







An enzymatic extract of the proliferative red seaweed Solieria chordalis reduces the virulence of Pseudomonas aeruginosa and stimulates the immune system of Caenorhabditis elegans.

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Photo taken in St Gildas de Rhuys, France, on July 2015



NANTES

Since 2002, more than 2 000 tons of Solieria chordalis had to be removed from beaches in South Brittany. (Neveu 2016)

Between 2002 and 2015: annual cost for seaweed collection : ~ 105 000 US\$ (Neveu 2016)

Economical and ecological problems

Objective

Active Water Soluble Extracts for the prevention of bacterial infections in animals, plants and humans

An environmentally friendly extraction method: Enzyme-Assisted Extraction (EAE)

Using the pathogenicity model Caenorhabditis elegans –

Pseudomonas aeruginosa

Proliferative macroalga: Solieria chordalis

Objective

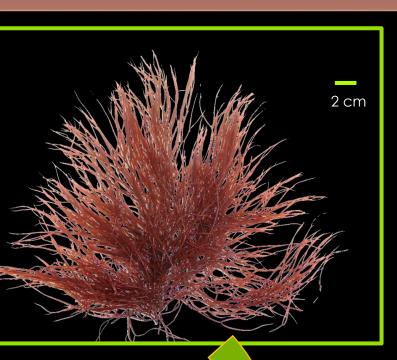
Proliferative macroalga: Solieria chordalis



Part I : Solieria chordalis





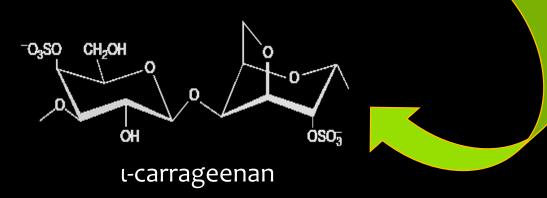


<u>Classification :</u>

Empire : Eukaryota Kingdom : Plantae Phylum : Rhodophyta Class : Florideophyceae Order : Gigartinales Family: Solieriaceae Genus : Solieria Species : Solieria chordalis.



Photos during scuba diving in South Brittany on october/november 2014 (Littoralg project)



Sulfated polysaccharides ~50 % of the dry matter with numerous applications.
(Bondu et al. 2010, Hardouin et al. 2013, Prajapati et al. 2014)



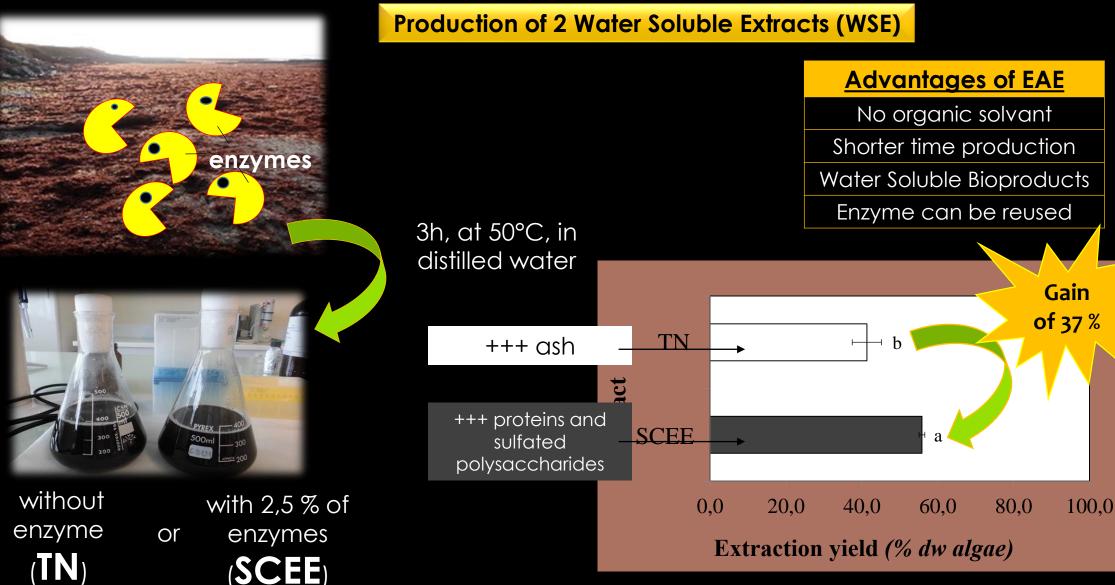
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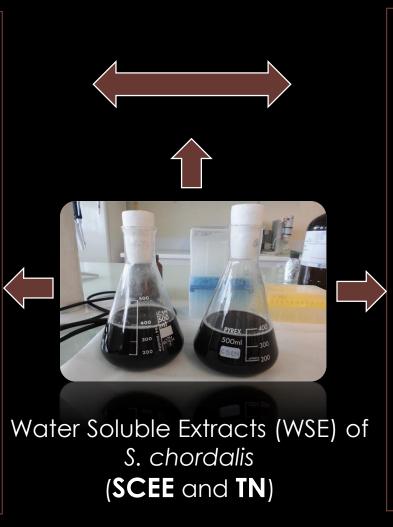
Pathogenicity model C. elegans -P. aeruginosa :

Antibacterial activity

lmmuno--stimulant activity Caenorhabditis elegans (nematode size: 1 mm)



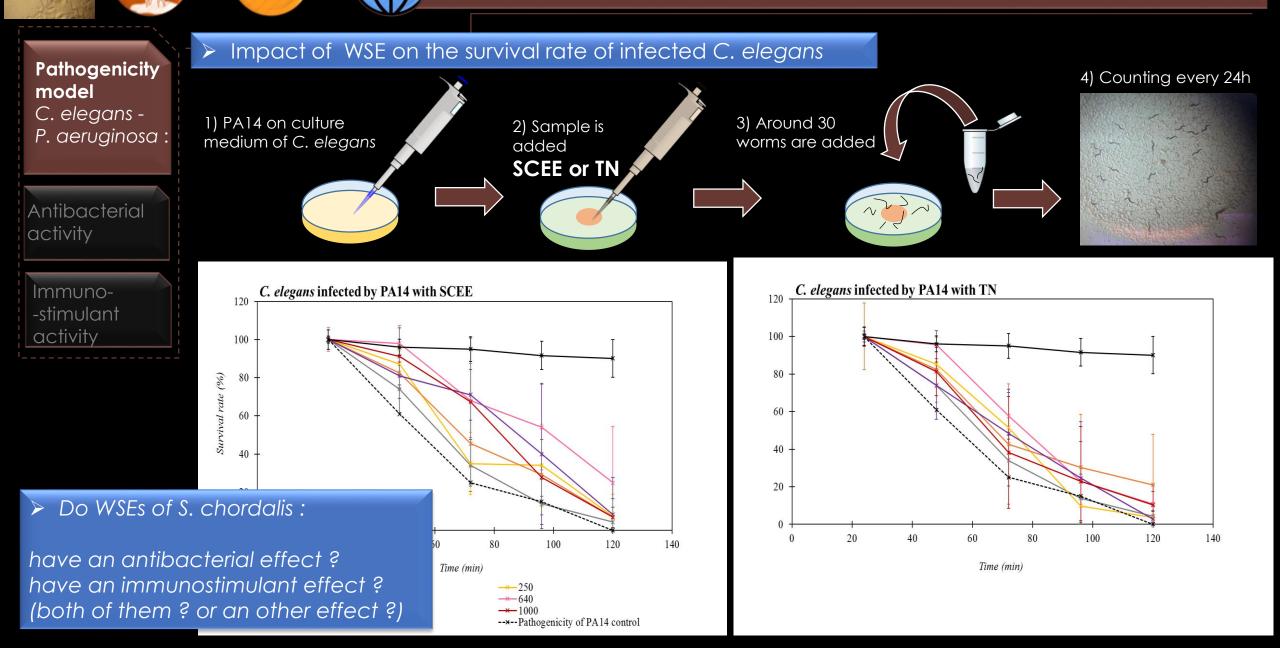
Innate Immune System					
Antibacterial effectors	Subsitence				
	(lipase, lectin)				
(antibacterial proteins, lysozyme)					

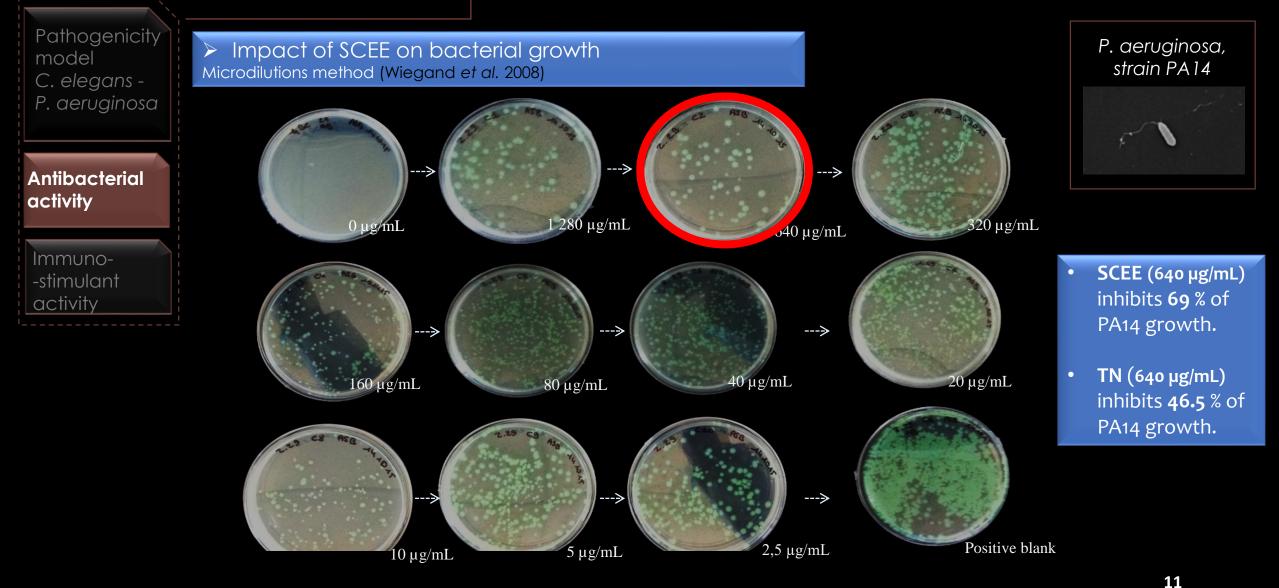


P. aeruginosa, strain PA14 (bacteria size 1-5 µm)



Bacterial virulence						
Virulence Factors (hydrogen cyanide, superoxyde dismutase)	Regulation system : quorum sensing					





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Pathogenicity model C. elegans -P. aeruginosa

$(\Delta\Delta Ct Analysis)$

Antibacterial activity

Immuno--stimulant activity

Impact of SCEE on gene expression of PA14 virulence

PA14 RNA extraction of PA14

640 μg/

mL

Reverse transcription into cDNA



Bacterial virulence analysis



640

sensing

factors

Pathogenicity model C. elegans -P. aeruginosa Ā

Antibacterial activity Immuno-

-stimulant activity

enicity	> Impac	t of SCEE on gene expression of PA14	4 virulence		μg/ mL
ans - ginosa	(∆∆Ct Anal	ysis) Exponential bacterial growth (6h of contact)	Stationary bacterial growth (24h of contact)	P	2A14 ↓
erial				RNA extrac	ction of PA14
endi	rhll	rhll		= =	= == 🗸
	rhIR	rhIR			ranscription cDNA
nt	lasl	lasl			CDNA .
¦	lasR	lasR			
				qF	PCR
	hcnC	hcnC			
	aroE	aroE			
	rpoN	rpoN			virulence Ilysis
	sodB	sodB		Genes	Genes
	sbe	sbe		coding for quorum	coding for virulence

640

Pathogenicity model C. elegans -P. aeruginosa

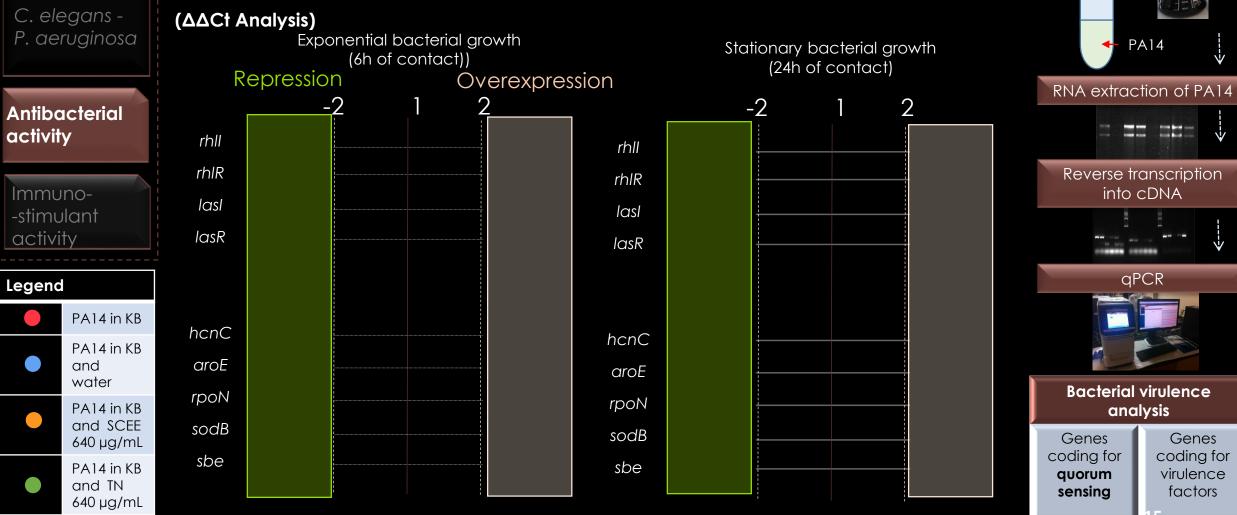




640 µg/

mL

Pathogenicity model C. elegans -P. aeruainosa



Impact of SCEE on gene expression of **PA14 virulence**

Pathogenicity model C. elegans -P. aeruginosa





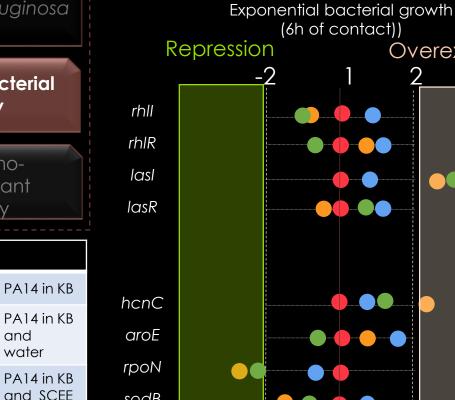
and

640 µg/mL

PA14 in KB

640 µg/mL

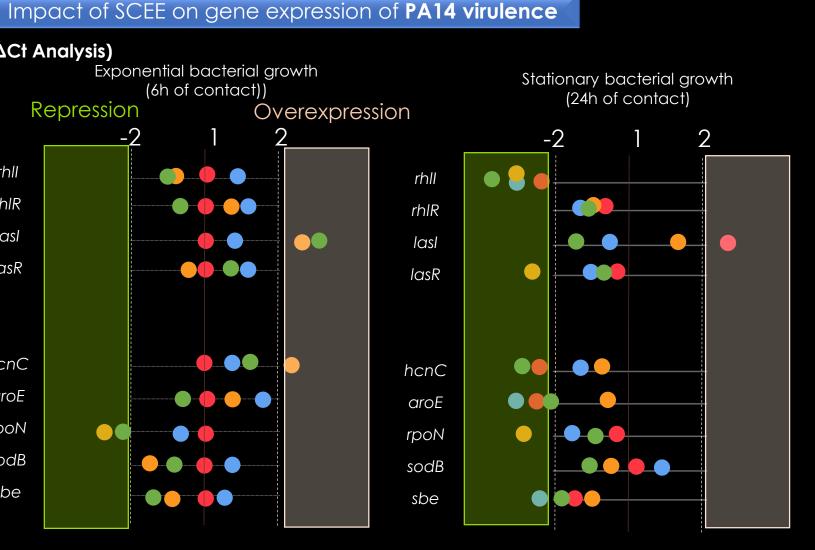
and TN

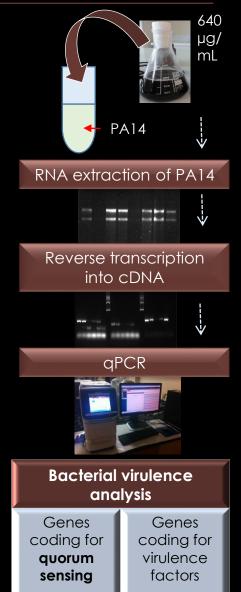


 $(\Delta\Delta Ct Analysis)$

sodB

sbe



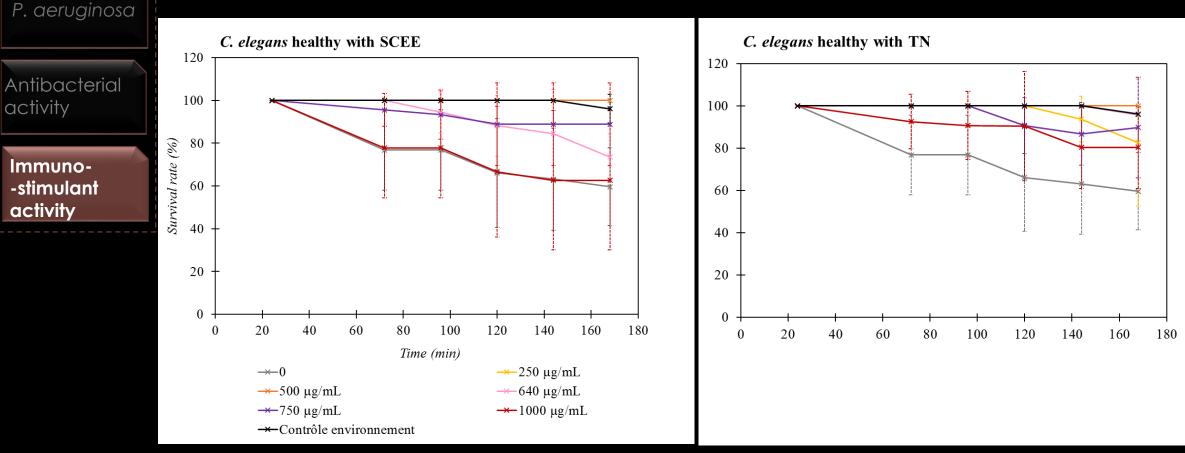






Pathogenicity model C. elegans -

Impact of WSE on the survival rate of healthy C. elegans

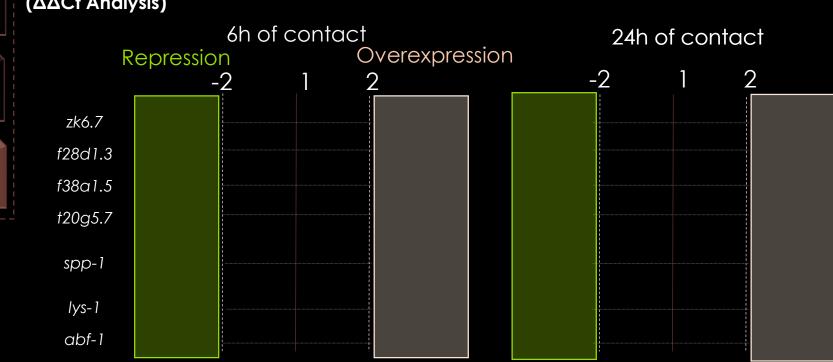




Impact of WSEs on gene expression of innate immune system of heahtly and infected C. elegans

$(\Delta\Delta Ct Analysis)$





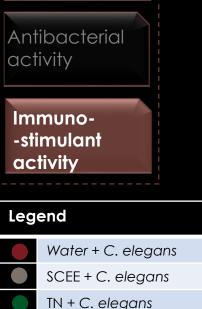
RNA extraction of C. elegans **Reverse transcription** into cDNA aPCR Innate Immune System analysis Antibacterial Stress resistance effector genes genes

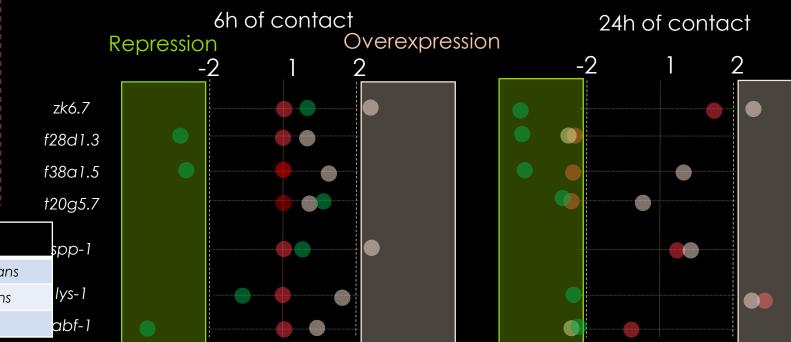
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Impact of WSEs on gene expression of innate immune system of heahtly and infected C. elegans

($\Delta\Delta$ Ct Analysis)



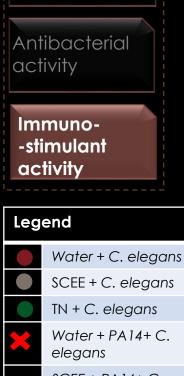


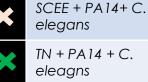
RNA extraction of C. elegans **Reverse transcription** into cDNA aPCR Innate Immune System analysis Antibacterial Stress resistance effector genes genes 19

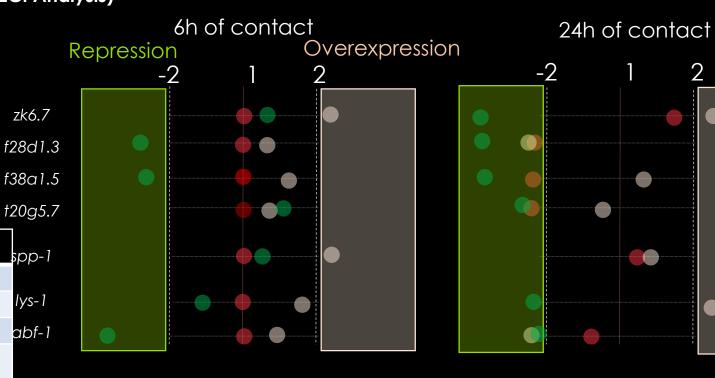


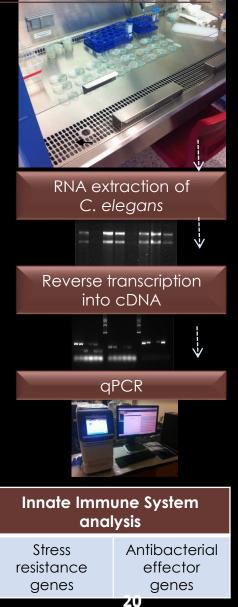


($\Delta\Delta$ Ct Analysis)





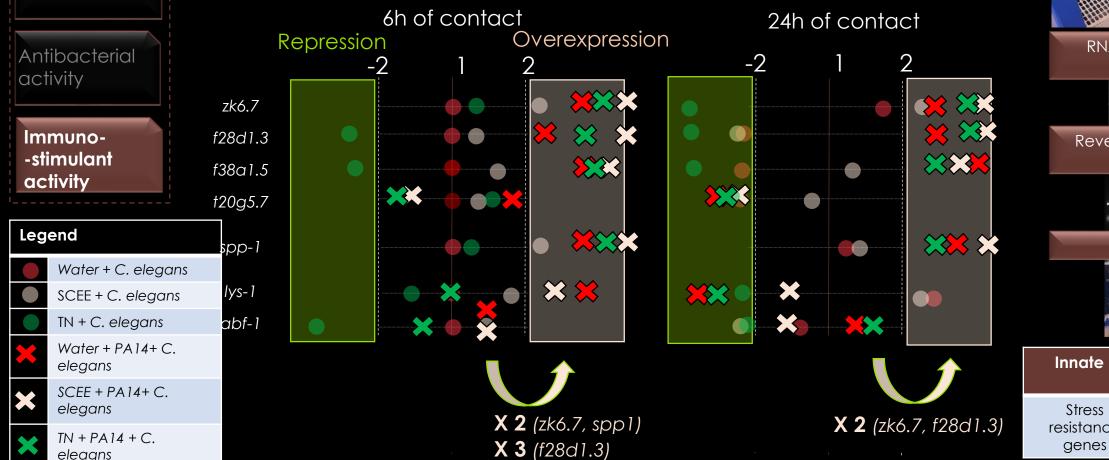






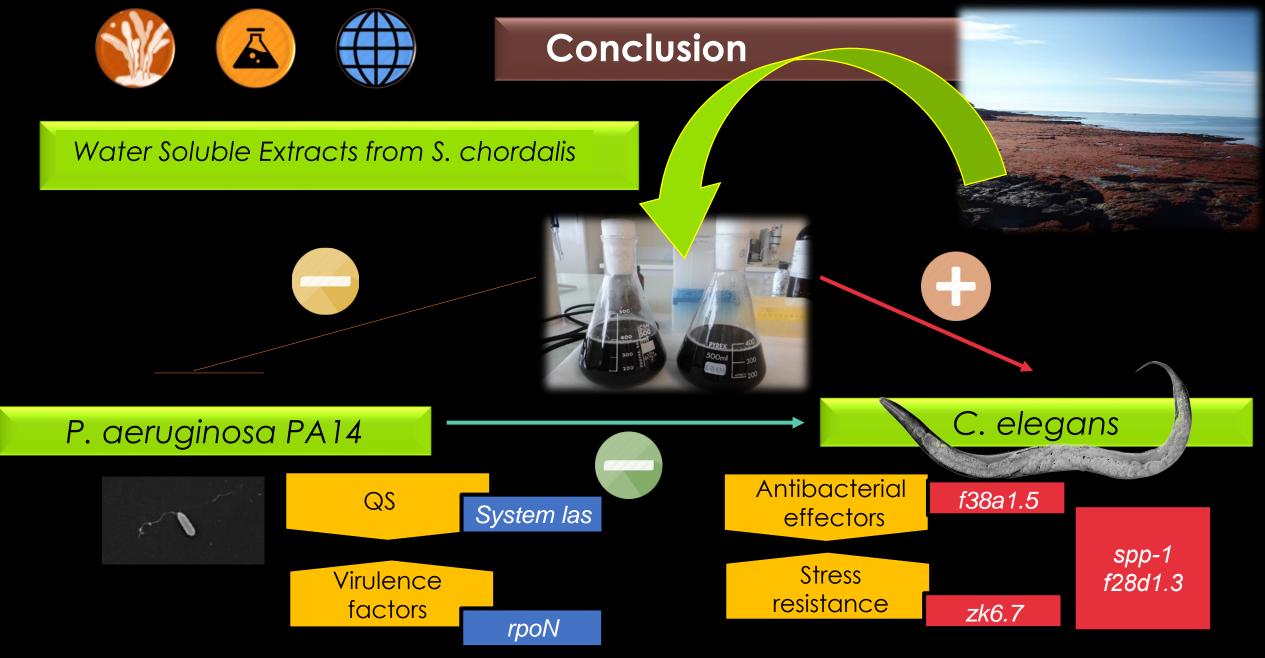
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($\Delta\Delta$ Ct Analysis)



RNA extraction of C. elegans **Reverse transcription** into cDNA aPCR Innate Immune System analysis Antibacterial resistance effector genes

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In accord with litterature (Kandasamy et al. 2012; Liu et al. 2013; Kandasamy et al. 2015 Kulshreshtha et al. 2016) and confirms the study of Bondu et al. 2010.





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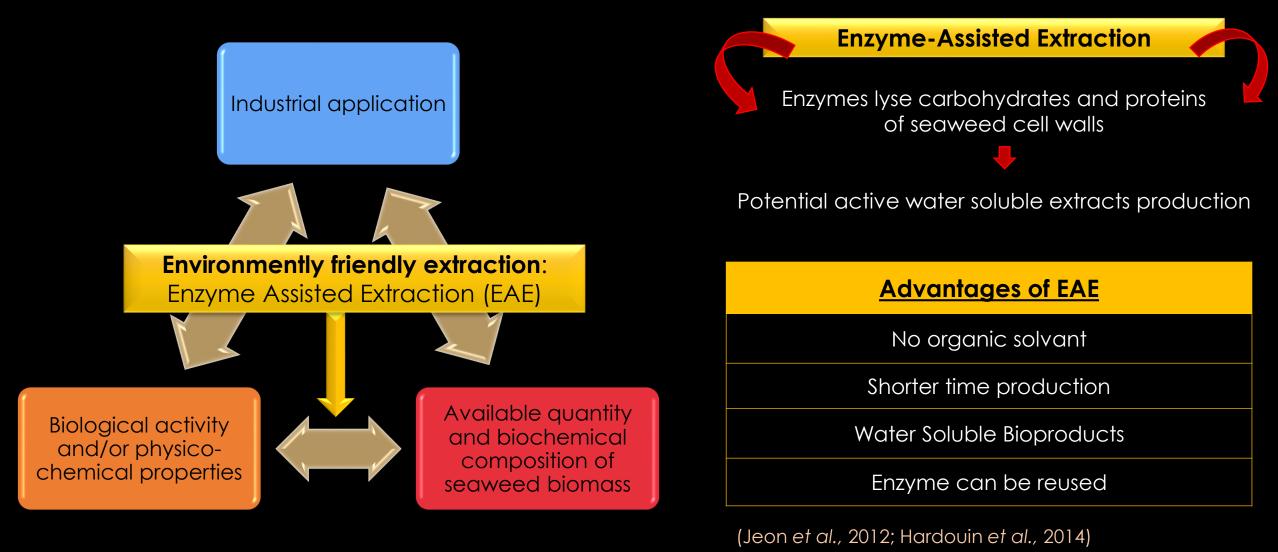






Part II : Enzyme-Assisted-Extraction







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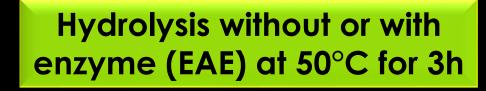
25





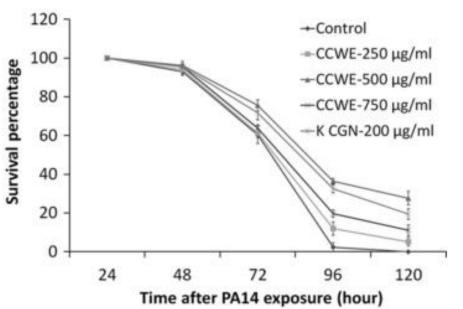
Biochemical composition of S. chordalis and of extracts SCEE and TN

	Dry matter Lipids		Ash content Proteins		Uronic acids	Sulfate groups	Neutral sugars
	(% FW)	(% DW)	(% DW)	(% DW)	(% DW)	(% DW)	(% DW)
S. chordalis	10.0 ± 0.6	3.0 ± 0.04	43.3 ± 3.1	22.0 ± 2.0	3.1 ± 0.1	14.7 ± 1.0	31.7 ± 0.7

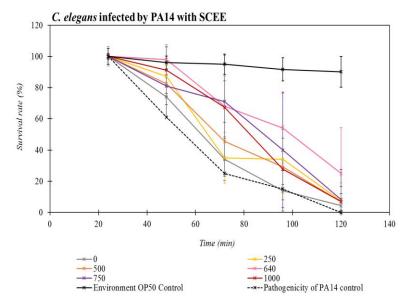


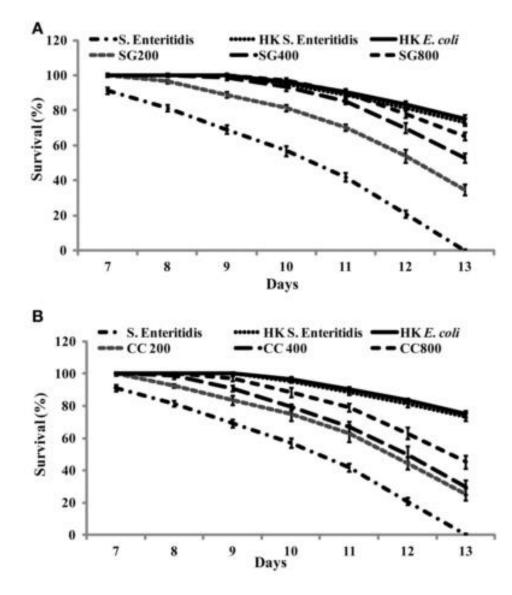
	Others	Ash content	Proteins	Uronic acids	Sulfate groups	Neutral sugars
(% DW)		(% DW)	(% DW)	(% DW)	(% DW)	(% DW)
TN	9.2 ± 1.7	70.9 ± 1.0	4.5 ± 0.1	1.0 ± 0.1	1.6 ± 0.4	12.8 ± 0.1
SCEE	3.6 ± 4.2	56.5 ± 1.6	8.9 ± 2.0	0.3 ± 0.2	3.2 <u>+</u> 0.1	27.5 ± 0.3

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Liu J, Hafting J, Critchley AT, et al (2013) Components of the Cultivated Red Seaweed *Chondrus crispus* Enhance the Immune Response of Caenorhabditis elegans to Pseudomonas aeruginosa through the pmk-1, daf-2/daf-16, and skn-1 Pathways. Appl Environ Microbiol 79:7343–7350. doi: 10.1128/AEM.01927-13





Kulshreshtha G, Borza T, Rathgeber B, et al (2016) Red Seaweeds *Sarcodiotheca gaudichaudii* and *Chondrus crispus* down Regulate Virulence Factors of Salmonella Enteritidis and Induce Immune Responses in Caenorhabditis elegans. Front Microbiol 7:421. doi: 10.3389/fmicb.2016.00421