

Vegetación marina alrededor del mundo: Patrones, importancia y evolución

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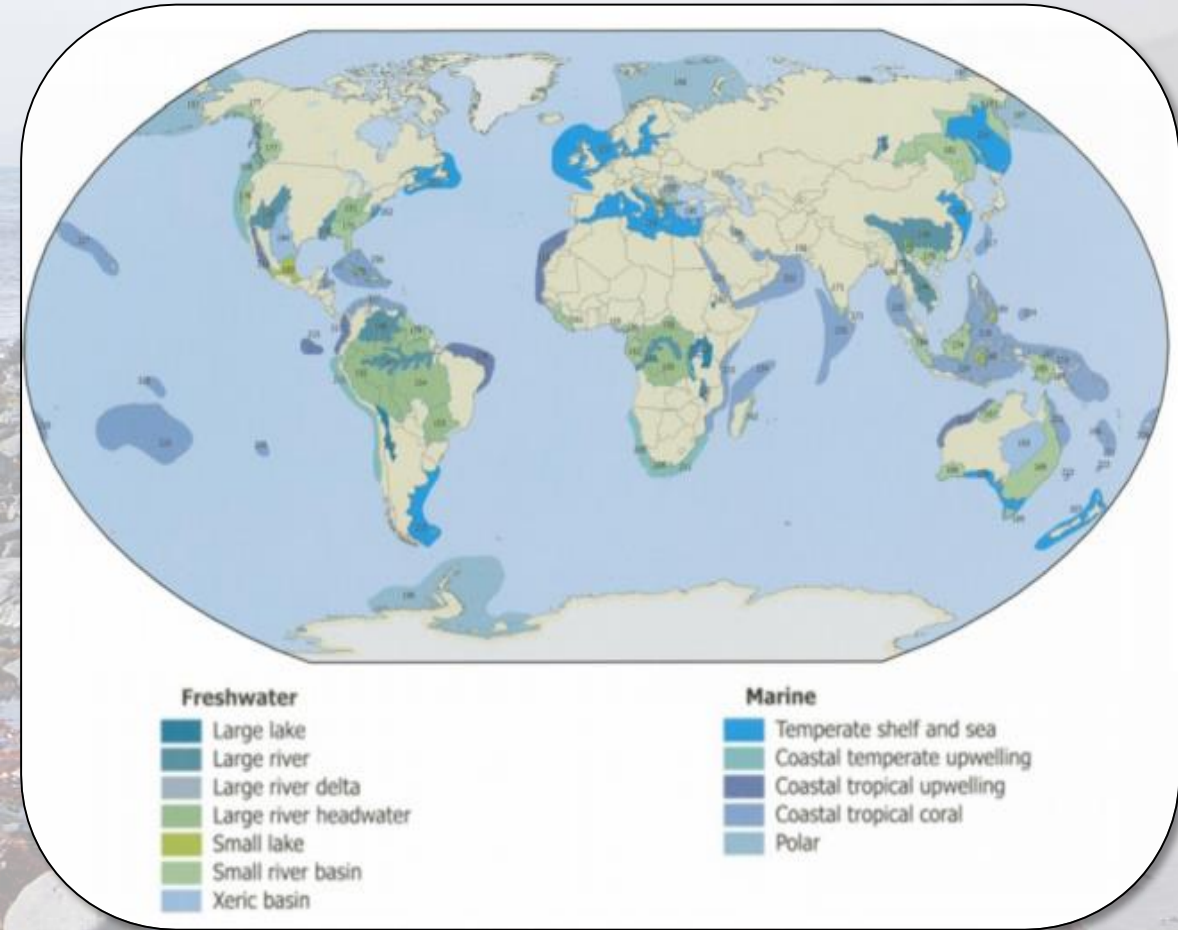
Laboratory of Botany - Department of Biology, Healthcare and the Environment - Faculty of Pharmacy
and Food Science - University of Barcelona



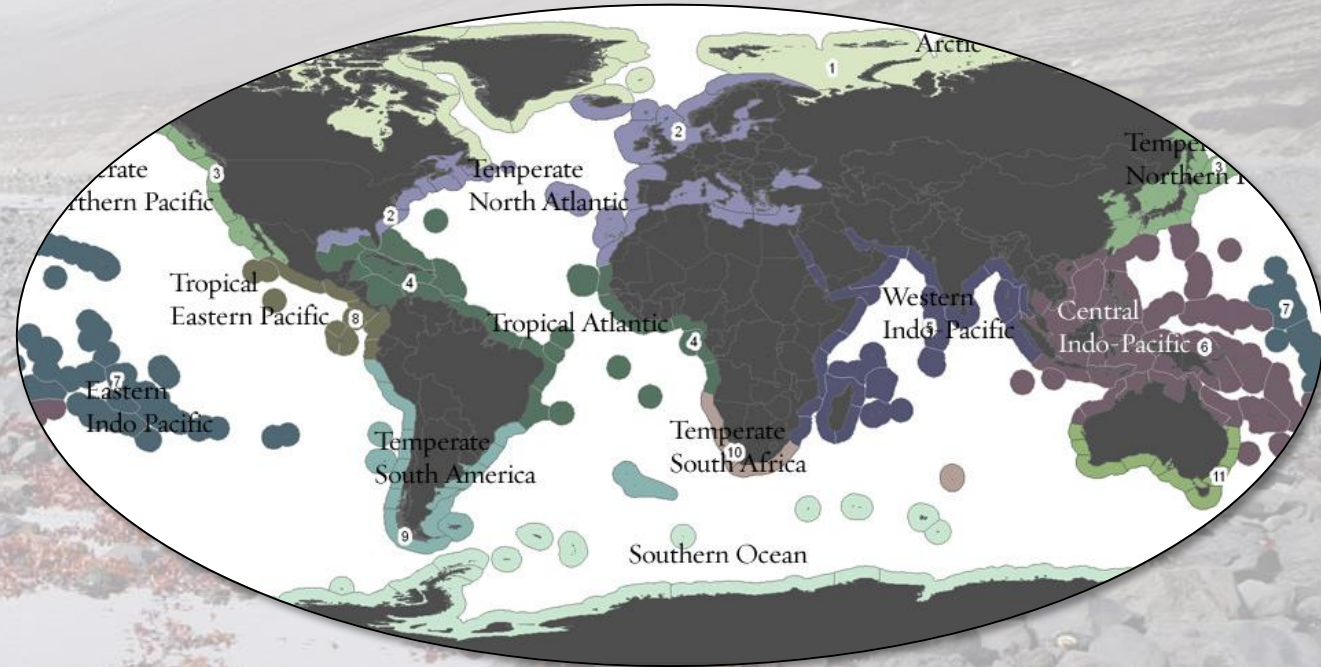
Macroalgae / Seaweeds







Freshwater and marine biomes as classified by Olson and Dinerstein (2002). In: *The Global 200: Priority ecoregions for global conservation*. *Annals of the Missouri Botanical Garden*, 89(2): 199-224



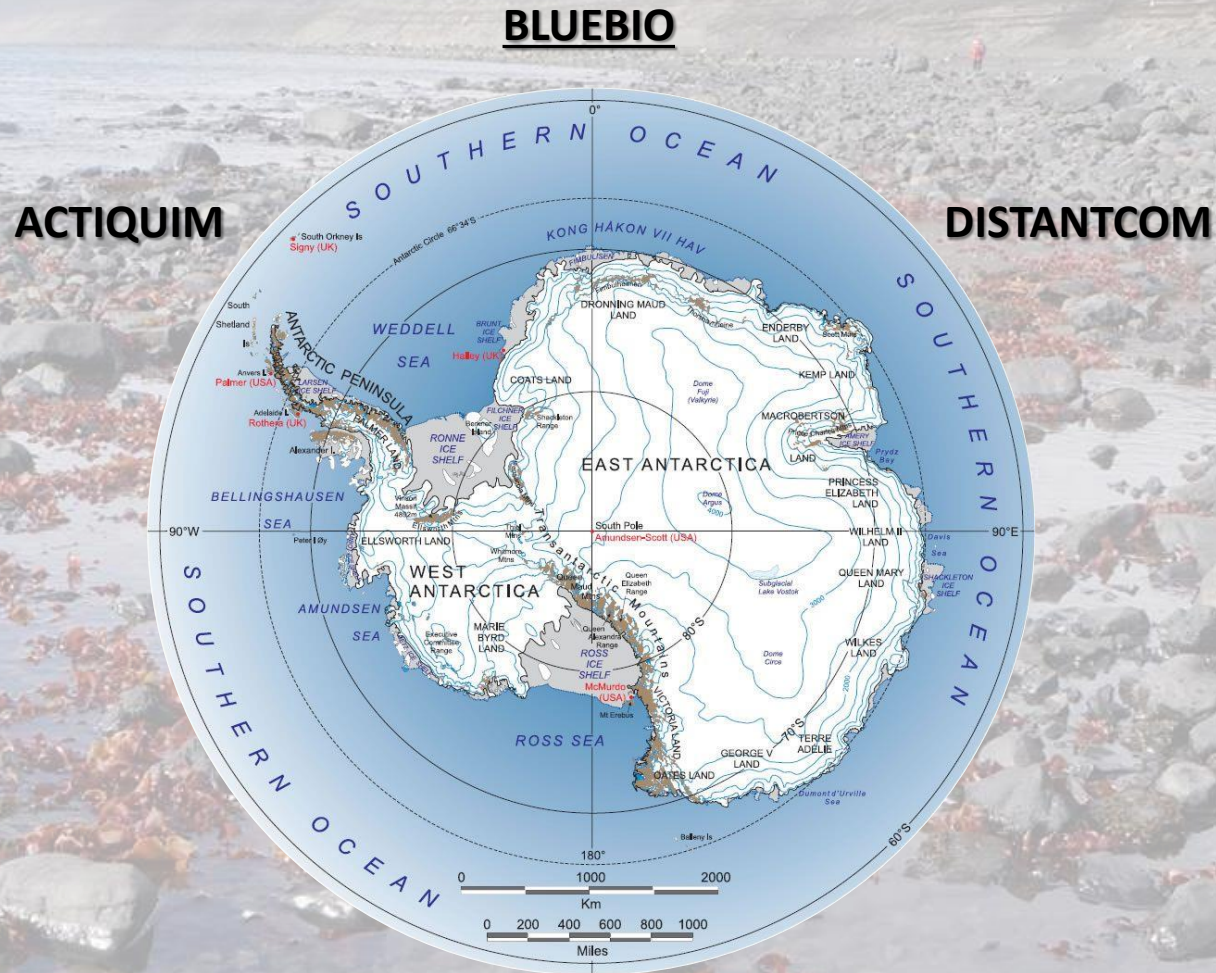
Marine realms as classified by Spalding et al. (2007). In: *Marine Ecoregions of the World: A Bioregionalization of Coastal and Shelf Areas*. *BioScience*, 57(7): 573-583.

Antarctic Marine Flora

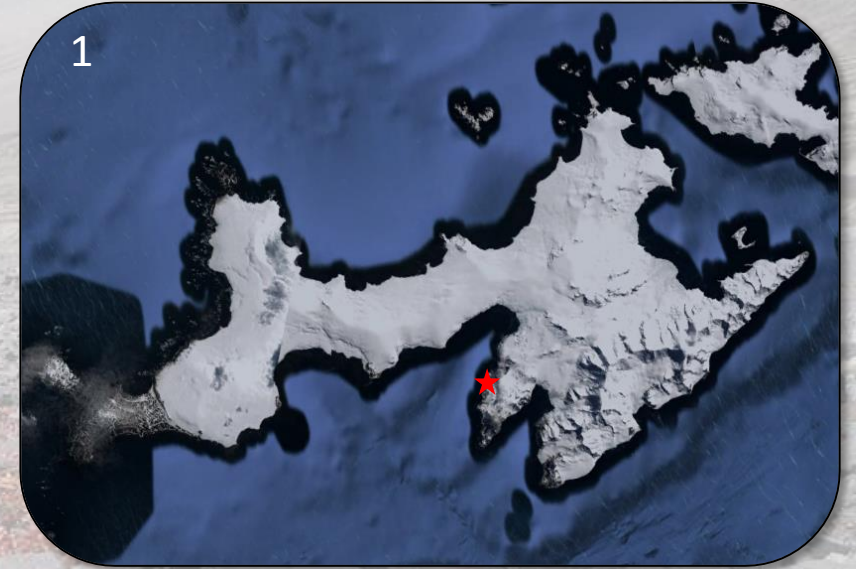
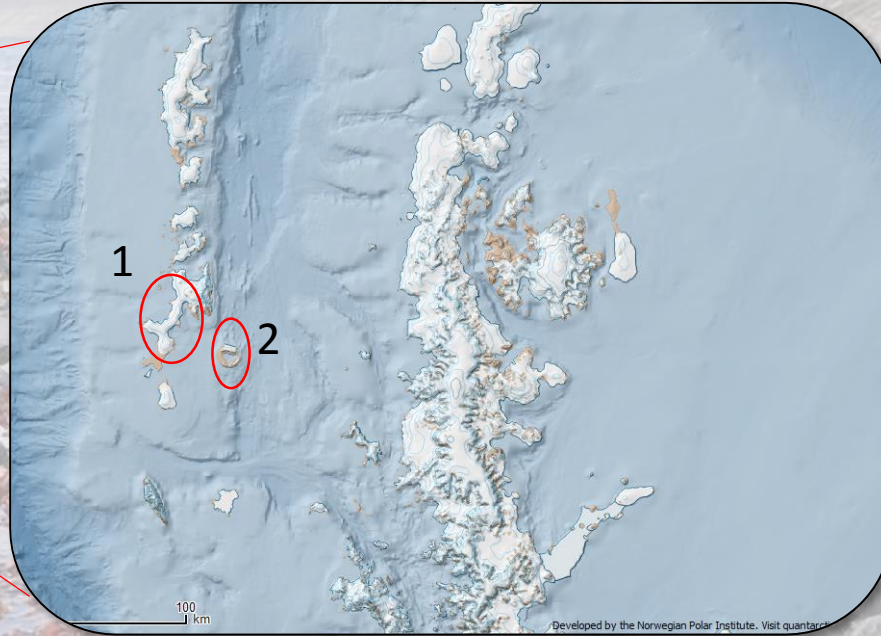
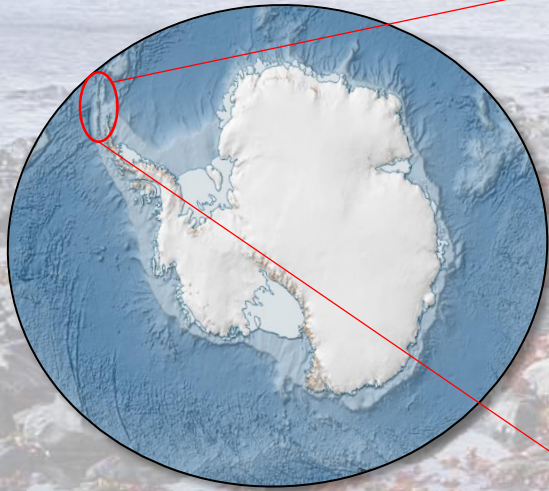
Small forms of *Fucus* from the European coasts



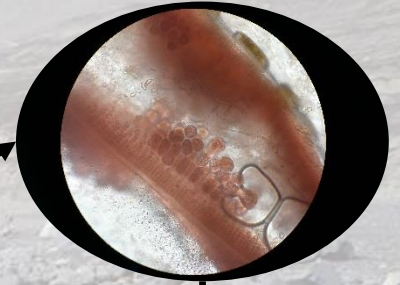
Antarctic Marine Flora



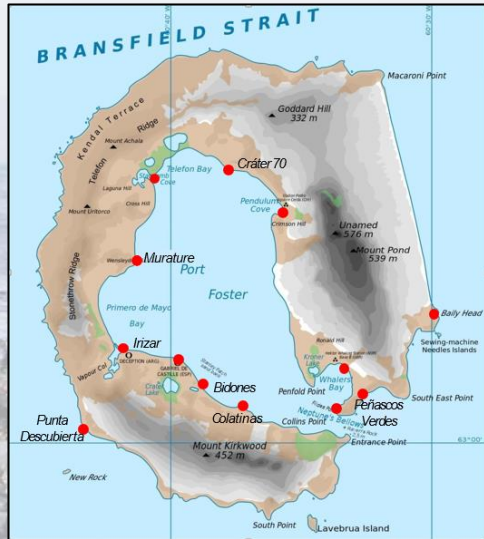
Sampling area



Sampling



Updated checklist of Deception Island



76 species

- 39 *Rhodophyta*
- 19 *Phaeophyceae*
- 18 *Chlorophyta*

63'3% of the known Antarctic flora

New cites for the Deception marine flora

1. *Austropugetia crassa* R. L. Moe
2. *Delisea pulchra* (Greville) Montagne
3. *Leniea lubrica* R. L. Moe
4. *Pantoneura plocamioides* Kylin
5. *Porphyra plocamiestris* R. W. Ricker
6. *Chordaria linearis* (J. D. Hooker & Harvey) A. D. Cotton

	NR	LA	PD	PV	CB	P	BA	CO	BI	BA	IR	MU	TEL	C7	PE	DC
	V	E	E	A	FILL			L	D	E	R			0	N	
CHLOROPHYTA																
<i>Acrosiphonia arcta</i> (Dillwyn) Gain							12			+		15				1
<i>Blidingia minima</i> (Nägeli & Kützing) Kylin									+			15				
<i>Chaetomorpha dubyana</i> Kützing							10									
<i>Chaetomorpha mawsonii</i> A.H.S. Lucas												15				
<i>Cladophora coelothrix</i> Kützing												15				
<i>Endophyton atroviride</i> O'Kelly												15				
<i>Monostruma grevillei</i> (Thuret) Wittrock												15				
<i>Monostruma harioti</i> Gain			5					10+		+		15	+	10	1,4	
<i>Prasiola crispa</i> (Lightfoot) Kützing							10					15				
<i>Rhizoclonium ambiguum</i> (J.D. Hooker & Harvey) Kützing												15				
<i>Rhizoclonium riparium</i> (Roth) Harvey												15				
<i>Spongomorpha pacifica</i> (Montagne) Kützing												15				
<i>Ulothrix australis</i> Gain												15				1
<i>Ulothrix flacca</i> (Dillwyn) Thuret				+					+			15	+			1

BCN-Phyc Herbarium

(Plant Biodiversity Resource Centre of the University of Barcelona).

Angulo-Preckler, *et al.*, (2018). Macrobenthic patterns at the shallow marine waters in the caldera of the active volcano of Deception Island, Antarctica. *Continental Shelf Research*, 157, 20-31.

Antibiotic properties of macroalgae from Antarctica

Methodology



Antibiotic properties of macroalgae from Antarctica

Methodology



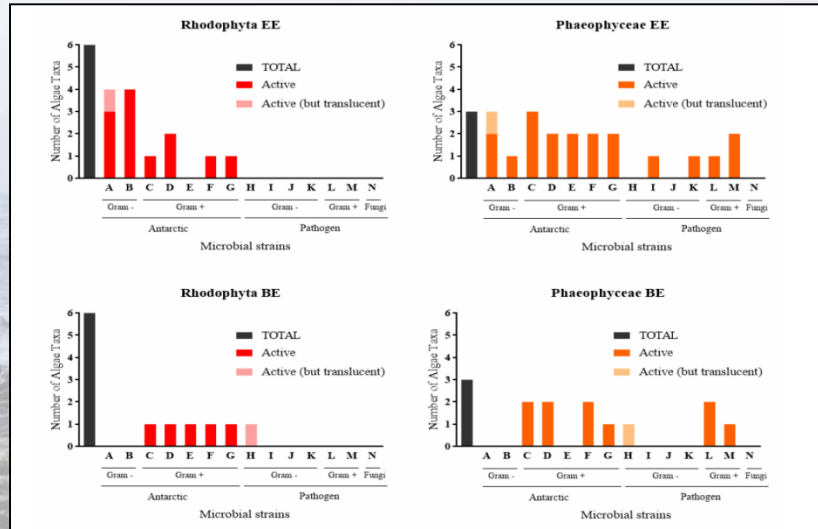
Acar (1980)

Microorganisms used in the antibiotic assay	
Antarctic	Microorganism Type
<i>Psychrobacter</i> sp.	Gram -
<i>Paracoccus</i> sp.	Gram -
<i>Arthrobacter</i> sp. (A)	Gram +
<i>Arthrobacter</i> sp. (B)	Gram +
<i>Oceanobacillus</i> sp.	Gram +
<i>Bacillus aquamaris</i>	Gram +
<i>Micrococcus</i> sp.	Gram +
Patogens	
<i>Vibrio cholerae</i> CECT-657	Gram -
<i>Escherichia coli</i> O157H7, ATCC 43888	Gram -
<i>Pseudomonas aeruginosa</i> NCTC 10332T	Gram -
<i>Escherichia coli</i> CECT515	Gram -
<i>Bacillus cereus</i> CECT 4014	Gram +
<i>Staphylococcus aureus</i> CECT 59	Gram +
<i>Candida albicans</i> CECT 1001	Fungus <i>Saccharomycetes</i>

Activity depending on Inhibition area radius	
No effect	0 (-)
Weak inhibition	0-1mm (+)
Moderate Inhibition	>1-3mm (++)
Trong Inhibition	>3-7mm (+++)
Very Strong Inhibition	>7-15mm (++++)

Lippert *et al.* (2003)

Antibiotic properties of macroalgae from Antarctica



22 Antarctic macroalgae tested

- 14 Rhodophyta (8 showed activity)
- 8 Phaeophyceae (4 showed activity)

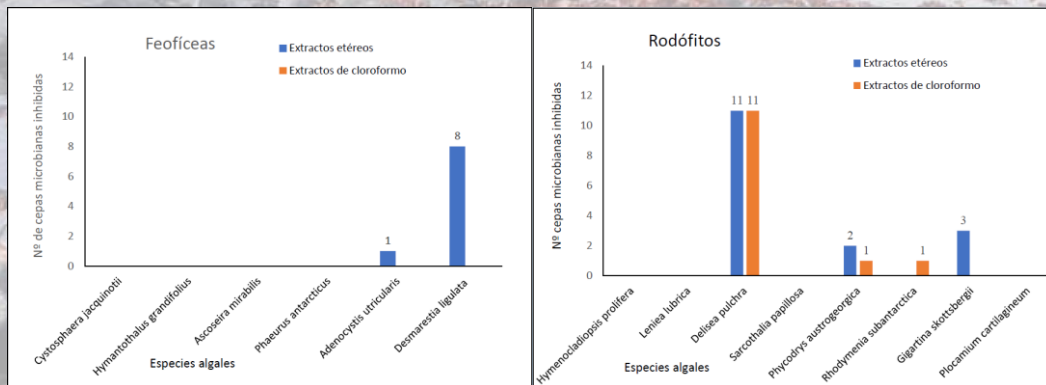
Delisea pulchra (Greville) Montagne

Desmarestia antarctica R.L.Moe & P.C.Silva

Tested microorganisms

- Antarctic bacteria were inhibited for most algae
- **Gram- were more resistant**
- *Escherichia coli*, *Pseudomonas aeruginosa* & *Candida albicans* were the most resistant

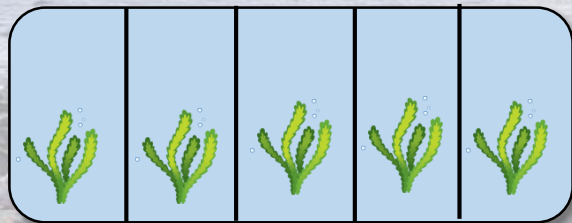
Source: Camacho (2017)



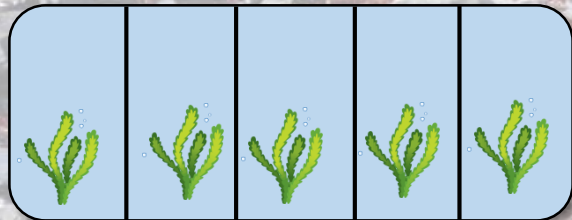
Source: Carcedo (2018)

Chemical activity in different climate change scenarios

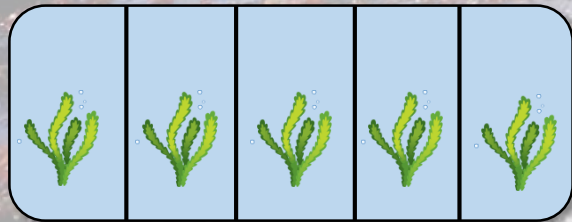
- Does Temperature changes affect to the chemical defences expression on macroalgae?
- Compare 3 different temperature scenarios in 3 different biomes (Antarctica, Mediterranean, Tropic)
- *Cystosphaera jacquinotii*, *Cystoseira compressa*, *Sargassum sp.*



T1 ~



T2 +



T3 ++



Genus Plocamium J.V.Lamouroux in Antarctica

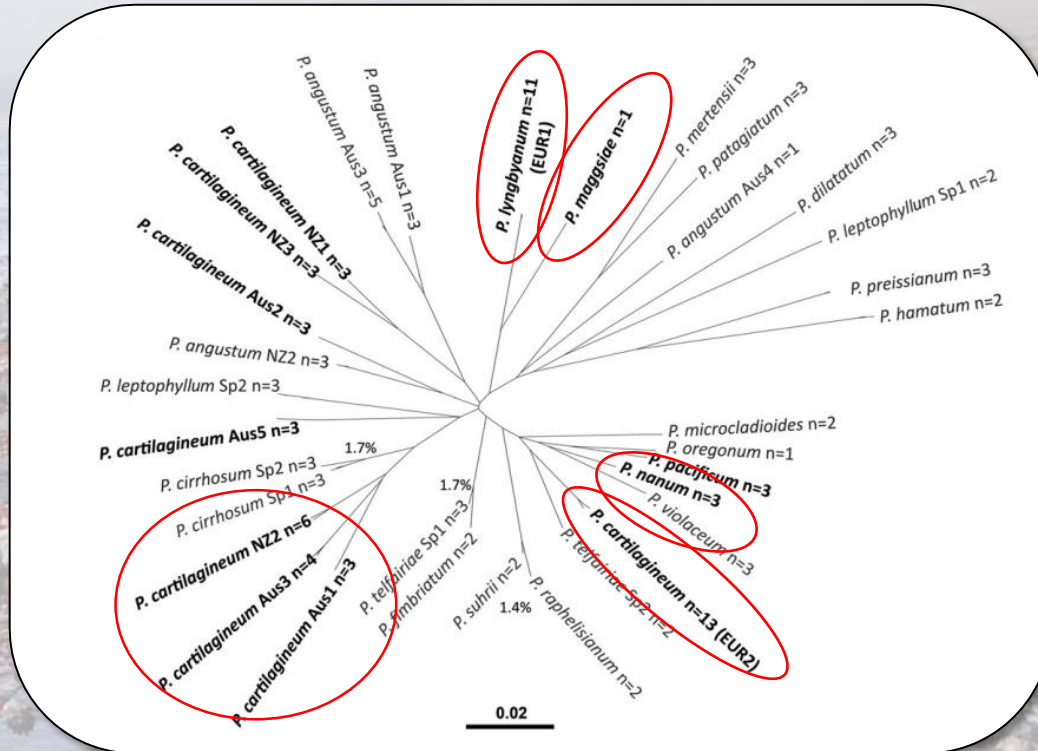
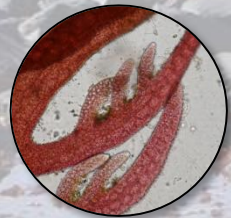
P. cartilagineum (Linnaeus) P.S.Dixon

P. hookery Harvey

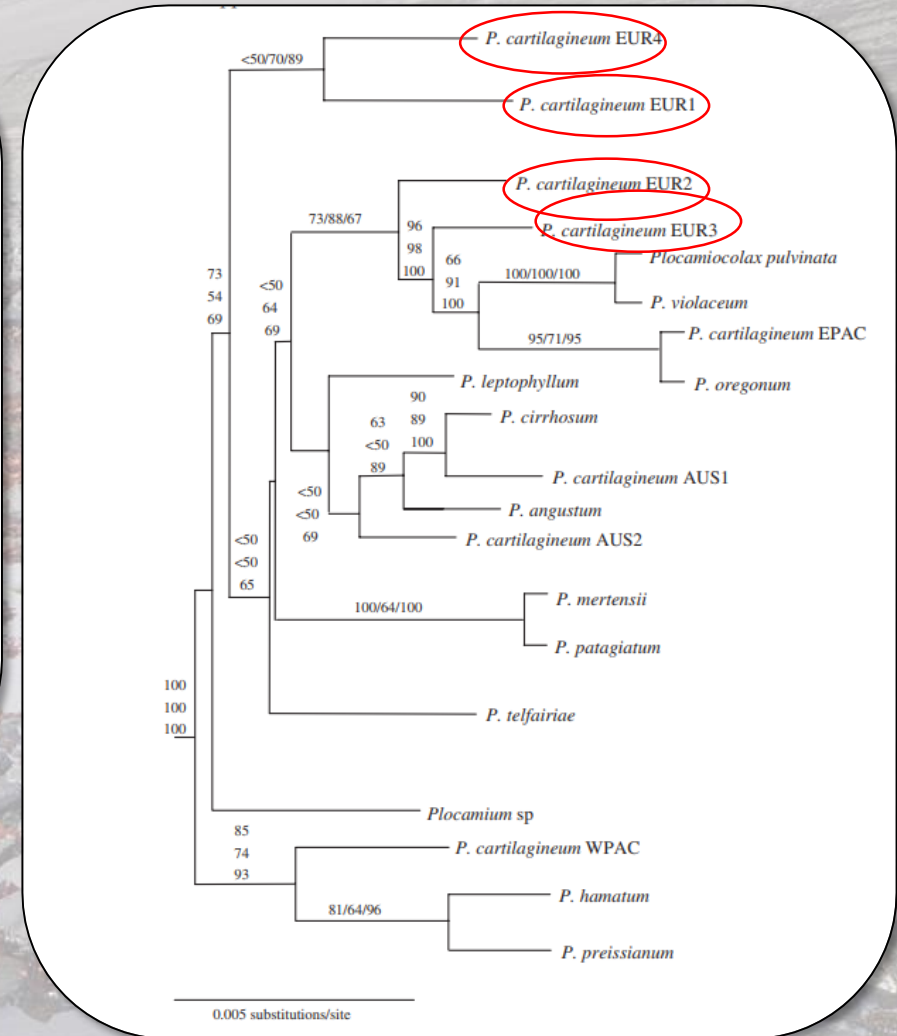
P. secundatum (Kützing) Kützing



Genus *Plocamium* J.V.Lamouroux in Antarctica



Source: Cremades *et al.* (2011).



Source: Saunders *et al.* (2005).

Small forms of *Fucus* from the European coasts



Fucus vesiculosus Linnaeus

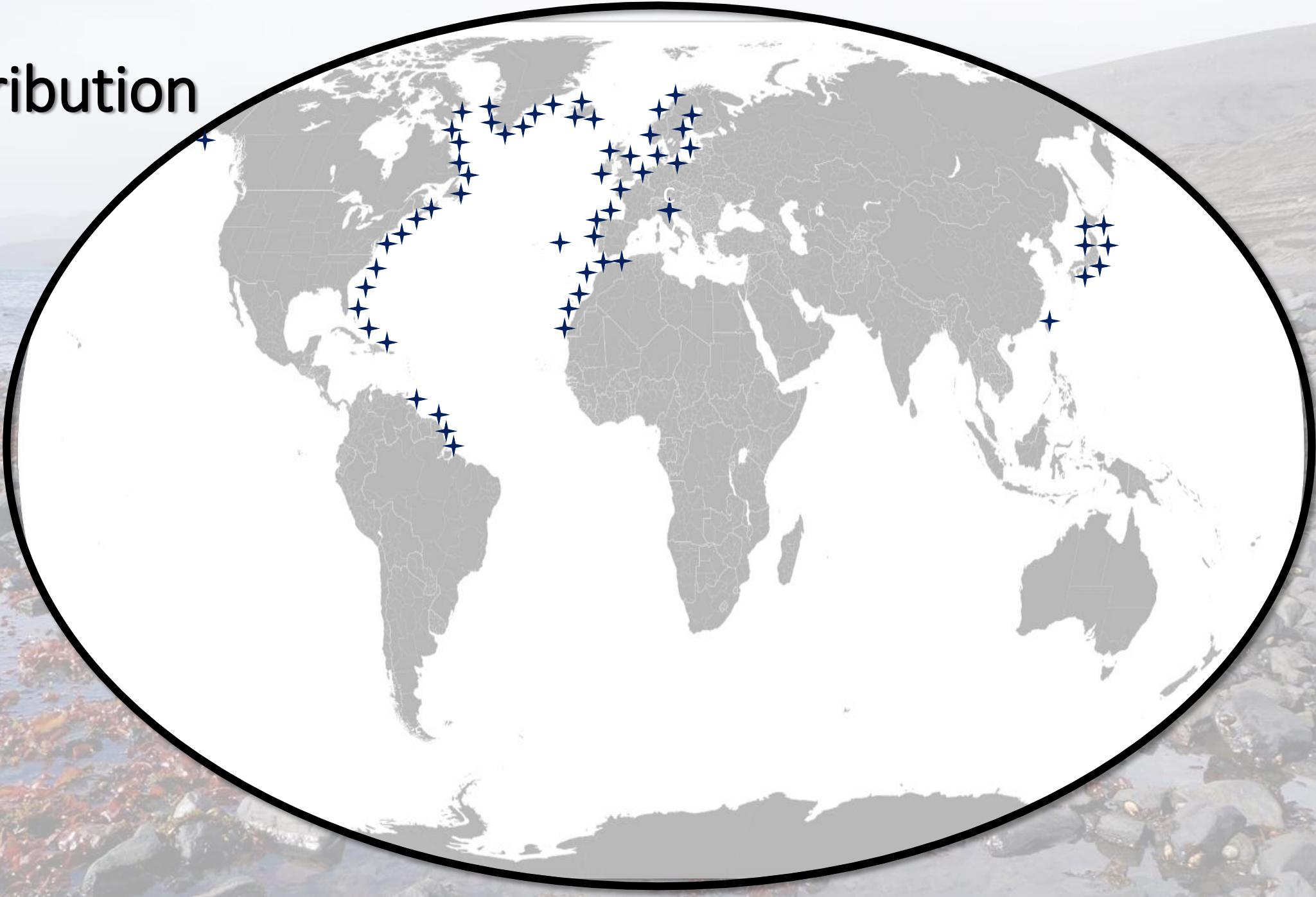
Source: aphotomarine



Fucus serratus Linnaeus

Source: wikipedia

Distribution



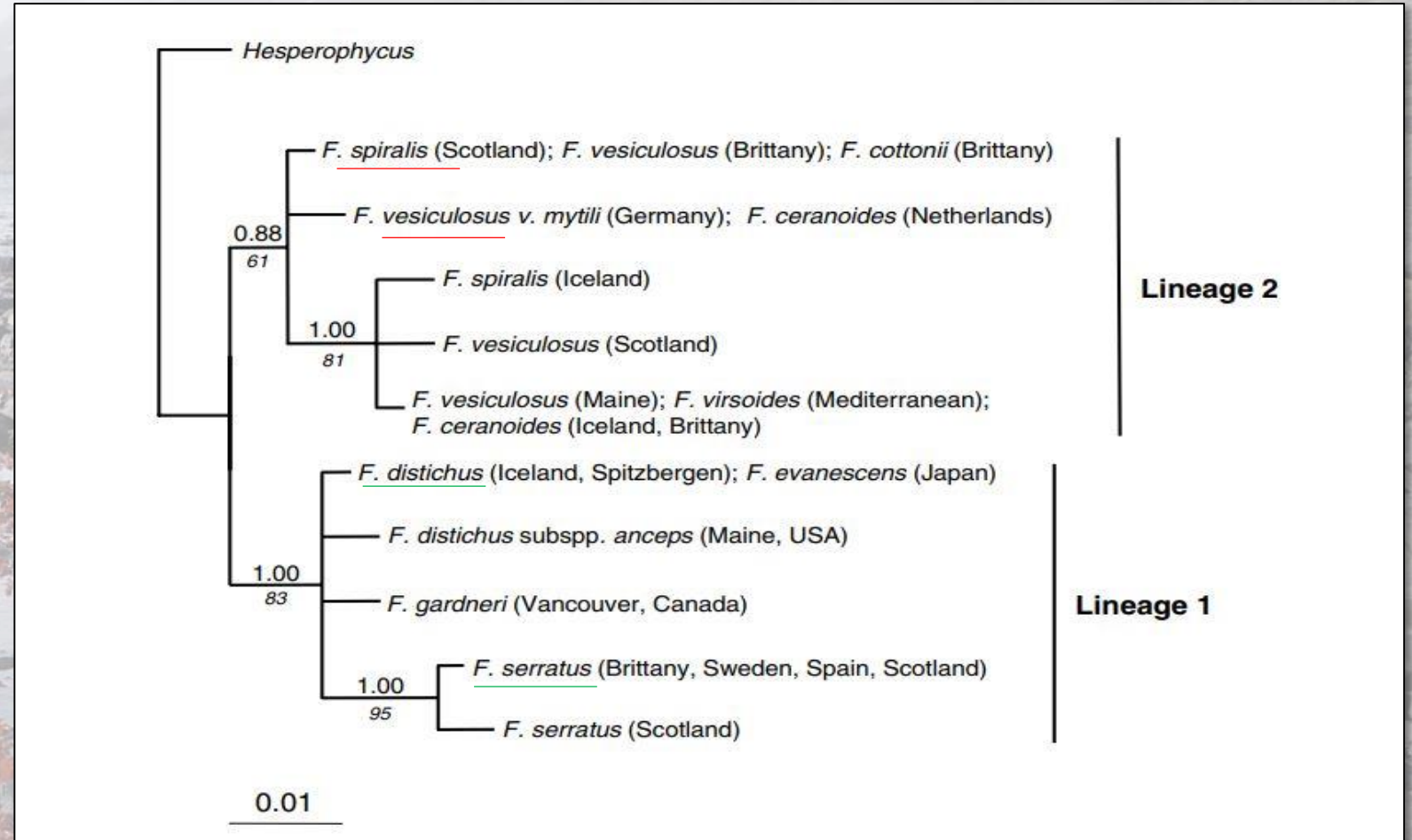
The genus *Fucus*

Fucus Linnaeus, 1753

About **9 accepted species**

- *F.chalonii*
- *F.cottonii*
- *F.vesiculosus*
- *F.spiralis*
- *F.serratus*
- *F.radicans*
- *F.distichus*
- *F.virsoides*
- *F.ceranoïdes*

Source: *Algaebase* (11-07-2018)



Bayesian phylogenetic tree based on **mtDNA 23S** sequences. Numbers above and below the line are Bayesian posterior probability and MP bootstrap values (1000 replications), respectively. Source: Coyer *et al.* (2006)



Fucus chalonii Feldmann



F. spiralis var. *nanus* (Stackhouse) Batters



Fucus cottonii M.J.Wynne & Magne



F. spiralis var. *limitaneus* (Montagne) I.M.Pérez-Ruzafa

Ecology



Fucus chalonii Feldmann



F. spiralis var. *nanus* (Stackhouse) Batters



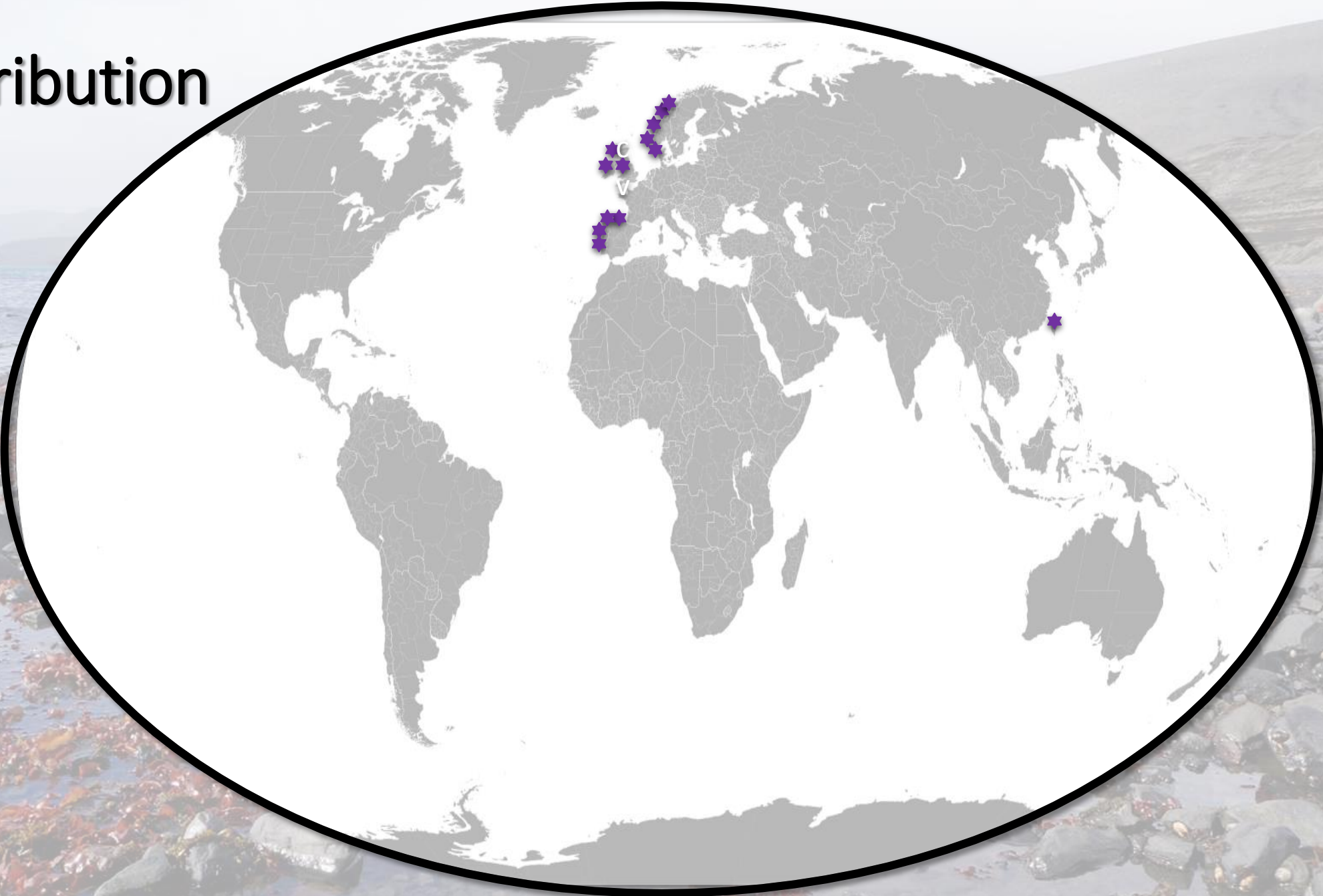
F. spiralis var. *limitaneus* (Montagne) I.M.Pérez-Ruzafa

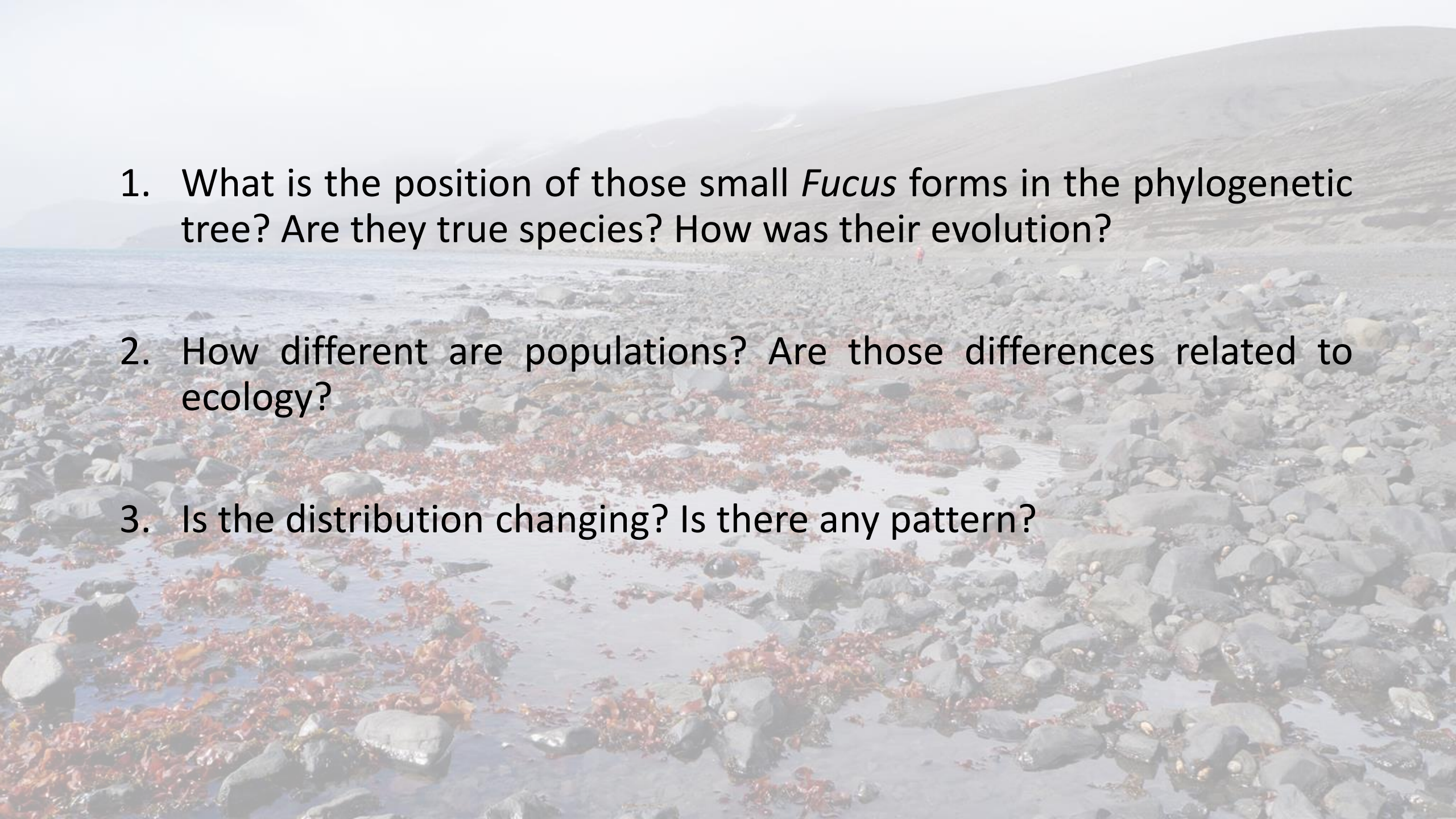
Ecology



Fucus cottonii M.J.Wynne & Magne

Distribution

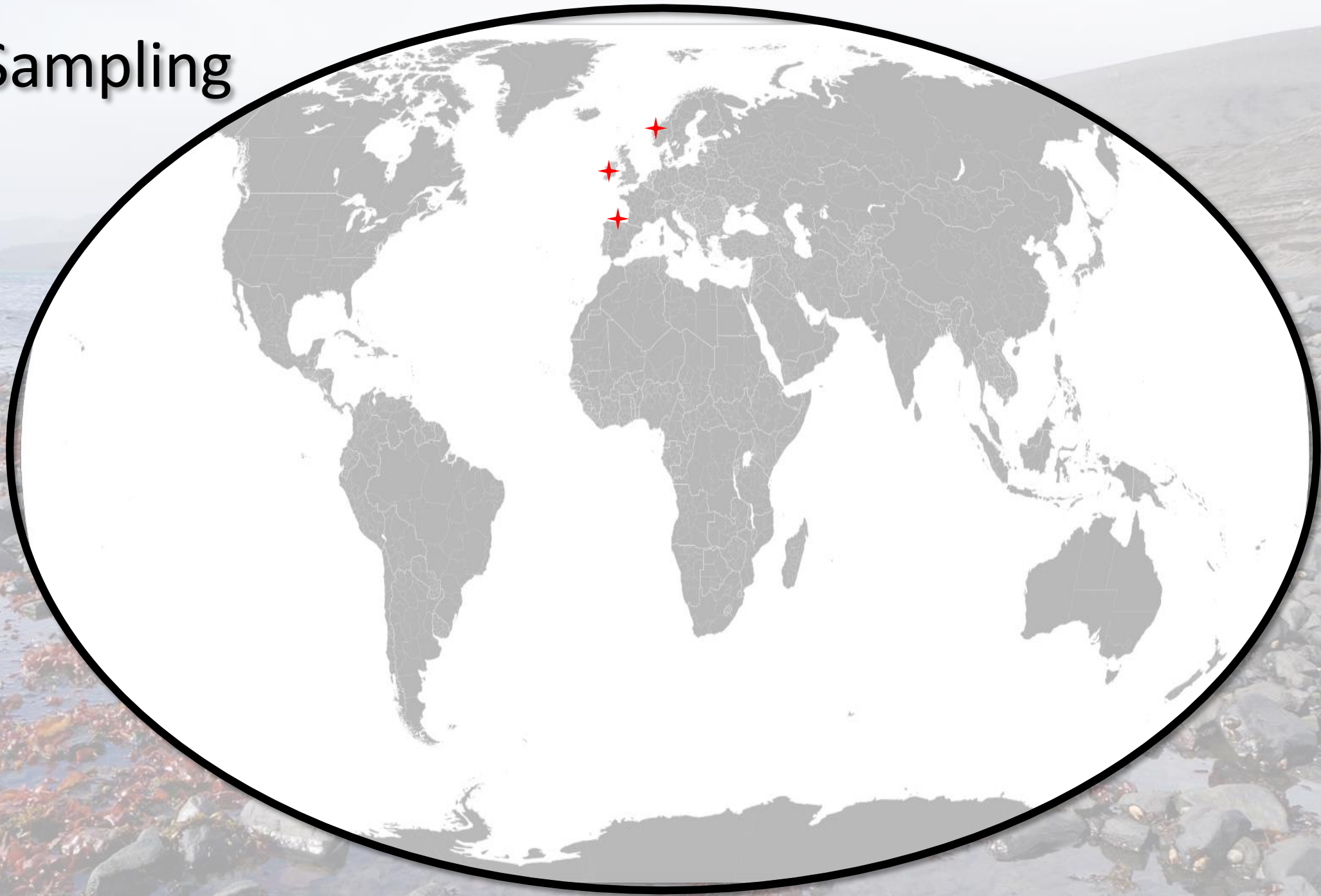


- 
- A photograph of a rocky beach with a large amount of brown seaweed scattered across the rocks. In the background, a person is visible walking on the beach, and a large, hazy hill or mountain rises in the distance. The sky is overcast and grey.
1. What is the position of those small *Fucus* forms in the phylogenetic tree? Are they true species? How was their evolution?
 2. How different are populations? Are those differences related to ecology?
 3. Is the distribution changing? Is there any pattern?

Methodologies

- **Phycological analysis:** classical study of morphology and anatomy.
- **Genetic analyses:** DNA-Barcoding & analysis of repetitive elements from the genome (microsatellites)
- **Measures of the DNA nuclear content (C-values):** current methodology by fluorimetry. Application of flux cytometry.
- **Karyotype analyses:** chromosome staining and Image analysis.
- **Species distribution analyses and niche comparison:** by maximum entropy algorithms and niche definition algorithms.

Sampling



Results



Conceptacle structure and germling growth. Fig. 6.

Mature oogonium containing egg cells ready to be released; Figs 7-8. Antheridia (Fig. 7) and germlings (Fig. 8) after 6 days in culture from Locality 1 (Illaunnginga) (possible damage to the attachment rhizoids due to transfer to microscope slide). Fig. 9. Immature oogonia from Locality 2 (Clifden). Fig. 10. Immature oogonia. Fig. 11. Antheridia both from Locality 3 (Achill Sound). Scale bar is 50 μm .

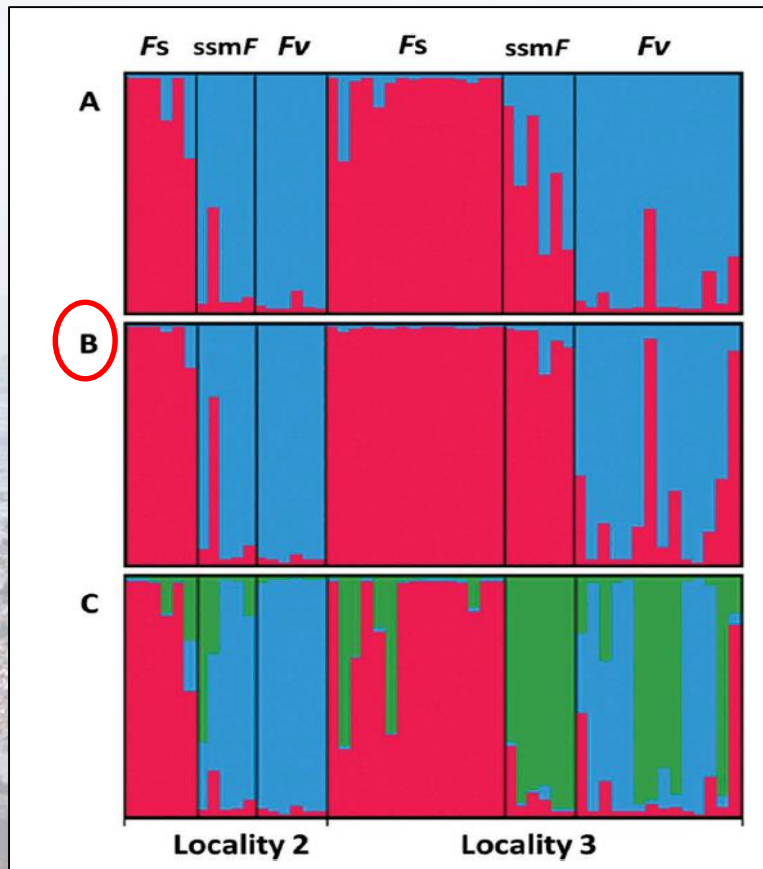
Table 2. Calculated F_{IS} , H_{exp} and allelic richness (\hat{A}) for n individuals from populations of *Fucus spiralis* (F_s), *F. vesiculosus* (F_v) and small salt marsh *Fucus* ($ssmF$) from Locality 2 (Clifden) and Locality 3 (Achill Sound).

Locality-taxon	n	F_{IS}	H_{exp}	\hat{A}
2- F_s	6	0.590*	0.327	1.400
2- $ssmF$	5	0.271	0.513	2.800
2- F_v	6	0.102	0.570	2.726
3- F_s	15	0.618*	0.295	1.472
3- $ssmF$	6	0.510*	0.374	2.600
3- F_v	14	0.504*	0.813	4.266

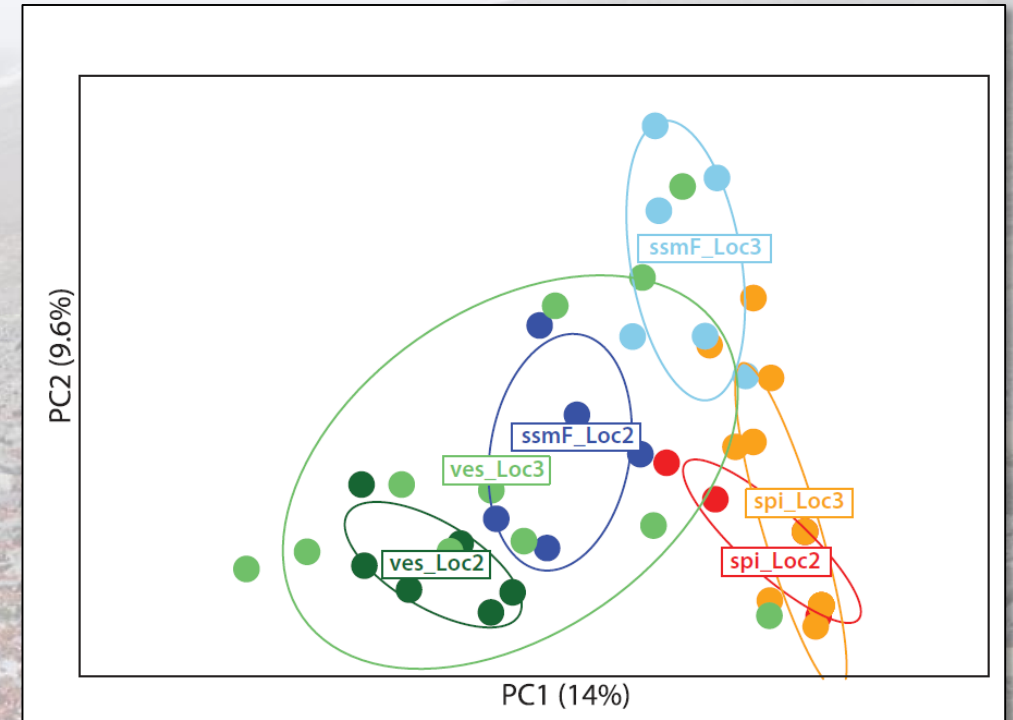
Significant F_{IS} values are marked with an asterisk. Allelic richness (\hat{A}) is standardized to $n = 5$.

Table 3. Pairwise F_{ST} values comparing populations of *Fucus spiralis* (F_s), *F. vesiculosus* (F_v) and small salt marsh *Fucus* ($ssmF$) from Locality 2 (Clifden) and Locality 3 (Achill Sound).

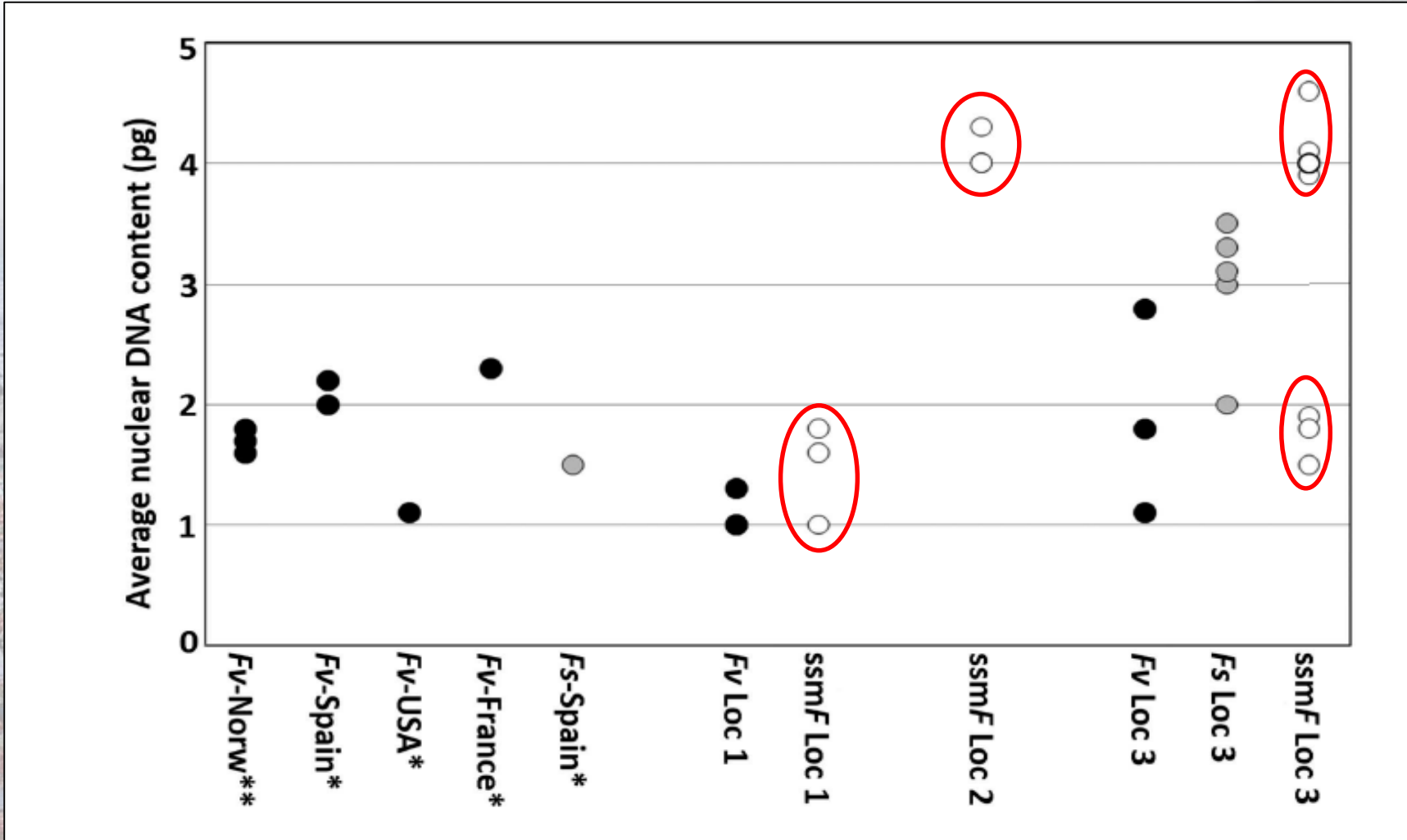
Locality-taxon	2- $ssmF$	2- F_v	3- F_s	3- $ssmF$	3- F_v
2- F_s	0.408	0.447	-0.059	0.469	0.238
2- $ssmF$		0.073	0.492	0.349	0.064
2- F_v			0.526	0.464	0.078
3- F_s				0.497	0.327
3- $ssmF$					0.182



Results from **STRUCTURE** (A) and **INSTRUCT** (B) analyses with $K = 2$; and from INSTRUCT analysis with $K = 3$ (C). Localities 2 and 3 consist of individuals sampled as *Fucus spiralis* (Fs), *F. vesiculosus* (Fv) and small salt marsh *Fucus* (ssmF). Each individual is represented by a bar and colours represent the proportional assignment to the STRUCTURE/INSTRUCT groups.



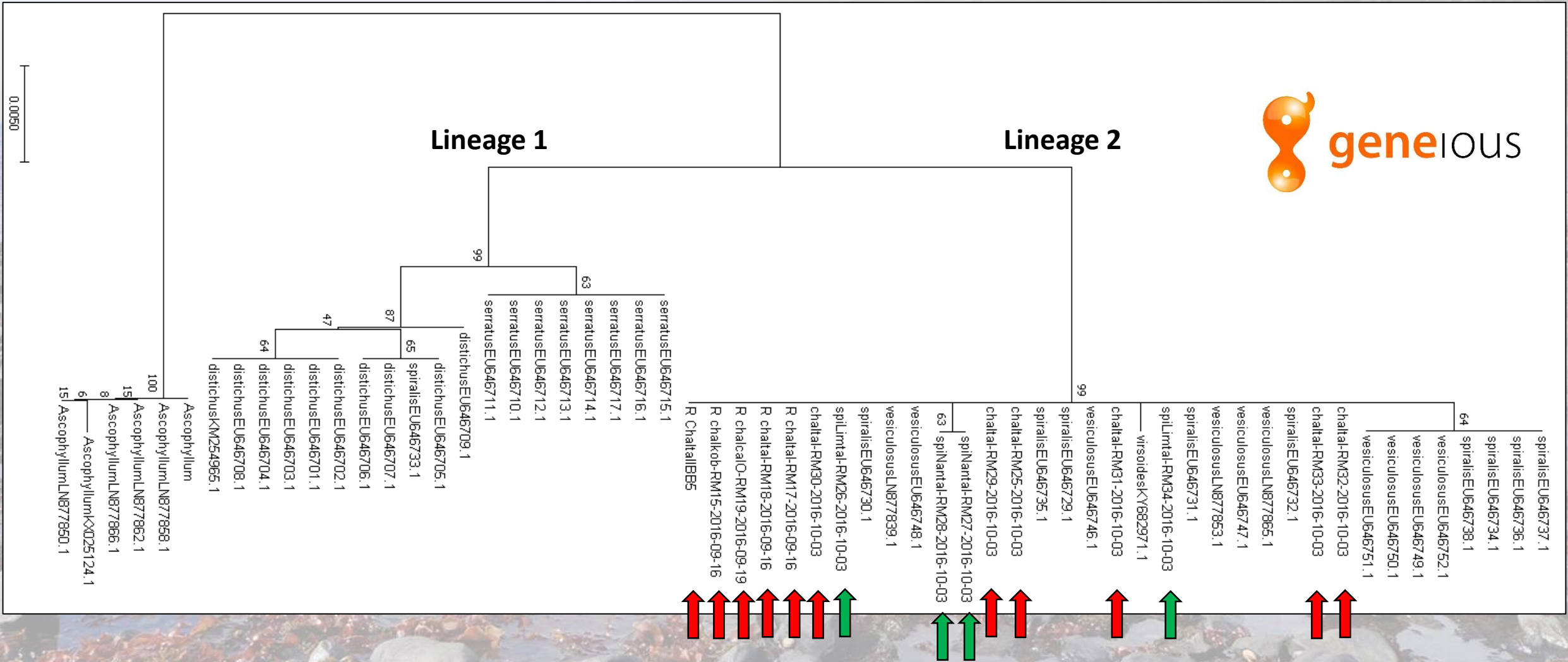
The two first principal components (PC) of a **Principal Component Analysis** showing genetic affiliation of haplotype composition of ssmF (blue), *F. vesiculosus* (ves) (green) and *F. spiralis* (spi) (orange-red) from Locality 2 (Loc2) and Locality 3 (Loc3). The circles represent 95% inertia ellipses for the populations, which characterizes the dispersion of each population around its centre of gravity. Percentages along axes indicate the proportion of overall variability explained by the principal components.



4x or 6x 8x?

Average nuclear DNA content (2C; pg) in *Fucus vesiculosus* (Fv, black circles), *F. spiralis* (Fs, grey circles) and small salt marsh *Fucus* (ssmF, white circles). Data from Illaunnginga (Locality 1), Clifden (Locality 2) and Achill Sound (Locality 3) are shown, together with unpublished data from Norway (**) and earlier published data (*) from Spain (Gómez Garreta *et al.*, 2010), USA (Kapraun, 2005) and France (Phillips *et al.*, 2011, recalculation from Peters *et al.*, 2004).

Source: Sjøtun *et al.* (2017)



Bayesian tree of COI-1 region for the studied species of *Fucus* plus several other species from the genus from GenBank

Discussion

- The **ssmF** showed high **variation** with respect to **reproduction mode, genetic affiliation** and **nuclear DNA content**.
- **ssmF can originate from different *Fucus* taxa** (Coyer *et al.*, 2006; Neiva *et al.*, 2012) . **This is supported by our microsatellite study** (in ex.: locality 2 ssmF derived mainly from *F. vesiculosus*, Locality 3 ssmF higher degree of hybridization/introgression)
- Kapraun, 2005; Gómez Garreta *et al.*, 2010; Phillips *et al.*, 2011 reported **variable genome size** for some brown algae, ¿**autopolyploids?**, ¿**allopolyploids?**

Near Future...

- Keep studying the composition of Antarctic flora quantitatively (Second BLUEBIO expedition)
- Finish the current Antarctic experiments and start new collaborations (role of seaweed in cloud formation & Passengers)
- Apply the methodologies showed to all the small *Fucus* forms and compare the results to understand the underlying processes of their evolution



Thank You!