup>®</sup>. The statistical analysis of two-ways ANOVA showed that there was a significant difference between the three species of *Sargassum* and between the types of extractant that affected the reducing power activity.

As for the antibacterial activity of two methods of extraction, the tested bacteria were sensitive only to solid-liquid extraction (data not shown). In other words, the enzymatic extracts showed no sign of inhibition against the bacterial growth. It was assumed that the sugar contained in our enzymatic extracts might provoke the growth of bacteria instead of inhibiting them.

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**Table 2.** Total phenolic content (TPC) of Sargassum (% of dry material) (mean  $\pm$  standard deviation, n=3).

Samples	Water	Methanol	Methanol 50 %	Ethanol 75%	Viscozyme <sup>®</sup>	Protamex <sup>®</sup>
S. aquifolium	$1.7 \pm 0.0^*$	$6.3 \pm 1.5^*$	$3.7 \pm 0.2^*$	$5.9 \pm 0.4^*$	$4.7 \pm 1.3^*$	$4.2 \pm 0.0^*$
S. ilicifolium	$2.1 \pm 0.0^*$	$7.5 \pm 2.4^*$	$5.3 \pm 0.3^*$	$5.9 \pm 0.4^*$	$3.6 \pm 0.1^*$	$5.0 \pm 0.3^*$
S. polycystum	$1.4 \pm 0.0^*$	$4.8 \pm 0.3^*$	$4.0 \pm 0.3^*$	$5.8 \pm 0.6^*$	$4.0 \pm 0.4^*$	$6.1 \pm 0.1^*$

<sup>\*</sup>Significantly different at p<0.05.

**Table 3.** Antioxidant activity of *Sargassum* expressed in IC50 (mg/ml) (mean  $\pm$  standard deviation, n=3).

	DPPH (IC 50 mg/ml)					
Samples	Water	Methanol	Methanol 50%	Ethanol 75%	Viscozyme®	<b>Protamex</b> ®
S. aquifolium	$14.6 \pm 5.7$	$23.8 \pm 9.0^*$	$22.3 \pm 8.0^*$	n.a	n.a	$5.7 \pm 7.9$
S. ilicifolium	n.a	$17.8 \pm 9.4^*$	$3.3 \pm 1.7^*$	n.a	$6.2 \pm 3.4$	$2.9 \pm 1.7$
S. polycystum	n.a	n.a	n.a	$5.2 \pm 5.9$	n.a	$1.9 \pm 1.1$

<sup>\*</sup>Significantly different at p<0.05.

**Table 4.** Reducing activity of *Sargassum* expressed in  $\mu$ M Equivalent Fe<sup>2+</sup> (mean  $\pm$  standard deviation, n=3).

Samples	FRAP (Concentration Equivalent Fe <sup>2+</sup> (μM))						
	Water	Methanol	Methanol 50%	Ethanol 75%	Viscozyme®	<b>Protamex</b> ®	
S. aquifolium	$14.9 \pm 1.7^*$	$12.6 \pm 2.5^*$	$22.4 \pm 4.9^*$	$14.4 \pm 0.3^*$	$33.2 \pm 2.5^*$	$17.0 \pm 0.7^*$	
S. ilicifolium	$23.7 \pm 1.6^*$	$11.1 \pm 0.8^*$	$14.9 \pm 0.8^*$	$7.9 \pm 1.24^*$	$13.4 \pm 2.2^*$	$11.7 \pm 1.2^*$	
S. polycystum	$43.5 \pm 1.8^*$	$12.8 \pm 3.1^*$	$16.9 \pm 2.6^*$	$13.8 \pm 1.9^*$	$27.7 \pm 4.9^*$	$15.9 \pm 3.0^*$	

<sup>\*</sup>Significantly different at p<0.05.

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#### 4. Discussion

Phlorotannins, class of phenolic compounds synthesized only in brown seaweed [71], is one of the most studied bioactive compounds due to their wide range of bioactivities, showing a promising potential for the nutraceutical, pharmaceutical and cosmeceutical industries. As reported by previous studies, the increasing interest towards phlorotannins occurs as they have been for their antioxidant [72–77], antimicrobial [78–80], antitumor [81,82], antidiabetic [83], antiallergic [84], anti-inflammatory [85], and antiviral effects [51,86,87].

SLE has become the most used method during the exploration of marine bioactive compounds, in this case, phlorotannins. This method mostly relies on the polarity of the solvents used due to the chemical nature of target compound [88]. In general, the process in SLE method can be categorized in three parts; (1) changing phase of the solute, in this case the target compounds, as it dissolve in the solvent, (2) its diffusion through the solvent in the pores of the solid to the outside of the particles, and (3) the transfer of the solute from the solution in contact with the particles to the main bulk of the solution [89]. The more dispersed the solute in the solid material, the more difficult the extraction will be and the extraction rate will fall. This is because the solvent will have to penetrate further in to the layer of solid material in order to reach the solute [89]. Furthermore, the efficiency of SLE highly depends on the particles size, temperature, and agitation during the process [90]. In the extraction of phenolic compounds, the storage time of dry materials and conditions, as well as the presence of interfering substances are other important factor to consider, influencing the extraction efficiency [91]. In marine seaweed, the main drawback during the extraction of phenolic compounds occurs due to the presence of complexes polysaccharides as the main component of seaweed's cell wall. Brown seaweed's phlorotannins are known to be incorporated in its cell wall [92] covalently bonded to the polysaccharides [35] and proteins [93]. Therefore, it requires a strong condition to degrade these bonds and extract the phlorotannins. The SLE method does not meet this requirement since it does not degrade such complex bonds since it works based on the solubility of target compound in to the solvents [89]. In addition, the polarity of target compound becomes the most important factor in choosing the solvent [94]. Methanol, ethanol, acetone, ethyl acetate, hexane [50,72,75,95,96] are some examples of organic solvents mostly used for phlorotannins extraction.

This study concerns about the possibility in applying alternative method that is not only environmentally friendly but also more efficient in extracting the phlorotannins. For that reason, enzyme-assisted extraction (EAE) is chosen as the alternative method. Additionally, the efficacy of SLE – as the conventional method, and EAE – as the alternative one, in extracting the phlorotannin from Sargassum were compared. For the SLE method, four different organic solvents were chosen, i.e water, methanol, methanol 50% (v/v), and ethanol 75% (v/v). Meanwhile, for the EAE method, there were two commercial enzymes, i.e. Viscozyme<sup>®</sup> – a carbohydrase, and Protamex<sup>®</sup> – a protease.

Based on the extraction yield of these two methods, the EAE yielded the dry algal material higher compared with the SLE (Table 1). As can be seen from this table, in three hours of extraction, the SLE extraction yield ranged from 3% to 15% of dry material. In the contrary to the SLE method, the extraction yield of EAE ranged from 21% to 38%. In overall, the average increase of extraction yield when using the Viscozyme® was around 62%. As for the Protamex®, the average increase of extraction was 58%. This result had shown the efficacy of EAE method in improving the extraction yield applied under the same circumstances as the SLE method. Interestingly, the efficiency of EAE in improving the extraction yield seems to be species-dependent. As can be seen on Table 1, Protamex® yielded more dry material in *S. polycystum* with 38% than Viscozyme®. In the contrary, the extraction yield of *S. aquifolium* and *S. ilicifolium* were higher in Viscozyme®, with 29% and27%, respectively. Viscozyme® was also yielded more soluble material in *S. fulvellum* (24%), *S. horneri* (30%), *S. correanum* (30%) and *S. thunbergii* (31%) than Protamex, with 17%, 27%, 21%, and 25%, respectively [97]. In *S. muticum*, the Viscozyme managed to extract 29% of soluble material [41,98]. It appears that the cellulose and the multicarbohydrase complex action of Viscozyme® were responsible for higher solid content in lyophilized extracts [98].

The EAE had also recovered more of phlorotannins represented by the total phenolic content. By using Viscozyme, the total phenolic content of Sargassum was 28% higher than using the organic solvents. Unfortunately, the phenolic content of Protamex® extracts of Sargassum were not in consistent with the Viscozyme® extracts. The Protamex did not seem to recover the phenolic content of Sargassum as they yielded lower content compared to the organic solvents, except for water. It implied that the Viscozyme<sup>®</sup> recovered more of phenolic compounds in Sargassum than the Protamex<sup>®</sup>. Phlorotannins are hydrophilic cell-wall-bound compounds [99] and despite of its soluble characteristic, compounds attached to the cell wall are not easily extracted using the typical extraction methods such as the SLE [56,61]. The application of Viscozyme<sup>®</sup>, as a carbohydrase, during the extraction allowed to facilitate the recovery of phenolic compound as this enzyme degrades the cell wall of seaweed mainly composed of polysaccharides. Viscozyme<sup>®</sup> has been extensively used for the recovery of phenolic compounds of terrestrial and marine plants as reported by many studies [41,65,100–102]. However, the efficiency of extraction relies on the optimum conditions applied. Selecting the appropriate hydrolytic enzymes is important to digest specific polymer bonds present in seaweed cell wall followed by the selection of suitable process conditions for maximum recovery of active compounds [56]. Combination of incubation time-temperature, pH, size of the interested molecules and agitation are mentioned to play a critical role during the process [14].

Phlorotannins have been reported to be responsible for many bioactivities, such as antioxidant, antimicrobial, antiviral and many more, as reported by previous studies [9,29,51,96]. The key of phlorotannins important bioactivities lies to its eight interconnected ring as they contain more hydroxyl groups than other tannins – hydrolysable tannin and condensed tannins [107,108]. As a consequence, they have greater antioxidant activity [107]. Hydrogen atom transfer is one of its mechanisms against the effects of unregaluted production of free radicals [36]. As performed in this study, *Sargassum* extracts – extracted with SLE and EAE method- showed their capability in scavenging the free radicals with best activity exhibited by the Protamex® extracts in spite of Viscozyme® extracted more phenolics. It suggested the presence of other molecules like polysaccharides or proteins as shown by the FTIR since the current study used crude extracts instead of the purified. Besides free radical scavenging, our results showed that the enzymatic extracts of *Sargassum* had the ferric reducing ability – a conversion of Fe<sup>III</sup> to Fe<sup>II</sup>. In addition, it showed quite promising potential as skin whitening antifouling coating. Such potential was based on the tyrosinase and biofilm inhibition activities of *Sargassum* enzymatic extracts.

Antioxidant activity of *S. horneri*, *S. fullvelum*, *S. correanum*, and *S. thunbergii* have been reported by Heo et al [65]. *S. correanum* contained higher phenolics when it was extracted with Viscozyme® and higher free radical scavenging activity compared to the others. Another study also showed that Viscozyme® used to extract *S. horneri* had the ability to scavenge the free radicals until 95% of inhibition [109]. Sanchez-camargo et al. [41] stated that the total phenols and antioxidant activity of *S. muticum* extracted with Alcalase® and Viscozyme® were improved compared to control – aqueous extraction. Therefore, it is obvious that enzymes used have facilitated the release of active compounds and in the same time it enhances the bioactivities especially the antioxidant – best known activity for phlorotannins.

# 5. Conclusion

A strong urge to explore marine bioactive compounds has brought green chemistry concept to the surface along the process. The principle of this concept is basically to apply a method that is environmentally friendly and favorable for human interest as well. In the exploration of marine bioactive compounds that can be further applied for the nutraceutical, pharmaceutical and cosmeceutical industries, the realization of this concept is crucial. Therefore, many studies have highlighted the possibilities of enzyme-assisted extraction as an alternative method that might replace the use of organic solvent to extract the active compounds.

This study has revealed that by using enzymes, the efficiency of the extraction increased compared to the solid-liquid extraction. This method not only improves the quality and quantity of extracts but it

also shows its promising potential by showing diverse bioactivities. Nevertheless, one should note that this study still works with crude extracts. Therefore, the interference of other compounds like polysaccharides and proteins should be considered. As a consequence, it is crucial to further evaluate the bioactivities of purified samples and characterize them.

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