

Non-indigenous seaweeds in the Northeast Atlantic Ocean, the Mediterranean Sea and Macaronesia: a critical synthesis of diversity, spatial and temporal patterns

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- 2 Atlantic Ocean, the Mediterranean Sea and
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⁵⁴**Abstract**

- ⁵⁵Effective monitoring and combatting the effect of non-indigenous seaweeds relies on a solid
- 56 confirmation of the non-indigenous status of the species. We critically analysed the status of
- 57 presumed non-indigenous seaweed species reported from the Mediterranean Sea, the Northeast
- ⁵⁸Atlantic Ocean and Macaronesia, resulting in a list of 140 species whose non-indigenous nature is
- ⁵⁹undisputed. For an additional 87 species it is unclear if they are native or non-indigenous
- ⁶⁰(cryptogenic species) or their identity requires confirmation (data deficient species). We discuss the
- ⁶¹factors underlying both taxonomic and biogeographic uncertainties and outline recommendations to
- 62 reduce uncertainty about the non-indigenous status of seaweeds. Our dataset consisted of over

⁶³19,000 distribution records, half of which can be attributed to only five species (*Sargassum*

⁶⁴*muticum*, *Bonnemaisonia hamifera*, *Asparagopsis armata*, *Caulerpa cylindracea* and *Colpomenia*

⁶⁵*peregrina*), while 56 species (40%) are recorded no more than once or twice. In addition, our

⁶⁶analyses revealed considerable variation in the diversity of non-indigenous species between the

67 geographic regions. The Eastern Mediterranean Sea is home to the largest fraction of non-

⁶⁸indigenous seaweed species, the majority of which have a Red Sea or Indo-Pacific origin and have

⁶⁹entered the Mediterranean Sea mostly via the Suez Canal. Non-indigenous seaweeds with native

70 ranges situated in the Northwest Pacific make up a large fraction of the total in the Western

⁷¹Mediterranean Sea, Lusitania and Northern Europe, followed by non-indigenous species with a

72 presumed Australasian origin. Uncertainty remains, however, regarding the native range of a

⁷³substantial fraction of non-indigenous seaweeds in the study area. In so far as analyses of first

74 detections can serve as a proxy for the introduction rate of non-indigenous seaweeds, these do not

75 reveal a decrease in the introduction rate, indicating that the current measures and policies are

76 insufficient to battle the introduction and spread of non-indigenous species in the study area.

⁷⁷**Highlights**

- ⁷⁸ Non-indigenous seaweed species in the Northeast Atlantic Ocean, the Mediterranean Sea 79 and Macaronesia are critically reanalysed.
- ⁸⁰ >19,000 distribution records revealed considerable variation in diversity of non-indigenous 81 seaweed species in the study area.

⁸²• Taxonomic and biogeographic uncertainties hamper a critical evaluation of the non-

83 indigenous status of many seaweed species.

⁸⁴**Keywords:** biodiversity, Chlorophyta, Europe, invasive alien species, non-indigenous species,

85 Phaeophyceae, Rhodophyta

⁸⁶**Introduction**

⁸⁷Over the course of several centuries, human-mediated transport has led to the introduction and

88 establishment of more than 14,000 non-indigenous species in Europe (EASIN, 2022). Some of

89 these non-indigenous species profoundly affect the abundance, diversity, interactions and evolution

90 of native biota and consequently affect ecosystem structure, functions and services (Simberloff *et*

⁹¹*al.*, 2013; Dawson *et al.*, 2017; Blakeslee *et al.*, 2020). The introduction of non-indigenous species

92 can also result in substantial negative economic impacts (Hulme *et al.*, 2009). The reported costs of

93 biological invasions, at a global level, were estimated to be at least 1.288 trillion US Dollars over ⁹⁴1970–2017 (Diagne *et al.*, 2021). Furthermore, biotic homogenisation and consequently also the ⁹⁵impact of non-indigenous species on native ecosystems are expected to increase in the context of

96 climate change (Bennett *et al.*, 2021).
97 The management of biological The management of biological invasions depends heavily on lists of reliably identified non-⁹⁸indigenous species. Such lists form an essential tool underpinning prevention, control, mitigation or 99 eradication strategies (Kolar & Lodge, 2001), and in particular to facilitate prevention and early 100 detection, which are the most cost-effective for management (Simberloff *et al.*, 2013). In addition, 101 government and management agencies use lists of non-indigenous species in their policies to 102 protect nature and reverse the degradation of ecosystems. For instance, the primary criterion for the 103 descriptor D2 dedicated to non-indigenous species under the European "Marine Strategy" ¹⁰⁴Framework Directive" is the rate of novel introductions per 6-year period (European Commission *et* ¹⁰⁵*al.*, 2021). Comprehensive and accurate lists of non-indigenous species, their respective origin, and ¹⁰⁶geographical and temporal spread are therefore crucial for an effective response and legislation to 107 battle threats imposed by non-indigenous species. Unfortunately, the compilation of such lists is ¹⁰⁸marred by the challenges involved (McGeoch *et al.*, 2012; Costello *et al.*, 2021). At local scales, ¹⁰⁹lists may be confounded by limited occurrence data and hence underestimate the number and spread 110 of non-indigenous species. At a more fundamental level, taxonomic uncertainty and the associated ¹¹¹lack of expertise in species identification are regarded as severe problems (Zenetos *et al.*, 2017). ¹¹²The effects of taxonomic uncertainty are likely more pronounced for less studied taxa and poorly ¹¹³sampled regions. For example, upon re-examination of about 100 potential non-indigenous taxa of ¹¹⁴marine molluscs, almost half of the records turned out to be misidentifications or the distributional ¹¹⁵data were incorrect (Zenetos *et al.*, 2017). While DNA-assisted identification has the potential to 116 solve identification problems, misidentifications of entries in genetic databases combined with ¹¹⁷geographic and taxonomic sampling bias make it a challenge in itself to correctly interpret gene ¹¹⁸sequence data (Viard *et al.*, 2019; Fort *et al.*, 2021; Tran *et al.*, 2022). In addition, taxonomic 119 knowledge is not static. Evolving taxonomic insights, often derived from genetic and biogeographic ¹²⁰studies, alter our views on the indigenous or non-indigenous nature of taxa, requiring checklists to 121 be continuously updated (Taylor, 2010; Guareschi & Wood, 2019). This problem is exacerbated in 122 the marine environment where many cryptic species have been documented (Appeltans et al., 123 2012).

¹²⁴The above-mentioned problems related to lists of non-indigenous species definitely apply to 125 seaweeds, which represent one of the largest groups of marine non-indigenous organisms, 126 constituting between 20 and 29% of all marine non-indigenous species in the Northeast Atlantic

127 Ocean, the Mediterranean Sea and Macaronesia (hereafter referred to as "the study area")

¹²⁸(Schaffelke *et al.*, 2006; Molnar *et al.*, 2008; Katsanevakis *et al.*, 2013) (Fig. 1; Fig. 2). The

129 consequences of non-indigenous species on native ecosystems have only been studied in a very

130 limited number of species. Although some non-indigenous species have been observed to have

¹³¹positive ecosystem effects (e.g. *Gracilaria vermiculophylla* in the Venice Lagoon and Northeast

132 Atlantic mudflats; Davoult *et al.*, 2017; Sfriso, 2020), impact studies on such seaweeds have mostly

133 detected negative ecological effects, with reduction in abundance of native biota being most

¹³⁴frequently reported (Williams & Smith, 2007; Weinberger *et al.*, 2008; Hammann *et al.*, 2013;

¹³⁵Katsanevakis *et al.*, 2014; Maggi *et al.*, 2015; Bulleri *et al.*, 2017; Anton *et al.*, 2019). However,

136 contrary to the evidence of substantial negative impact on coastal ecosystems of many non-

¹³⁷indigenous seaweeds (e.g. *Caulerpa cylindracea*, *Caulerpa taxifolia*, *Codium fragile*), so far

¹³⁸*Rugulopteryx okamurae* is the only seaweed included in the list of invasive alien species of Union

139 concern (COMMISSION IMPLEMENTING REGULATION (EU) 2022/1203 of 12 July 2022

140 amending Implementing Regulation (EU) 2016/1141). This EU regulation enforces member states

141 to adopt measures to prevent, minimise or mitigate the adverse impact of those species.

Regional lists of non-indigenous seaweed species have been regularly published until recently. For the Mediterranean Sea, which has been disproportionately affected by non-indigenous species as well as other stressors (Lejeusne *et al.*, 2010; Katsanevakis *et al.*, 2014), non-indigenous seaweeds have been critically revised on a regular basis (Verlaque *et al.*, 2015; Zenetos *et al.*, 2017; Galil *et al.*, 2021). Non-indigenous seaweeds of Macaronesia were included in Borges *et al.* (2010), Chainho *et al.* (2015), Gallardo *et al.* (2016) and Castro *et al.* (2022). Bárbara *et al.* (2005) and Brodie *et al.* (2016) provided a list of non-indigenous seaweeds as part of a revised check-list of Galician and British seaweeds, respectively. However, there are gaps and uncertainties for some regions, and more importantly, a critical compilation encompassing the Northeast Atlantic Ocean, Macaronesian and Mediterranean regions is currently lacking. The absence of a critically revised list in the study area not only impedes a comprehensive overview of non-indigenous seaweeds, but 153 may also introduce ambiguity related to the status of specific taxa due to differences in the criteria used to define non-indigenous species (see Materials and Methods). In addition, in the absence of a 155 comprehensive list, spatial and temporal patterns of introductions are difficult to deduce.

¹⁵⁶To address this knowledge gap, we compiled a database of non-indigenous seaweeds in the 157 Northeast Atlantic Ocean, the Mediterranean Sea and Macaronesia with their distribution records, 158 their likely origin and putative introduction vectors. These data are used to provide a quantitative ¹⁵⁹assessment of the spatio-temporal dynamics of primary and secondary introductions and to detect

160 shortcomings in the monitoring and legislation required to tackle the introduction of non-indigenous

161 species more effectively.

163 Materials and methods

¹⁶⁴**Data compilation**

We compiled a database of non-indigenous marine seaweed species records from three regions, 166 namely the Northeast Atlantic Ocean (excluding Greenland), the Mediterranean Sea and ¹⁶⁷Macaronesia (Fig. 2). For some of the analyses we subdivided the Northeast Atlantic Ocean into ¹⁶⁸Lusitania and Northern Europe and the Mediterranean Sea into a Western and Eastern part. With 169 respect to Macaronesia, the compilation includes records from the Azores, Canary Islands, Madeira 170 and the Salvagen Islands, but not Cape Verde. The dataset builds on previous lists by Mineur *et al.* ¹⁷¹(2010) and Verlaque *et al.* (2015), and includes published records of species occurring in a natural 172 environment and flagged as non-indigenous in the study area irrespective of taxonomic confidence 173 and biogeographic status (see below). In addition, we included unpublished records produced by 174 various research projects conducted by, amongst others, the Station Biologique de Roscoff (France), 175 National Biodiversity Data Centre (Ireland), Stichting ANEMOON (the Netherlands), Scottish 176 Natural Heritage (Scotland), the ICES Working Group on Introductions and Transfers of Marine 177 Organisms 2004, as well as the European Alien Species Information Network (EASIN, 2022) 178 records, collection data, GBIF records and personal data. All records were added to the database 179 under the name they were reported as. Names were updated according to the most recent taxonomic 180 consensus (AlgaeBase, Guiry & Guiry, 2023).

181 The species listed as non-indigenous include those that are naturalised (i.e. having 182 established permanent, self-maintaining populations), as well as species for which no information is 183 available on population status (i.e. species referred to as 'alien' by Verlaque *et al.* 2015). Species ¹⁸⁴that have been demonstrated to be misidentifications or unsupported records are excluded from the 185 list. To promote consistency in definitions and criteria used to determine whether a species is non-¹⁸⁶indigenous, we have adopted the criteria for assessing the biogeographic status proposed by Essl *et* ¹⁸⁷*al.* (2018) (Fig. 3). This framework stresses 1) the need for crossing a biogeographic barrier, 2) the ¹⁸⁸involvement of direct or indirect human agencies in the physical movement of individuals, spores or ¹⁸⁹fragments, and 3) the ability of the species to reproduce without human assistance in the introduced 190 range. The combination of these criteria excludes records of species which are in the process of

191 expanding their range naturally, for example as a result of global warming. Species entering the Mediterranean Sea through the Suez Canal (i.e. Lessepsian migrants), on the other hand are 193 considered non-indigenous because of the anthropogenic nature of the dispersal corridor. In contrary, species entering the Mediterranean Sea through the strait of Gibraltar , without a human-195 vector, are not considered as non-indigenous. The dataset also includes species indigenous to the study area that have demonstrably become displaced within he study area as a result of human-mediated exchanges. Examples include exchanges of species between Atlantic and Mediterranean shores. Species for which the area of origin is unknown are assigned as 'cryptogenic' (sensu Carlton, 1996). In cases where there is not sufficient information to be conclusive on their biogeographic status, species are labelled as 'data deficient'. Species with low uncertainty, for which there is no doubt about their non-indigenous status, have been labelled 'non-indigenous'.

202 Added to these criteria but highly relevant with respect to seaweeds, where a solid 203 taxonomic framework is often lacking for many taxa, is taxonomic confidence. We assigned a 204 'high-confidence score' (score $= 1$) to accepted nominal species that had not been shown to be a 205 species complex based on molecular studies in their introduced or native ranges. A high score was 206 also assigned to species for which, so far, there is sufficient confidence in unambiguous 207 identification based on morphology. We acknowledge, however, that the latter does not rule out the 208 potential existence of cryptic species hidden under the accepted nominal species. Conversely, ²⁰⁹species that belong to an understudied complex of cryptic species were assigned a low-confidence 210 score (score $= 0$). A cryptic species is defined here as a taxon composed of two or more species that 211 have been classified as a single nominal species, because they were initially not distinguished based 212 on their morphological characteristics (Bickford *et al.*, 2007; Pante *et al.*, 2015).

²¹³Recognising we cannot be conclusive about the non-indigenous status of many seaweed ²¹⁴species, we explicitly acknowledge the uncertainty in the assessment of the taxonomic as well as 215 biogeographic status of putative non-indigenous seaweeds in the study area (Fig. 3). The status of 216 each species is concisely described in Suppl. Material Table S1.

²¹⁷For every species we determined the year when the species was first reported in the 218 Northeast Atlantic Ocean, Mediterranean Sea and/or Macaronesia. Where possible, this date refers 219 to the year the species was detected (i.e. collection date) rather than when the record was published 220 (i.e. publication date). We acknowledge that detection dates may not portray the actual date the 221 species was introduced. For each species an estimate is provided for its native biogeographic range. 222 If the native range could not be assessed, we indicated 'uncertain'. The putative distribution of the

223 species was based on literature reports included in AlgaeBase (Guiry & Guiry, 2023). Species traits

²²⁴(e.g. thallus size) were obtained from AlgaeTraits (Vranken *et al.*, 2022). For spatial and temporal

225 analyses, distribution records were filtered on a combination of unique year, coordinates and

226 species name to eliminate potential duplicate records. The complete dataset has been archived at

227 Zenodo and is available at DOI: 10.5281/zenodo.7798640. This dataset contains the following

228 information for each record: currently accepted scientific name, the scientific name under which it

²²⁹was originally reported, year of record, location, country, coordinates and reference.

²³¹**Results and Discussion**

²³²A total of 19,724 records of non-indigenous seaweeds were collected dating from 1808 to 2022 ²³³(Fig. 2). Of these, 17,104 were retained after removing duplicates and incomplete data. The ²³⁴geographical distribution of the records highlights considerable sampling of non-indigenous 235 seaweeds from all coastlines in the study area (Fig. 2). The list contains 227 species (Table 1). Non-236 indigenous species make up approximately 10% of the seaweed flora in the Mediterranean Sea, 6% 237 in the Northeast Atlantic Ocean and 4% of the Macaronesian flora. The total number of 227 238 includes all species regardless of taxonomic and biogeographic uncertainty. For 84 species neither 239 their non-indigenous status nor their taxonomy is challenged (Fig. 4). These species make up 83% 240 of the distribution records in the database. Half of the distribution records can be attributed to only ²⁴¹five species (*Sargassum muticum*, *Bonnemaisonia hamifera*, *Asparagopsis armata*, *Caulerpa* ²⁴²*cylindracea* and *Colpomenia peregrina*). Fifty-six species are most likely non-indigenous, but 243 decisions are hampered by taxonomic uncertainties (Fig. 4). On the other hand, 87 of the 227 244 species have a cryptogenic or data deficient status (30 species with an uncertain biogeographic 245 status, and 57 species for which both the geographic status and taxonomic confidence are 246 uncertain), meaning that the evidence for a non-indigenous status is mediocre to weak (Fig. 4).

²⁴⁷We emphasise that the categorisation of putative non-indigenous species according to ²⁴⁸biogeographic uncertainty and taxonomic confidence emerged as a consensus among the authors of 249 this paper. A literature search will undoubtedly reveal several additional species names that could 250 potentially be added to the list of cryptogenic or data deficient species. However, there is little 251 added value in incorporating species names which are wholly unsupported or most likely result 252 from misidentifications or other mistakes. Evidently, both taxonomic and biogeographic uncertainty 253 plague the compilation of lists and databases of non-indigenous species. Below we discuss how the

254 level of sophistication of systematic and biogeographic knowledge translates to uncertainty in the

255 number of non-indigenous seaweeds in the study area.

257 Taxonomic confidence

258 For 140 species the non-indigenous nature of the species itself is not disputed. However, the reliable 259 identification of 56 of those species is challenging, and therefore their current distribution as well as 260 their putative region of origin are questionable. In most cases this uncertainty can be attributed to a 261 poorly established taxonomic framework. Taxonomic uncertainty is rife in seaweeds. In the absence 262 of DNA sequence data the identification of many seaweed species is particularly difficult (e.g. Van ²⁶³Oppen *et al.*, 1996; Maggs *et al.*, 2007; Cianciola *et al.*, 2010; De Clerck *et al.*, 2013; Verbruggen, 264 2014). Taxonomic uncertainty is much higher among small-sized species: 67% of species smaller 265 than 5 cm are flagged as taxonomically uncertain, compared to 34% of species larger than 5 cm. Of 266 the taxa larger than 5 cm with high taxonomic uncertainty are many that belong to genera that are ²⁶⁷widespread in tropical and warm-temperate regions (e.g. *Avrainvillea, Caulerpa, Codium,* ²⁶⁸*Dichotomaria, Ganonema, Hypnea*). From a biogeographic perspective, taxonomic uncertainty 269 plagues "only" 20% of species with a Northwest Pacific origin (11 of 56 species), but 62% of 270 species with a likely Lessepsian or tropical Indo-Pacific origin (26 of 42 species) (Table 1).

271 Recent advances in the taxonomy of several genera, nearly always assisted by DNA 272 sequence data, have demonstrated that many so-called wide-ranging (or cosmopolitan) seaweeds 273 actually consist of species complexes of morphologically almost indistinguishable species ²⁷⁴(pseudocryptic species), or even truly cryptic species which are indistinguishable based on 275 morphological criteria. The individual species are often confined to specific geographic areas (e.g. ²⁷⁶Won *et al.*, 2009; Vieira *et al.*, 2017; Diaz-Tapia *et al.*, 2018; Leliaert *et al.*, 2018; Diaz-Tapia *et al.*, 277 2020). A more refined taxonomic framework therefore alters our understanding of the biogeography 278 of the species in many cases and consequently our interpretation of their native versus non-279 indigenous status. The *Caulerpa racemosa* complex is highly representative of how evolving 280 insights into species diversity alter our views of the taxa being non-indigenous in the study area. ²⁸¹While initially *Caulerpa* specimens with vesiculate branchlets collected in the Mediterranean Sea ²⁸²were identified as *C. racemosa*, the latter proved to be a complex of at least eight species, three of ²⁸³which (*C. chemnitzia*, *C. cylindracea* and *C. requienii*) are currently considered non-indigenous in ²⁸⁴the Mediterranean Sea (Verlaque *et al.*, 2000; Verlaque *et al.*, 2003; Draisma *et al.*, 2014; Verlaque ²⁸⁵*et al.*, 2015). Similarly, a better understanding of the taxonomy of foliose *Grateloupia* species

286 resulted not only in the recognition that *G. turuturu* was introduced in the study area from the

²⁸⁷Northwest Pacific, as opposed to *G. doryphora* whose distribution is likely to be restricted to the

288 Pacific coast of South America (Gavio & Fredericq, 2002), but also revealed that so-called non-

289 indigenous foliose *Grateloupia* species in the study area were actually a mixture of two non-

²⁹⁰indigenous species, *G. lanceolata* and *G. turuturu,* and a native species, *G. lanceola*, which had

²⁹¹been regarded a synonym of *G. doryphora* (Verlaque *et al.*, 2005; Figueroa *et al.*, 2007).

292 In many other instances, however, conspecificity of populations from the native and non-293 native regions remains to be demonstrated. There are also examples where several non-indigenous ²⁹⁴species are thought to be conspecific by some authors but regarded as distinct species by others. For 295 example, some authors consider *Antithamnion hubbsii* distinct from *A. nipponicum* (Athanasiadis, 296 1996), while others treat the former as a synonym of the latter (e.g. Kim & Lee, 2012). Similarly, 297 records of *Polysiphonia morrowii* and *P. senticulosa* likely belong to the same species, even though 298 both species are regarded as distinct (D'Archino *et al.*, 2013; Stegenga & Karremans, 2015; ²⁹⁹Piñeiro-Corbeira *et al.*, 2020). Given the widespread nature of cryptic and pseudocryptic diversity 300 in seaweeds, continuous efforts of DNA-assisted identifications through Sanger sequencing will

301 probably continue to revise our view on non-indigenous species.

³⁰²Although DNA sequence data are in many cases a great help in verifying species identities, 303 this does not mean DNA solves every single problem like a magic wand. Apart from reference 304 sequences in repositories not being available or reliable, patterns of genetic divergence can be 305 complicated and prone to different interpretations. For example, differences in the interpretation of ³⁰⁶genetic patterns and species boundaries in the genus *Melanothamnus* led to the recognition of a ³⁰⁷single species, *M. harveyi* s.l. (McIvor *et al.*, 2001) or by contrast to the recognition of at least three ³⁰⁸separate species, including *M. akkeshiensis*, *M. japonicus* and *M. harveyi* s.s. (Savoie & Saunders, ³⁰⁹2015). The narrower species concept would result in an interpretation whereby *M. harveyi* is native 310 to the Northeast Atlantic Ocean rather than a non-indigenous species introduced to the study area 311 from the Northwest Pacific Ocean. Under the alternative scenario which recognises a single ³¹²genetically diverse species, *M. harveyi* is widely distributed globally with both cryptogenic and ³¹³non-indigenous haplotypes in the study area (Piñeiro-Corbeira *et al.*, 2019). One should note that 314 despite the availability of a good number of sequences of these species/haplotypes, the potential 315 native area of the species (Northwest Pacific Ocean) has been scarcely sampled. Therefore, it is still 316 possible that *M. harveyi* s.s. can be present in this region but remained undetected. Distribution ³¹⁷records of *M. harveyi* and *M. japonicus* are included as *M. harveyi/japonicus* in our dataset.

319 Biogeographic uncertainty

³²⁰A lack of baseline data with respect to the global distribution of seaweeds is the major contributor

321 to biogeographic uncertainty reported for 87 taxa (Fig. 4). Brown and green seaweeds display

³²²slightly less biogeographic uncertainty, 28% and 34%, respectively, compared to 42% for red

- ³²³seaweeds. Baseline data of seaweed diversity along coastlines in the study area as well as the
- 324 putative native regions in the form of herbarium collections, censuses and historical checklists can
- ³²⁵serve as a reference for the presence of species in a given area. Here as well, low confidence in the
- 326 taxonomy and identification of seaweeds makes the interpretation of species lists exceedingly
- 327 difficult. If a species is not reliably identified, its distribution is not reliable. As a result,
- 328 biogeographic and taxonomic uncertainties usually go hand-in-hand. Three different categories of
- 329 factors that lead to biogeographic uncertainty are discussed below.

³³⁰**Pseudo-indigenous species**

- ³³¹Several seaweed species have been described from the study area that were presumed native, but
- 332 later turned out to be non-indigenous species. Carlton (2009) named such species pseudo-
- ³³³indigenous. For example, *Dictyota cyanoloma* was described as a new species from the
- ³³⁴Mediterranean Sea and Macaronesia (Tronholm *et al.*, 2010), but subsequent collecting efforts
- 335 revealed that the species most likely represents a cryptic introduction (Aragay Soler *et al.*, 2016;
- ³³⁶Steen *et al.*, 2017; Tran *et al.*, 2021). Similarly, *Porphyra olivii* described from Greece (Brodie *et*
- ³³⁷*al.*, 2007a) turned out to be conspecific with *Neopyropia koreana* (Vergés *et al.*, 2013; Yang *et al.*,
- ³³⁸2020), a species native to the Northwest Pacific. Such insights invariably result from DNA-assisted
- 339 species identification and subsequent interpretations of biogeographic patterns. Hereby widely
- 340 disjunct distribution ranges are interpreted as non-natural and therefore the result of human-
- 341 mediated dispersal.

³⁴²Evidently, determining the non-indigenous nature of a species becomes more difficult for 343 historic introductions. In such cases we fully rely on DNA signatures which can point toward a non-344 indigenous nature of the species. DNA-assisted identification of historic voucher specimens of ³⁴⁵*Codium fragile* revealed that the invasive (sub)species was already introduced into the study area as 346 early as 1845 (Provan *et al.*, 2008), which is roughly a century before phycologists realised the 347 species was actually native to the Northwest Pacific Ocean and non-indigenous to the study area as 348 well as several other parts of the world. In the case of *Cutleria multifida*, described from Norfolk,

³⁴⁹England as early as 1801, genetic signatures point toward an introduction of the Mediterranean Sea

350 populations from the Northwest Pacific. The Northeast Atlantic Ocean populations, however, are

351 genetically more diverse and well-differentiated from those in Japan and are therefore considered

³⁵²native (Kawai *et al.*, 2016). It remains to be determined if *Cutleria multifida* is native to the

353 Northeast Atlantic Ocean as well as the Pacific, or whether an even more complex history of

354 historic introductions underlies this pattern.

³⁵⁵**Discerning natural from human-mediated dispersal**

356 Eventually, the possibility of introductions needs to be evaluated against historic and ongoing ³⁵⁷natural dispersal events. The recent observation of *Flabellia petiolata* from the south coast of ³⁵⁸England confronts researchers with exactly this question (Díaz-Tapia *et al.*, 2020). Despite a long 359 tradition of seaweed studies and regular surveys, *F. petiolata* was never recorded from the British 360 Isles prior to 2013. The closest populations of the species are found in the Mediterranean Sea and 361 the Canary Islands. A recent introduction would be the most obvious explanation. However, the ³⁶²English populations of *F. petiolata* could also be interpreted as a relic of a formerly more 363 widespread Atlantic Ocean distribution. The species' range might have been continuous during 364 warmer periods in the Holocene, but persisted in the Northeast Atlantic Ocean in a handful of ³⁶⁵refugia during colder periods. Afterall, several native species, e.g. *Cladophora battersii, Codium* ³⁶⁶*bursa* and *Halopithys incurva* display similar distribution patterns (Maggs & Hommersand, 1993; ³⁶⁷Brodie *et al.*, 2007b).

Quaternary climatic cycling probably also facilitated dispersal of temperate species across the tropical Atlantic Ocean connecting southern Africa with Europe. The presence of *Schizymenia apoda* in the Azores, the British Isles and Namibia may have resulted from natural amphi-equatorial dispersal events in recent geological times, but also a human-mediated introduction in the Atlantic Ocean cannot be ruled out given the presence of *S. apoda* in Australia and China (Gabriel *et al.*, 2019; Gunnarsson *et al.*, 2020). Natural dispersal events from the Northeast Pacific Ocean to 374 Northern Europe through the Bering Strait may be difficult to discern from introductions (Lindstrom, 2001; Bringloe & Saunders, 2019). For instance, *Schizymenia jonssonii*, a species 376 recently described from Iceland, may have colonised the northern Atlantic Ocean naturally via the Bering Strait but it is equally possible the species is a relatively recent introduction (Gunnarsson *et al.*, 2020).

379 Population-level sampling and the application of genetic markers with sufficient

380 intraspecific resolution (e.g. fast evolving spacer regions, microsatellite markers or SNP data) have

381 the potential to shed light on natural versus human-mediated dispersal events, and more generally to

382 help in reconstructing introduction history (Viard $&$ Comtet, 2015), but are rarely used in studies of

383 putative seaweed introductions. Notable exceptions include the invasive history of *Fucus* species,

³⁸⁴*Sargassum muticum* and *Gracilaria vermiculophylla*.

³⁸⁵Coyer *et al.* (2011) demonstrated a North Pacific origin of *Fucus distichus* followed by at ³⁸⁶least two separate colonisation events of the North Atlantic Ocean prior to the last glacial 387 maximum, which makes this species native to Europe. However, the taxon, having a predominantly 388 northern distribution, was accidentally introduced in the Oslofjord followed by further expansion in the Kattegat region as a result of an introduction event in the late $19th$ century (Coyer *et al.*, 2002, as ³⁹⁰*F. evanescens*, currently regarded as a subspecies of *F. distichus*). *Fucus serratus* was also exported 391 from mainland Europe to Atlantic North America, Iceland and the Faroes (Coyer *et al.*, 2006; ³⁹²Brawley *et al.*, 2009). Discharging of ballast stones in destination harbours is considered the prime source of introductions in Atlantic North America and Iceland in the 19th century. The *F. serratus* 394 population in the Faroes is of more recent origin (late $20th$ century) and was most likely introduced ³⁹⁵from Iceland (Coyer *et al.*, 2006). Many marine benthic organisms including seaweeds but also ³⁹⁶many invertebrates display amphi-Atlantic distribution (Haydar, 2012). For such disjunct 397 distributions, distinguishing scenarios of post-glacial relicts or natural long-distance dispersal from ³⁹⁸human-assisted dispersal (and introduction) is challenging and most often requires a combination of 399 life-history traits assessment and high resolution molecular markers. The power of genome-wide ⁴⁰⁰genetic variation was demonstrated for *Sargassum muticum* (Le Cam *et al.*, 2020); whereas 401 microsatellite markers failed to reveal any genetic variation in the invaded range of the species, a 402 panel of single-nucleotide polymorphisms (SNPs) obtained from ddRAD sequencing confirmed a 403 secondary introduction to the Northeast Atlantic Ocean from the Northeast Pacific Ocean, but also ⁴⁰⁴revealed two additional cryptic introductions to Europe. Similarly, Krueger-Hadfield *et al.* (2017) 405 identified the areas in the native region that most likely contributed to the European invasions of the ⁴⁰⁶red alga *Gracilaria vermiculophylla*. Subsequent work used SNPs to refine the origins and 407 understand evolution during invasion (Flanagan *et al.*, 2021).

408 In Macaronesia, several confounding factors further complicate the interpretation of the non-⁴⁰⁹indigenous nature of species. The geographical location of Macaronesia, bordering the tropical 410 Atlantic Ocean, contributes significantly to this difficulty in interpretation. Several tropical and

411 subtropical taxa are, probably erroneously, attributed a pantropical distribution, which not only contributes to high taxonomic uncertainty, but the latter also translates into biogeographic uncertainty. In addition, it is not always evident to preclude natural dispersal to explain the presence of particular species. *Halimeda incrassata*, a species that naturally occurs in the tropical western Atlantic Ocean (Verbruggen *et al.*, 2006), was reported from Porto Santo, Madeira, by Wirtz & 416 Kaufmann (2005) and more recently from the Balearic Islands by Alós et al. (2016). Further surveys indicated the species is also present in the Canary Islands (Sangil *et al.*, 2018) and Azores (Costa *et al.*, 2017). Even though in the Mediterranean Sea the species displays typical invasive 419 behaviour, the involvement of human activities in its establishment is not clear. Another species 420 from tropical western Atlantic Ocean, *Caulerpa ashmeadii*, was recently reported from Porto Santo, Madeira, and may represent a similar case of natural range expansion across the Atlantic (Ribeiro *et al.*, 2023). Amphi-Atlantic Ocean distributions have been confirmed using molecular markers for several seaweed taxa (*Cladophoropsis membranacea*, Leliaert *et al.*, 2009; *Laurenciella marilzae*, Cassano *et al.*, 2012; *Laurencia catarinensis*, Machin-Sanchez *et al.*, 2012; Tronholm *et al.*, 2012; *Vertebrata foetidissima*, Díaz-Tapia *et al.*, 2013; *Dictyota* spp., Tronholm *et al.*, 2013; *Caulerpa prolifera*, Varela-Álvarez *et al.*, 2015; *Laminaria digitata*, Neiva *et al.*, 2020; *Lobophora* spp., Vieira *et al.*, 2020). In these examples presumed natural distribution ranges have not been

428 challenged.

Also of note is that Macaronesia covers a large geographic area, encompassing several biologically diverse archipelagos. Several taxa that have long been reported from the Canary Islands 431 and Madeira and are considered native in those areas, have recently been reported from the Azores. The geographic position of the Azores and the actual oceanographic current circulation in the North Atlantic Ocean would not seem favourable for natural range expansions from the Canary Islands and Madeira. In some cases, initial reports from anthropogenic habitats, such as harbour environments (e.g. *Caulerpa webbiana*), favour the hypothesis of an anthropogenic factor in the range expansion, although evidence is lacking for several other species which are presumed non-indigenous (e.g. *Halimeda incrassata*, *Xiphosiphonia pinnulata*, *Hypoglossum heterocystideum*).

A final category of uncertainty in natural vs. human-mediated dispersal mechanisms concerns 439 those non-indigenous species which have been displaced within the study area. The biogeographic history of the Mediterranean Sea biota is closely intertwined with the Northeast Atlantic Ocean to which it is connected by the narrow Strait of Gibraltar (Bianchi & Morri, 2003; Patarnello *et al.*, 2007; Le Gall *et al.*, 2021), which results in a subset of species being shared between both regions. However, several Northeast Atlantic Ocean species have been recently introduced into the

Mediterranean Sea, often in lagoons with extensive aquaculture facilities, e.g. *Ascophyllum*

nodosum, *Chorda filum*, *Fucus spiralis* and *Grateloupia minima* (Petrocelli *et al.*, 2013). In some

446 cases however, patterns become more complex, for example, when native and non-indigenous

populations co-occur as is the case for *Chondria coerulescens*, *Vertebrata fucoides* and possibly

also *Ganonema farinosum* (Verlaque *et al.*, 2015). It is worth remembering that many

Mediterranean Sea species naturally dispersed from the Atlantic Ocean after the Zanclean flood

450 which occurred after the Messinian salinity crisis about 5.33 myr (Blondel *et al.*, 2010).

Species of unresolved origin

In several cases the non-indigenous nature of certain seaweeds remains unresolved. Some recently

described species in the study area, such as *Polysiphonia radiata* and *P. delicata*, are mainly known

454 from marinas and are probably non-indigenous but their origin remains unknown because it is likely

that they remained undescribed in their native area (Díaz-Tapia *et al.*, 2017). Morphological

similarity of putative non-indigenous species to native species can also complicate interpretation of

non-indigenous patterns as demonstrated by *Anotrichium furcellatum*. The latter was originally

458 described from Naples, but considered non-indigenous in the Northeast Atlantic Ocean. The

Mediterranean Sea populations, however, may have been largely replaced by a cryptic introduction

460 of *A. okamurae*, originally from the Northwest Pacific Ocean (Verlaque *et al.*, 2015). The status of

A. furcellatum and *A. okamurae* has not yet been tested with molecular data.

Similar arguments could be made for species that are considered native in the study area. In the case of *Lobophora delicata*, which is not considered as non-indigenous, a lack of baseline data makes it difficult to be conclusive on its status as a native species. As pointed out by Vieira *et al.* (2019), the first records of *Lobophora* in the Mediterranean Sea date back to 1955 (Edelstein, 1960), which is surprising for a distinctive seaweed genus which can be easily found in many places 467 growing at a depth of 0.5 m. In contrast, other genera of Dictyotales were invariably reported from the Mediterranean Sea in the 18th or early 19th century. Has *L. delicata* been overlooked or does 469 the late discovery of the species correspond to a more recent introduction? Without proper baseline 470 data, e.g. herbarium records, this is difficult to test, and if the species does not display typical 471 invasive behaviour its native status may simply never come into question.

472 A puzzling case is presented by several taxa with clear Indo-Pacific affinities which 473 appeared in the Mediterranean Sea prior to the opening of the Suez Canal in 1869, e.g.

Acanthophora nayadiformis, Asparagopsis taxiformis and *Ganonema farinosum.* For instance, *Asparagopsis taxiformis* was first described from Alexandria in the Mediterranean Sea as *Fucus taxiformis* Delile (1813), and thus reported as a native species. However, further molecular work 477 revealed that this accepted species was made of five distinct lineages, possibly corresponding to two cryptic species (Ní Chualáin *et al.*, 2004; Andreakis *et al.*, 2007; Dijoux *et al.*, 2014), one of them 479 presumably present in the Mediterranean Sea prior to the opening of the Suez Canal, and one more recently introduced. Similar complexity was revealed for the closely related species *Asparagopsis armata*, supposedly introduced in the study area, for which novel sampling in the South Pacific 482 Ocean showed the existence of two highly divergent clades, presumably corresponding to two 483 cryptic species, one of them distributed in Europe, South Africa and Tasmania, and one restricted (so far) to Western Australia, New Zealand and Tasmania (Dijoux *et al.*, 2014). Such cases highlight the difficulty in establishing whether a species is non-indigenous in the absence of large

486 sampling encompassing the global distribution of the targeted presumably non-indigenous species.

488 The spatial patterns and origins of non-indigenous seaweeds

⁴⁸⁹Analysis of the distribution of non-indigenous seaweeds in the study area reveals clear patterns in 490 richness and the number of species shared among regions. The large-scale spatial patterns are 491 discussed below in a context of dispersal vectors that determine spread and establishment of non-492 indigenous species.

⁴⁹³The Eastern Mediterranean Sea is home to the highest number of non-indigenous seaweeds ⁴⁹⁴(77 species), followed by the Western Mediterranean Sea (47 species) and Lusitania (45 species). ⁴⁹⁵Macaronesia and Northern Europe harbour somewhat lower numbers (36 and 40 species, 496 respectively). These numbers refer to high-confidence non-indigenous species only. Adding ⁴⁹⁷cryptogenic and data deficient species further underscores the higher number of non-indigenous 498 species in the Eastern Mediterranean Sea. In the latter region an extra 47 species are flagged as ⁴⁹⁹cryptogenic or data deficient, which is considerably higher compared to the other regions which 500 typically host 20 or less cryptogenic and data deficient species. In all regions, most of the non-501 indigenous seaweeds belong to Rhodophyta (between 63-76% of the species non-indigenous in each 502 region), followed by brown seaweeds (18-22%), while green seaweeds contribute to 8-18% of the 503 species recorded (Fig. 5).

⁵⁰⁴We did not detect a significant correlation between the number of non-indigenous species 505 and the number of records (Pearson correlation $= -0.18$, p-value $= 0.74$), which indicates that 506 differences of non-indigenous species between regions are not a mere artefact of sampling eff differences of non-indigenous species between regions are not a mere artefact of sampling effort. ⁵⁰⁷The number of non-indigenous species also does not scale with the length of the coastline (Pearson 508 correlation = 0.10, p-value = 0.88). As will be argued below, the number of non-indigenous species 509 in a given region and the fraction of species shared between regions is a complex function including 510 the efficiency of primary and secondary dispersal vectors combined with abiotic (and potentially 511 biotic) ecological factors that determine the establishment of non-indigenous species in the recipient 512 ecosystems (reviewed in Maitner *et al.*, 2021).

The Mediterranean Sea and the Northeast Atlantic Ocean share 45 high-confidence non-514 indigenous species, while Macaronesia shares roughly an equal number of non-indigenous species 515 with the Mediterranean Sea (24 species) and the Northeast Atlantic Ocean (21 species) (Fig. 6A). Within the Northeast Atlantic Ocean, Macaronesia and Lusitania share 22 high-confidence non-indigenous species, while Macaronesia and Northern Europe share none other than the 14 non-518 indigenous species present in all three Northeast Atlantic regions (Fig. 6B). A relatively low 519 number, 18 high-confidence non-indigenous species out of 140, are shared between the Northeast 520 Atlantic Ocean, Macaronesia and the Mediterranean Sea (Fig. 6A). The broad distribution of these 521 non-indigenous species is noteworthy for it reflects a very wide amplitude in abiotic and biotic parameters. At least nine of these widely distributed non-indigenous species (*Antithamnion hubbsii/nipponicum, Antithamnionella spirographidis, Asparagopsis armata, Bonnemaisonia hamifera, Codium fragile* subsp. *fragile, Colpomenia peregrina, Dictyota cyanoloma, Grateloupia turuturu, Scytosiphon dotyi*) are reported from all five regions. The remaining species have a more 526 restricted distribution range, being only present in the three central regions Macaronesia, Lusitania and the Western Mediterranean Sea (*Phycocalidia suborbiculata, Spongoclonium caribaeum, Symphyocladia marchantioides*), or being absent from either of the two peripheral regions, i.e. the Eastern Mediterranean Sea (*Antithamnionella ternifolia, Sargassum muticum*) or Northern Europe (*Asparagopsis taxiformis, Rugulopteryx okamurae, Pachymeniopsis gargiuloi*).

531 The number of high-confidence non-indigenous species currently only found in the ⁵³²Mediterranean Sea (66 species) is striking. Including cryptogenic or data deficient species brings 533 this number to 115. However, there is considerable differentiation of non-indigenous species 534 between the Western and Eastern Mediterranean Sea. Only 34 non-indigenous species are shared 535 between both regions, representing 38% of the total non-indigenous species diversity in the

536 Mediterranean Sea (Fig. 6C). In addition, the fraction of non-indigenous species unique to the

⁵³⁷Eastern Mediterranean Sea (27 species) is considerably larger than for the Western Mediterranean

538 Sea (8 species) (Fig. 6C). The latter pattern is largely the result of dispersal of warm-adapted

539 species from the Red Sea and by extension the Indo-Pacific Ocean via the Suez Canal. At present,

540 only a fraction of these have spread to the Western Mediterranean Sea resulting in a higher diversity

541 of non-indigenous species in the Eastern Mediterranean Sea.

542 The non-indigenous species reported from the Thau Lagoon in France, and the Mar Piccolo 543 and Venice Lagoon in Italy are quite distinct compared to those of surrounding Mediterranean ⁵⁴⁴waters. A combination of anthropogenic disturbances and intense aquaculture activities (trade and 545 exchanges), more specifically import of shellfish, has resulted in a very high diversity of non-546 indigenous species in these lagoon systems (66 species), comparable to that of the surrounding 547 Mediterranean Sea locations despite their much smaller area. The non-indigenous species in ⁵⁴⁸Mediterranean lagoons have, moreover, more affinities with the Atlantic Ocean than with the ⁵⁴⁹Mediterranean Sea, likely to be due to exchanges between shellfish production areas (see below). 550 Nearly half of the non-indigenous species encountered in Mediterranean aquaculture lagoons are 551 not (yet) reported from surrounding Mediterranean water, while 36 are shared with the Northeast 552 Atlantic Ocean. Of these 36 non-indigenous species, 13 have only been recorded in aquaculture ⁵⁵³lagoons within the Mediterranean Sea. These mainly include species with a Northwest Pacific ⁵⁵⁴Ocean origin, e.g. *Dasysiphonia japonica*, *Neopyropia yezoensis* and *Nitophyllum* ⁵⁵⁵*stellatocorticatum*. Differences in the abiotic physico-chemical environment between ⁵⁵⁶Mediterranean lagoons and surrounding coastlines probably underlie the failure of these species to 557 spread widely in the Mediterranean Sea. *Rugulopteryx okamurae*, however, presents a striking 558 counterexample of this trend. The species was collected in the Thau Lagoon for the first time in ⁵⁵⁹2002 (Verlaque *et al.*, 2009). Initially *R. okamurae* appeared to be rather non-invasive, but in 2015 560 it was reported from the Strait of Gibraltar (Ocaña *et al.*, 2016; El Aamri *et al.*, 2018; García-561 Gómez *et al.*, 2020), where the species forms dense stands rapidly overgrowing most native 562 seaweed species. More recently the same alarming invasive behaviour of *R. okamurae* has been ⁵⁶³noted in the Marseille area (Ruitton *et al.*, 2021) as well as southwest Portugal (Liulea *et al.*, 2023) 564 and Macaronesia (Faria et al., 2022).

⁵⁶⁵A Northwest Pacific origin of the largest part of the non-indigenous species present in the ⁵⁶⁶Mediterranean lagoons and Northern Europe is well established (Fig. 7, Table 1) (Boudouresque *et* ⁵⁶⁷*al.*, 2010). Regular monitoring and surveys demonstrated that many of those species had been first

568 accidentally introduced in the Mediterranean lagoons (notably the Thau lagoon) and were then 569 transported to the Northeast Atlantic Ocean hitchhiking with oyster transfers (Mineur *et al.*, 2007a). ⁵⁷⁰*Undaria pinnatifida* presents a notable exception to this pattern. Following its accidental 571 introduction in the Thau lagoon, the species was deliberately introduced in Brittany for aquaculture 572 purposes (Floc'h *et al.*, 1991) from which it rapidly spread and established itself as one of the 573 dominant non-indigenous species in artificial as well as natural habitats in the Northeast Atlantic ⁵⁷⁴Ocean (Voisin *et al.*, 2005; Guzinski *et al.*, 2018). Note that commercial transfers of oysters 575 between the Mediterranean Sea and the Atlantic coasts of France is still fully allowed under French 576 and European regulations, which results in a quasi-continuous series of secondary introduction 577 events. Aquaculture- and fisheries-associated transport (e.g. nets, packing material) between the 578 Atlantic Ocean and Mediterranean Sea is also undoubtedly responsible for the introduction of a 579 range of native Atlantic Ocean species to the Mediterranean lagoons (e.g. *Ascophyllum nodosum*, ⁵⁸⁰*Chorda filum, Grateloupia minima, Fucus spiralis* and *Vertebrata fucoides*).

581 The Mediterranean Sea, Macaronesia and Lusitania are also home to 35 high-confidence 582 non-indigenous species with presumed or established Australasian origin (Fig. 7). Species with 583 Australasian origin are much less represented in Northern Europe, reflecting the warm temperate 584 nature of these species. The introduction vectors for this category of species remain, however, ⁵⁸⁵largely elusive. For *Acrothamnion preissii* and *Womersleyella setacea* ship traffic has been 586 suggested as vector based on their first observation close to a major harbour (Livorno, Italy), but 587 accidental release from scientific laboratories and public or private aquaria is also a possibility. As ⁵⁸⁸with the notorious case of *Caulerpa taxifolia* (Verlaque *et al.*, 2015), aquarium releases have likely 589 resulted in the introduction of other seaweeds, mainly in the Mediterranean Sea (Vranken *et al.*, ⁵⁹⁰2018).

591 Despite the abovementioned clear categories of non-indigenous species and associated 592 pathways, for circa one third of the species there is considerable uncertainty regarding the area of 593 origin and the potential dispersal vectors. Complicating identification of native range and vectors 594 even further, population-level molecular studies on several non-indigenous species have unveiled 595 multiple independent introductions possibly involving different vectors (McIvor *et al.*, 2001; Voisin ⁵⁹⁶*et al.*, 2005; Geoffroy *et al.*, 2016; Le Cam *et al.*, 2020).

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⁵⁹⁹**Introduction hotspots**

600 Of the 140 high-confidence non-indigenous species, 65% have been reported for the first time in the ⁶⁰¹Mediterranean Sea, 26% in the Northeast Atlantic Ocean and 9% in Macaronesia. The distributions 602 of the first record of each species in the study area underscore the importance of aquaculture for ⁶⁰³introductions of seaweeds. The Thau lagoon, with 30 reports of first introductions in the study area, 604 is one of the major introduction hotspots. In total 58 species, constituting 32% of the total seaweed ⁶⁰⁵diversity or 48-99% of the biomass, have become established in this coastal lagoon (Boudouresque ⁶⁰⁶*et al.*, 2010). The Thau lagoon is an important centre of oyster cultivation in the Mediterranean Sea. ⁶⁰⁷However, the oyster farmers rely on the import of oyster spat produced in other regions because the 608 lagoon is not particularly suitable for oyster reproduction. Since 1977, only spat of Pacific oysters ⁶⁰⁹spat produced in the Atlantic Ocean is allowed to be laid in the French Mediterranean lagoons ⁶¹⁰(Verlaque *et al.*, 2007). However, it is likely that some non-official imports from outside of Europe 611 occur, as reported by Verlaque (1996). These continuous transfers across different biogeographic 612 regions result in astonishingly high numbers of non-indigenous species. A low native diversity due 613 to the low occurrence of natural hard substrata in lagoons and relatively recent construction of hard 614 substrata for aquaculture purposes, concomitant with transfers of livestock which seed the new 615 substrata, makes these lagoons hotspots for non-indigenous species establishment (Mineur *et al.*, 616 2015).

⁶¹⁷The Southeast Mediterranean Sea constitutes another introduction hotspot, which accounts 618 for 24 first reports and a total of 32 non-indigenous species. The inauguration of the Suez Canal in 619 1869 resulted in an open connection between the Northern Red Sea and the Eastern Mediterranean 620 Sea. As a result, more than 500 marine species are believed to have invaded the Mediterranean Sea 621 through the Suez Canal, so-called Lessepsian migrants (Zenetos *et al.*, 2010; Zenetos *et al.*, 2012; ⁶²²Galil *et al.*, 2021). With respect to non-indigenous seaweeds, many species were first reported in a 623 series of papers by the Egyptian phycologist Anwar Aleem (Aleem, 1948; Aleem, 1950; Aleem, ⁶²⁴1951; Aleem, 1993). Recent efforts by Greek, Lebanese, Israeli and Turkish phycologists have 625 expanded the list of Lessepsian seaweeds considerably (e.g. Tsiamis, 2012; Hoffman, 2013; Bitar *et* ⁶²⁶*al.*, 2017; Israel & Einav, 2017; Çinar *et al.*, 2021; Galil *et al.*, 2021). Nevertheless, a paucity of 627 historic baseline data makes it often difficult to establish the Lessepsian origin of many species or to 628 point to the exact date of introduction. As outlined above, records of species with clear Indo-Pacific ⁶²⁹affinities which predate the opening of the Suez Canal (e.g. *Ganonema farinosum* and ⁶³⁰*Acanthophora nayadiformis*) still puzzle phycologists. In addition, the identities of many species ⁶³¹reported for the first time by Aleem and others (e.g. *Gracilaria arcuata, G. disticha, Hypnea*

flagelliformis, *Solieria dura, Spatoglossum variabile*) have never been confirmed using molecular markers and are highly uncertain. In general, a detailed understanding of past and contemporary 634 temporal dynamics of seaweed introductions in the Eastern Mediterranean Sea remains a challenge. More than in any other region it remains difficult to link the observation of a new seaweed species with the introduction date. This uncertainty has bearing on the monitoring of migration through the Suez Canal which has been regarded as an ongoing process (Boudouresque, 1999; Por, 2012). The 638 current construction of the new Suez Canal, doubling the capacity of the current corridor, is expected to further increase the influx of Red Sea species (Galil *et al.*, 2015) and contribute to further tropicalisation of the Mediterranean Sea (Bianchi & Morri, 2003; Coll *et al.*, 2010).

641 Compared to the two Mediterranean Sea introduction hotspots, first records appear less ⁶⁴²localised in the Northeast Atlantic Ocean. The English Channel (Brittany, southern English coast) 643 and the Scheldt estuary (the Netherlands) are most prominent as introduction hotspots. To what ⁶⁴⁴extent this spatial pattern reflects the true locations of primary introductions or whether the 645 locations of first records are biased by the distribution of preferred study areas of phycologists and 646 research institutes is difficult to assess. There is, indeed, a strong correlation between the 647 introduction hotspots (i.e. locations from which a high number of first records for the study area ⁶⁴⁸were reported) and the density map of all records of non-indigenous species, which is indicative of 649 high monitoring activities in areas where many non-indigenous species are found. In addition, it is 650 noteworthy that the English Channel and Scheldt estuary are important areas for oyster farming, and 651 many non-indigenous species have been accidentally introduced with oyster transfer from the 652 Mediterranean Sea.

⁶⁵⁴**Introduction rates: temporal trends**

⁶⁵⁵Disentangling the factors underpinning temporal trends in the accumulation rate of non-indigenous 656 seaweeds may improve our understanding of introductions and result in better-informed predictions 657 of future trajectories (Seebens *et al.*, 2018). Deducing temporal trends in the rate of introduction of ⁶⁵⁸non-indigenous species, however, assumes a correlation between the date of introduction and the 659 moment the species was detected. Although seemingly straightforward, for several species the 660 timespan between introduction and detection is probably considerable and unpredictable. Detection 661 obviously depends on collecting effort, but as highlighted in previous sections, the taxonomic and ⁶⁶²biogeographic framework will also determine if a species is considered non-indigenous.

663 Furthermore, Costello & Solow (2003) demonstrated that an increasing rate of detection need not ⁶⁶⁴imply an increasing rate of introductions even when collection effort is constant. Given this 665 complexity, reports that introduction rates have increased or decreased in specific time windows 666 should be treated with caution.

667 Acknowledging this uncertainty, the detection of non-indigenous species shows two distinct 668 phases, one prior to 1950-1970 characterised by low accumulation rates, followed by another much
669 higher accumulation rate from then onward (Fig. 8). Irrespective of taxonomic and biogeographic higher accumulation rate from then onward (Fig. 8). Irrespective of taxonomic and biogeographic uncertainties, there is little to no indication for a decline in the rate at which non-indigenous species are reported. The observation that the detection, and presumably also the introduction, of non-indigenous species has not reached saturation (Fig. 8), is in line with the observations by Seebens *et al*. (2017) for other taxonomic groups. For seaweeds, the absence of a decline in the first-record rate may point to the inefficiency of measures aimed to prevent and mitigate new introductions. Alternatively, the community of phycologists involved in monitoring of non-indigenous seaweeds has become larger and more efficient at detecting incoming species, e.g. through the use of DNA-677 barcoding methods.

678 According to Mineur *et al.* (2007b) who sampled the hulls of several commercial cargo 679 vessels, hull fouling seems to play a relatively minor role in the displacement of seaweeds across 680 the globe. Similarly, ballast water, one of the prime sources of introductions for marine invertebrates and microalgae (Bolch & de Salas, 2007; Gollasch *et al.*, 2015), is relatively 682 unimportant with respect to introduction of seaweeds. Yet there is evidence of introduction with 683 commercial ships not hulls or ballast water but other components such as anchors, for instance in Australia for *U. pinnatifida* (South *et al.*, 2017). Leisure boats, however, likely contribute to the local spread of non-indigenous species within the study area. Their role has certainly been underestimated so far, at least for secondary introductions (Mineur *et al.*, 2008). Contrary to the Eastern Mediterranean Sea where the opening of the Suez Canal resulted in an ongoing and steady influx of non-indigenous species (Galil *et al.*, 2015), in the Western Mediterranean Sea a disproportionate number of non-indigenous seaweed species appears to have been introduced through import of oyster stocks (Verlaque *et al.*, 2007). In the late 1960s and early 1970s, disease 691 caused by Asian oysters importation outbreaks in the study area affecting oyster populations caused a major disruption of production (Mineur *et al.*, 2015). Mitigation procedures involved massive imports in the 1970s of Pacific oyster from its native range in the Northwest Pacific Ocean, or via from the Puget Sound in the Northeast Pacific Ocean where the species is also cultivated on a large 695 scale (Mineur *et al.*, 2014). *Sargassum muticum* introduced into the study area was shown to have 22 ⁶⁹⁶several origins including a primary introduction from Asia and a secondary introduction from the

- 697 Northeast Pacific Ocean (Le Cam et al., 2020).
- 698

699 Conclusion and perspectives

700 In conclusion, our critical synthesis of non-indigenous seaweed diversity in the Northeast Atlantic 701 Ocean, Mediterranean Sea and Macaronesia revealed widespread taxonomic and biogeographic ⁷⁰²uncertainty. This finding negatively impacts efforts to evaluate the effectiveness of measures to 703 reduce non-indigenous species influx, manage their risks and impacts, and devising potential 704 control strategies. This uncertainty can be addressed through the progressive use of molecular ⁷⁰⁵markers, particularly standard DNA barcoding approaches, which can in most cases confirm the 706 identification of presumed non-indigenous seaweeds (Viard $&$ Comtet, 2015). Importantly, 707 however, a taxonomic and biogeographic reference framework should also be established for 708 putative indigenous regions. DNA-based identification is relatively well-developed for species with ⁷⁰⁹a Northwest Pacific origin. For tropical taxa such a framework lags behind, resulting in higher 710 levels of uncertainty regarding the identity of non-indigenous species with presumed tropical origin. ⁷¹¹Such a reference framework will also be necessary for early detection of non-indigenous species 712 and monitoring with the use of bulk sample and eDNA metabarcoding (Darling *et al.*, 2017; Keck ⁷¹³& Altermatt, 2023).

⁷¹⁴However, standard DNA barcoding may not be sufficient to interpret more complex 715 introduction histories, such as cases where non-indigenous and indigenous populations co-occur, as 716 suggested for several species in the Mediterranean Sea (Verlaque *et al.*, 2015) or where recent 717 human-mediated dispersal needs to be evaluated against ongoing and natural dispersal events linked 718 to Pleistocene or Holocene climatic oscillations (Neiva *et al.*, 2016). In these cases a combination of 719 population-level sampling strategies and molecular markers that capture intraspecific diversity is ⁷²⁰needed to shed light on the number and directions of dispersal events and re-evaluate the status of 721 taxa currently considered cryptogenic.

⁷²²In parallel with DNA-barcoding efforts, historical and reliable baseline data of seaweed 723 diversity is needed to reduce the subjective interpretation of non-indigenous species and determine ⁷²⁴more precisely their date of introduction. It is likely that the introduction of many non-indigenous 725 species occurred significantly earlier than the time of their first detection, especially for species that ⁷²⁶have morphologically similar congeners in the study area (such as species from the genera *Dictyota*, *Gracilaria*, *Polysiphonia* and *Ulva*). Herbarium collections can serve as a crucial source of primary 728 data to address this issue, and advances in sequencing technologies make it possible to obtain 729 genetic data from voucher specimens tens or even hundreds of years old as demonstrated for the *Codium fragile*-complex (Provan *et al.*, 2008). We anticipate that herbaria will play an increasingly important role in documenting spatio-temporal patterns of non-indigenous seaweeds, alongside large-scale digitization efforts for these collections. More precise estimates of the date of 733 introduction will also reduce uncertainty regarding the accumulation rate of non-indigenous species. Our analyses suggest that the rate of introduction in the study area has not decreased. However, it is unclear whether this trend reflects a steady accumulation of non-indigenous species or increased 736 and more efficient detection through monitoring or advanced identification methods. Based on reliable baselines, temporal surveys will allow to uncover the trends in non-indigenous species 738 introduction rates. Diversifying the type of survey, from punctual surveys (e.g. Bio-Blitzes or Rapid Assessment surveys) to systematic comprehensive surveys (e.g. full inventories), including morphological-based and DNA-based assessments, such as metabarcoding (see above), is a need for effective prevention and early detection and also to monitor trends over time in new species 742 introductions.

743 Our study also reveals significant differences in the geographical distribution of non-744 indigenous seaweed species across the study area, with only 18 species shared between the three 745 main regions. Non-indigenous species distribution reflects both their abiotic niche and the
746 efficiency of primary and secondary dispersal vectors. It is expected that current patterns v efficiency of primary and secondary dispersal vectors. It is expected that current patterns will ⁷⁴⁷become increasingly homogenous with time due to various factors. First, recently introduced ⁷⁴⁸species may continue to expand as part of the ongoing invasion process. Secondly, evolutionary 749 processes such as selection, genetic admixture and hybridisation may occur during the invasion ⁷⁵⁰process, leading to adaptation and expansion of the species in new ecological conditions. Last, 751 further range expansions are expected for non-indigenous species with affinities for warm temperate 752 to tropical temperatures as a result of ongoing ocean warming. This could result in an influx of 753 warm-water adapted species into regions where conditions are currently unfavourable, while ⁷⁵⁴warming may also render regions unfavourable for non-indigenous species that currently thrive 755 there. It is anticipated that under ocean warming, Eastern Mediterranean Sea non-indigenous 756 species would likely expand their range in western direction, while species ranges in Lusitania ⁷⁵⁷would shift northward. The accurate estimation of the rate at which non-indigenous species expand 758 their ranges, either as invasion fronts or more erratically by jump dispersals, rely on detailed

759 monitoring. Such estimates can inform on dispersal vectors at various spatial scales and guide 760 policy makers to take effective measures to prevent or limit the spread of these species.

⁷⁶¹We advocate that combining the efforts of taxonomists who provide a reliable framework on 762 the number and likely geographic origin of non-indigenous seaweeds, together with environmental ⁷⁶³monitoring, offers the best strategy to identify species of concern, characterise their life history 764 traits, and develop effective management strategies. Especially for the species entering the
765 Mediterranean Sea through the Suez Canal, a combination of horizon scanning exercises, in Mediterranean Sea through the Suez Canal, a combination of horizon scanning exercises, intensive 766 monitoring and rapid-response eradication efforts at the local level may be the only tools to try and 767 control the establishment of Lessepsian migrants. With respect to non-indigenous species that ⁷⁶⁸hitchhike with shellfish transport, effectively limiting imports into Europe and controlling 769 translocations between regions in Europe should effectively reduce the rate of primary and 770 secondary introductions (Mineur *et al.*, 2014). Alternatively, immersion for shorter periods (3 771 seconds) at temperatures of $80-85^{\circ}\text{C}$ is effective in killing macroalgal propagules (Mineur *et al.*, ⁷⁷²2007a). In addition, hull fouling has an important role for introductions, especially towards ⁷⁷³secondary spread of already introduced non-indigenous seaweeds (Clarke Murray *et al.*, 2011). The 774 related guidance developed in the context of the Marine Environmental Protection Committee ⁷⁷⁵(MEPC, 2011) is a step forward, but we stress the need for more enforceable control of this 776 pathway. Mitigating the negative effects of non-indigenous species that have already established ⁷⁷⁷will likely prove to be even more difficult. Solid baseline data will allow us to detect introduction 778 patterns and non-indigenous species range shifts in early stages and act accordingly.

⁷⁷⁹Combatting the effects of non-indigenous seaweeds will require coordinated action at the ⁷⁸⁰European and international level to prevent the introduction of species, to quickly detect and rapidly 781 eradicate species to prevent them from establishing, and to manage established species to minimise 782 their ecological and economic impact (IAS Regulation (EU) 1143/2014 on invasive alien species). ⁷⁸³These measures will require significant efforts and collaboration between science, management, 784 policy, and society. Our dataset supports these regulatory actions by providing a solid baseline on 785 non-indigenous seaweeds. This baseline contributes to the assessment of the current situation, helps 786 authorities to identify new introductions and monitor the status of already established species, and ⁷⁸⁷importantly identifies the current knowledge gaps concerning taxonomic and biogeographic 788 uncertainties.

790

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803 Disclosure statement

804 No potential conflict of interest was reported by the authors.

806 Supplementary information

807 Supplementary Table S1: concise description of the status of species flagged as non-indigenous in 808 the Northeast Atlantic Ocean, the Mediterranean Sea, and Macaronesia.

809 The complete dataset containing all records is available at Zenodo (DOI:10.5281/zenodo.7798640).

811 Author contributions

- 812 LM. van der Loos + Q. Bafort +S. Bosch: concept, data acquisition, analyses, writing; F. Leliaert +
- 813 O. De Clerck: concept, data acquisition, analyses, writing. Other authors: data acquisition, writing.

814

- ⁸¹⁵**Table 1.** Overview of the non-indigenous seaweed species (Charophyta, Chlorophyta,
- 816 Phaeophyceae, Xanthophyceae, Rhodophyta) reported from the Northeast Atlantic Ocean, the 817 Mediterranean Sea, and Macaronesia.
- Mediterranean Sea, and Macaronesia.

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819

⁸¹⁹**FIGURE 1.** Illustration of selected non-indigenous species in the study area. A) *Colpomenia*

820 *peregrina* growing attached to nylon fishing net in a harbour (photo: Frank Perk, the Netherlands);
821 B) The green non-indigenous species *Ulva australis* (photo: Ignacio Bárbara, Atlantic coast Spain);

- ⁸²¹B) The green non-indigenous species *Ulva australis* (photo: Ignacio Bárbara, Atlantic coast Spain);
- ⁸²²C) A dense *Codium fragile* subsp. *fragile* reef (photo: Mick Otten, the Netherlands); D)
- ⁸²³*Caulacanthus okamurae* often grows high in the intertidal (photo: Ignacio Bárbara, Atlantic coast
- ⁸²⁴Spain); E) *Rugulopteryx okamurae* has been introduced in the Northeast Atlantic Ocean,
- 825 Mediterranean Sea and in Macaronesia (photo: Sandrine Ruitton, Mediterranean coast France); F)
- ⁸²⁶*Asparagopsis armata* is often regarded as a high-nuisance invasive species (photo: Ignacio Bárbara,

- 827 Atlantic coast Spain); G) *Gelidium vagum* has only been reported from the Netherlands but is
- 828 locally very abundant (photo: Mart Karremans, the Netherlands); H) A *Sargassum muticum* forest (photo: Rob Aarsen, the Netherlands); I) *Grateloupia turuturu, Codium fragile* subsp. *fragile*, and
- ⁸²⁹(photo: Rob Aarsen, the Netherlands); I) *Grateloupia turuturu*, *Codium fragile* subsp. *fragile*, and
- ⁸³⁰*Sargassum muticum* covering the seabed (photo: Ad Aleman, the Netherlands); J) A fertile
- 831 specimen of *Nitophyllum stellatocorticatum* (photo: Mart Karremans, the Netherlands); K) *Undaria*
- ⁸³²*pinnatifida* growing attached to aquaculture facilities (photo: Ron Offermans, the Netherlands); L)
- ⁸³³A tidal pool with *Grateloupia turuturu* (photo: Ignacio Bárbara, Atlantic coast of Spain).

834
835 **FIGURE 2.** Flowchart for assessing the taxonomic confidence and biogeographic status of putative

836 non-indigenous species. This figure builds on the concepts from Essl *et al.* (2018).

- 837
838 ⁸³⁸**FIGURE 3.** Map of the study area (Northern Europe, Lusitania, Macaronesia, Western
- 839 Mediterranean Sea, and Eastern Mediterranean Sea), with indication of the number of recorded non--
- 840 indigenous species, number of records and the length of the coastline. The number of species
- 841 includes cryptogenic and data deficient species.

842
843 ⁸⁴³**FIGURE 4.** Biogeographic status (either non-indigenous or uncertain, the latter category including 844 the 'cryptogenic' and 'data deficient' species) and taxonomic confidence of seaweed species 845 flagged as non-indigenous in the study area. Circle surface area corresponds to the number of 846 species.

species.

847
848 849 (Rhodophyta) non-indigenous species in Northern Europe, Lusitania, Macaronesia, the Eastern and
850 Western Mediterranean Sea, and aquaculture sites in the Mediterranean Sea. Dark shaded colours Western Mediterranean Sea, and aquaculture sites in the Mediterranean Sea. Dark shaded colours 851 represent numbers of high-confidence non-indigenous species; light shaded colours represent

852 cryptogenic and data deficient species.

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853 854 FIGURE 6. Number of non-indigenous species shared among regions. A) Mediterranean Sea,
855 Macaronesia, and Northeast Atlantic Ocean; B) Northern Europe, Lusitania, and Macaronesia; Macaronesia, and Northeast Atlantic Ocean; B) Northern Europe, Lusitania, and Macaronesia; C) 856 Western Mediterranean Sea, Eastern Mediterranean Sea, and aquaculture sites in the Mediterranean 857 Sea. Numbers display the high-confidence non-indigenous species (excluding cryptogenic and data

858 deficient species), with total number of non-indigenous species including cryptogenic and data

859 deficient species in brackets.

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 B_6 ⁸⁶⁰**FIGURE 7.** 861 The origin of non-indigenous species. For each of the recipient study regions, the Sankey diagram

862 displays what proportion of non-indigenous species have likely originated from a specific area

863 (Northwest Pacific Ocean, Northeast Pacific Ocean, Indo Pacific Ocean/Red Sea, Australasia,

864 Mediterranean Sea, Northeast Atlantic Ocean, Western Atlantic Ocean, or uncertain). The width of

865 the flow arrows is proportional to the number of non-indigenous species (including non-indigenous,

866 cryptogenic and data deficient species).
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species to be evaluated

 \bigcirc Mediterranean Sea

- Macaronesia \bigcap
- Northeast Atlantic Ocean C

- Northern Europe \bigcirc
- О Lusitania
- Macaronesia \bigcirc

 $C)$

West. Med. Sea East. Med. Sea \bigcirc Aquaculture Med. Sea

Origin

Recipient region

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