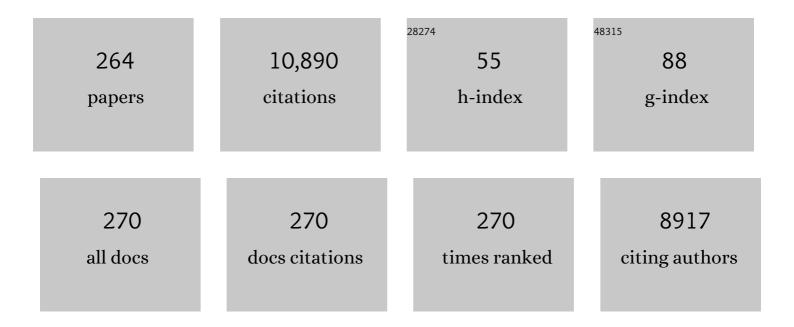
List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7811222/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Phenotypic Plasticity in Sargassum Forests May Not Counteract Projected Biomass Losses Along a Broad Latitudinal Gradient. Ecosystems, 2023, 26, 29-41.	3.4	1
2	Global biodiversity patterns of marine forests of brown macroalgae. Global Ecology and Biogeography, 2022, 31, 636-648.	5.8	22
3	Biogeographic Population Structure of Chimeric Blades of Porphyra in the Northeast Atlantic Reveals Southern Rich Gene Pools, Introgression and Cryptic Plasticity. Frontiers in Plant Science, 2022, 13, 818368.	3.6	1
4	Major Expansion of Marine Forests in a Warmer Arctic. Frontiers in Marine Science, 2022, 9, .	2.5	16
5	Ocean currents shape the genetic structure of a kelp in southwestern Africa. Journal of Biogeography, 2022, 49, 822-835.	3.0	9
6	eDNA metabarcoding for diet analyses of green sea turtles (Chelonia mydas). Marine Biology, 2022, 169, 1.	1.5	14
7	<strong>The genus <em>Cystoseira s.l.</em> (Ochrophyta, Fucales, Sargassaceae) in the Black Sea: morphological variability and molecular taxonomy of <em>Congolaria</em> <em>barbata</em> and endemic <em>Ericaria crinita</em> f. <em>bosphorica comb. nov.&lt;:/em&gt;&lt;:/strong&gt;:. Phytotaxa, 2021, 480, 1-21.</em></strong>	0.3	15
8	Bottom Trawling Threatens Future Climate Refugia of Rhodoliths Globally. Frontiers in Marine Science, 2021, 7, .	2.5	27
9	Phylogeographic Analysis Suggests a Recent Population Bottleneck in the Rare Red Sea Tridacna squamosina. Frontiers in Marine Science, 2021, 8, .	2.5	0
10	Mates Matter: Gametophyte Kinship Recognition and Inbreeding in the Giant Kelp, <i>Macrocystispyrifera</i> (Laminariales, Phaeophyceae). Journal of Phycology, 2021, 57, 711-725.	2.3	16
11	Genomes Vary in Size and Spatial Patterns Within Chimeric Blades of Porphyra spp Frontiers in Marine Science, 2021, 8, .	2.5	5
12	New Records of Fish Species from the Coast of Luanda, Angola. Thalassas, 2021, 37, 803-811.	0.5	0
13	Climateâ€induced range shifts shaped the present and threaten the future genetic variability of a marine brown alga in the Northwest Pacific. Evolutionary Applications, 2021, 14, 1867-1879.	3.1	12
14	Charting a course for genetic diversity in the UN Decade of Ocean Science. Evolutionary Applications, 2021, 14, 1497-1518.	3.1	19
15	Spatiotemporal patterns of phenology of the alien Phaeophyceae Sargassum muticum on the Atlantic coast of Morocco. Scientia Marina, 2021, 85, 103-111.	0.6	3
16	The collapse of marine forests: drastic reduction in populations of the family Sargassaceae in Madeira Island (NE Atlantic). Regional Environmental Change, 2021, 21, 1.	2.9	14
17	The microbiome of the habitatâ€forming brown alga <i>Fucus vesiculosus</i> (Phaeophyceae) has similar crossâ€Atlantic structure that reflects past and present drivers <sup>1</sup> . Journal of Phycology, 2021, 57, 1681-1698.	2.3	17
18	Characterization and Comparison of Bacterial Communities of an Invasive and Two Native Caribbean Seagrass Species Sheds Light on the Possible Influence of the Microbiome on Invasive Mechanisms. Frontiers in Microbiology, 2021, 12, 653998.	3.5	10

#	Article	IF	CITATIONS
19	Predicted regime shift in the seagrass ecosystem of the Gulf of Arguin driven by climate change. Global Ecology and Conservation, 2021, 32, e01890.	2.1	8
20	Development of tools to rapidly identify cryptic species and characterize their genetic diversity in different European kelp species. Journal of Applied Phycology, 2021, 33, 4169-4186.	2.8	2
21	Microbial Surface Biofilm Responds to the Growth-Reproduction-Senescence Cycle of the Dominant Coral Reef Macroalgae Sargassum spp Life, 2021, 11, 1199.	2.4	0
22	Potential Biodiversity Connectivity in the Network of Marine Protected Areas in Western Africa. Frontiers in Marine Science, 2021, 8, .	2.5	9
23	Warming Threatens to Propel the Expansion of the Exotic Seagrass Halophila stipulacea. Frontiers in Marine Science, 2021, 8, .	2.5	13
24	Environmental drivers of rhodolith beds and epiphytes community along the South Western Atlantic coast. Marine Environmental Research, 2020, 154, 104827.	2.5	38
25	How experimental physiology and ecological niche modelling can inform the management of marine bioinvasions?. Science of the Total Environment, 2020, 700, 134692.	8.0	10
26	Brazil oil spill response: Protect rhodolith beds. Science, 2020, 367, 156-156.	12.6	24
27	Congruence between fine-scale genetic breaks and dispersal potential in an estuarine seaweed across multiple transition zones. ICES Journal of Marine Science, 2020, 77, 371-378.	2.5	12
28	Linking Ecology to Genetics to Better Understand Adaptation and Evolution: A Review in Marine Macrophytes. Frontiers in Marine Science, 2020, 7, .	2.5	14
29	High Coral Bycatch in Bottom-Set Gillnet Coastal Fisheries Reveals Rich Coral Habitats in Southern Portugal. Frontiers in Marine Science, 2020, 7, .	2.5	14
30	Phylogeny and Evolution of the Brown Algae. Critical Reviews in Plant Sciences, 2020, 39, 281-321.	5.7	82
31	Spatial patterns of microbial communities across surface waters of the Great Barrier Reef. Communications Biology, 2020, 3, 442.	4.4	30
32	Ecological traits, genetic diversity and regional distribution of the macroalga Treptacantha elegans along the Catalan coast (NW Mediterranean Sea). Scientific Reports, 2020, 10, 19219.	3.3	11
33	Genetic structure of amphi-Atlantic <i>Laminaria digitata</i> (Laminariales, Phaeophyceae) reveals a unique range-edge gene pool and suggests post-glacial colonization of the NW Atlantic. European Journal of Phycology, 2020, 55, 517-528.	2.0	13
34	Spatiotemporal variation of the epifaunal assemblages associated to Sargassum muticum on the NW Atlantic coast of Morocco. Environmental Science and Pollution Research, 2020, 27, 35501-35514.	5.3	3
35	Thermal traits for reproduction and recruitment differ between Arctic and Atlantic kelp Laminaria digitata. PLoS ONE, 2020, 15, e0235388.	2.5	19
36	Genetic diversity of a marine foundation species, <i>Laminaria hyperborea</i> (Gunnerus) Foslie, along the coast of Ireland. European Journal of Phycology, 2020, 55, 310-326.	2.0	7

#	Article	IF	CITATIONS
37	Golden carbon of Sargassum forests revealed as an opportunity for climate change mitigation. Science of the Total Environment, 2020, 729, 138745.	8.0	68
38	A fine-tuned global distribution dataset of marine forests. Scientific Data, 2020, 7, 119.	5.3	45
39	The Small Giant Clam, Tridacna maxima Exhibits Minimal Population Genetic Structure in the Red Sea and Genetic Differentiation From the Gulf of Aden. Frontiers in Marine Science, 2020, 7, .	2.5	8
40	Microbiome dynamics in the tissue and mucus of acroporid corals differ in relation to host and environmental parameters. PeerJ, 2020, 8, e9644.	2.0	30
41	Genetic Affinities and Biogeography of Putative Levantine-Endemic Seaweed Treptacantha rayssiae (Ramon) M.Mulas, J.Neiva & Āilsrael, comb. nov. (Phaeophyceae). Cryptogamie, Algologie, 2020, 41, .	0.9	4
42	Hybrid vigour for thermal tolerance in hybrids between the allopatric kelps <i>Laminaria digitata</i> and <i>L. pallida</i> (Laminariales, Phaeophyceae) with contrasting thermal affinities. European Journal of Phycology, 2019, 54, 548-561.	2.0	32
43	Sex-dependent and -independent transcriptional changes during haploid phase gametogenesis in the sugar kelp Saccharina latissima. PLoS ONE, 2019, 14, e0219723.	2.5	15
44	Fineâ€scale genetic structure and flowering output of the seagrass <i>Enhalus acoroides</i> undergoing disturbance. Ecology and Evolution, 2019, 9, 5186-5195.	1.9	11
45	Integrating reproductive phenology in ecological niche models changed the predicted future ranges of a marine invader. Diversity and Distributions, 2019, 25, 688-700.	4.1	30
46	Toward a Coordinated Global Observing System for Seagrasses and Marine Macroalgae. Frontiers in Marine Science, 2019, 6, .	2.5	123
47	Open Coast Seagrass Restoration. Can We Do It? Large Scale Seagrass Transplants. Frontiers in Marine Science, 2019, 6, .	2.5	50
48	Increased evolutionary rates and conserved transcriptional response following allopolyploidization in brown algae. Evolution; International Journal of Organic Evolution, 2019, 73, 59-72.	2.3	11
49	Gene pool and connectivity patterns of <i>Pinna nobilis</i> in the Balearic Islands (Spain, Western) Tj ETQq1 1 Marine and Freshwater Ecosystems, 2019, 29, 175-188.	0.784314 2.0	rgBT /Overloci 9
50	Canopy microclimate modification in central and marginal populations of a marine macroalga. Marine Biodiversity, 2019, 49, 415-424.	1.0	23
51	Sexual reproduction vs. clonal propagation in the recovery of a seagrass meadow after an extreme weather event. Scientia Marina, 2019, 83, 357.	0.6	15
52	Genetic diversity increases with depth in red gorgonian populations of the Mediterranean Sea and the Atlantic Ocean. PeerJ, 2019, 7, e6794.	2.0	10
53	Genetic and oceanographic tools reveal high population connectivity and diversity in the endangered pen shell Pinna nobilis. Scientific Reports, 2018, 8, 4770.	3.3	31
54	Glacial vicariance drives phylogeographic diversification in the amphi-boreal kelp Saccharina latissima. Scientific Reports, 2018, 8, 1112.	3.3	61

#	Article	IF	CITATIONS
55	Bioâ€ORACLE v2.0: Extending marine data layers for bioclimatic modelling. Global Ecology and Biogeography, 2018, 27, 277-284.	5.8	567
56	Isolation and characterization of microsatellite markers for the red alga <i>Porphyra umbilicalis</i> . Plant Genetic Resources: Characterisation and Utilisation, 2018, 16, 390-393.	0.8	5
57	Small scale temporal patterns of recruitment and hatching of Atlantic horse mackerel (L.) at a nearshore reef area. Fisheries Oceanography, 2018, 27, 505-516.	1.7	5
58	Predicted extinction of unique genetic diversity in marine forests of Cystoseira spp Marine Environmental Research, 2018, 138, 119-128.	2.5	43
59	Projected climate changes threaten ancient refugia of kelp forests in the North Atlantic. Global Change Biology, 2018, 24, e55-e66.	9.5	140
60	Host Differentiation and Compartmentalization of Microbial Communities in the Azooxanthellate Cupcorals Tubastrea coccinea and Rhizopsammia goesi in the Caribbean. Frontiers in Marine Science, 2018, 5, .	2.5	25
61	The paranthurid isopod crustacean Paranthura nigropunctata (Lucas, 1846): first record from the Atlantic coast of Morocco. Acta Oceanologica Sinica, 2018, 37, 190-194.	1.0	2
62	Summer shifts of bacterial communities associated with the invasive brown seaweed Sargassum muticum are location and tissue dependent. PLoS ONE, 2018, 13, e0206734.	2.5	57
63	Individual-based genetic analyses support asexual hydrochory dispersal in Zostera noltei. PLoS ONE, 2018, 13, e0199275.	2.5	4
64	Differentiation in fitness-related traits in response to elevated temperatures between leading and trailing edge populations of marine macrophytes. PLoS ONE, 2018, 13, e0203666.	2.5	28
65	Harnessing positive species interactions as a tool against climate-driven loss of coastal biodiversity. PLoS Biology, 2018, 16, e2006852.	5.6	91
66	The introduction of <i>Sargassum muticum</i> modifies epifaunal patterns in a Moroccan seagrass meadow. Marine Ecology, 2018, 39, e12507.	1.1	8
67	Marine forests of the Mediterranean-Atlantic Cystoseira tamariscifolia complex show a southern Iberian genetic hotspot and no reproductive isolation in parapatry. Scientific Reports, 2018, 8, 10427.	3.3	25
68	Climate Change Impacts on Seagrass Meadows and Macroalgal Forests: An Integrative Perspective on Acclimation and Adaptation Potential. Frontiers in Marine Science, 2018, 5, .	2.5	149
69	Dramatic loss of seagrass habitat under projected climate change in the Mediterranean Sea. Global Change Biology, 2018, 24, 4919-4928.	9.5	140
70	Kelps' Long-Distance Dispersal: Role of Ecological/Oceanographic Processes and Implications to Marine Forest Conservation. Diversity, 2018, 10, 11.	1.7	34
71	Postglacial expansion of the Arctic keystone copepod Calanus glacialis. Marine Biodiversity, 2018, 48, 1027-1035.	1.0	15
72	Past climate changes and strong oceanographic barriers structured low″atitude genetic relics for the golden kelp <i>Laminaria ochroleuca</i> . Journal of Biogeography, 2018, 45, 2326-2336.	3.0	44

#	Article	IF	CITATIONS
73	Re-assessing the origins of the invasive mussel Mytilus galloprovincialis in southern Africa. Marine and Freshwater Research, 2018, 69, 607.	1.3	22
74	Polyploid lineages in the genus Porphyra. Scientific Reports, 2018, 8, 8696.	3.3	21
75	Seaweed Loads Cause Stronger Bacterial Community Shifts in Coastal Lagoon Sediments Than Nutrient Loads. Frontiers in Microbiology, 2018, 9, 3283.	3.5	25
76	Unraveling seaweeds bacteriomes. , 2018, , 95-113.		2
77	Acidification increases abundances of <i>Vibrionales</i> and <i>Planctomycetia</i> associated to a seaweed-grazer system: potential consequences for disease and prey digestion efficiency. PeerJ, 2018, 6, e4377.	2.0	16
78	Similar Epiphytic Macrofauna Inhabiting the Introduced <i>Sargassum muticum</i> and Native Fucoids on the Atlantic Coast of Morocco. Cryptogamie, Algologie, 2018, 39, 269-292.	0.9	5
79	Entangled fates of holobiont genomes during invasion: nested bacterial and host diversities in <i>Caulerpa taxifolia</i> . Molecular Ecology, 2017, 26, 2379-2391.	3.9	42
80	Population dynamics of temperate kelp forests near their low-latitude limit. Aquatic Botany, 2017, 139, 8-18.	1.6	9
81	The interaction between the proliferating macroalga Asparagopsis taxiformis and the coral Astroides calycularis induces changes in microbiome and metabolomic fingerprints. Scientific Reports, 2017, 7, 42625.	3.3	23
82	Cryptic diversity, geographical endemism and allopolyploidy in NE Pacific seaweeds. BMC Evolutionary Biology, 2017, 17, 30.	3.2	18
83	Interactions of daylength, temperature and nutrients affect thresholds for life stage transitions in the kelp Laminaria digitata (Phaeophyceae). Botanica Marina, 2017, 60, .	1.2	43
84	Regional Genetic Structure in the Aquatic Macrophyte Ruppia cirrhosa Suggests Dispersal by Waterbirds. Estuaries and Coasts, 2017, 40, 1705-1716.	2.2	16
85	Accounting for uncertainty in predictions of a marine species: Integrating population genetics to verify past distributions. Ecological Modelling, 2017, 359, 229-239.	2.5	19
86	Larval development and allometric growth of the blackâ€faced blenny <i>Tripterygion delaisi</i> . Journal of Fish Biology, 2017, 90, 2239-2254.	1.6	2
87	Major shifts at the range edge of marine forests: the combined effects of climate changes and limited dispersal. Scientific Reports, 2017, 7, 44348.	3.3	87
88	Development and characterization of twelve microsatellite markers for Porphyra linearis Greville. Genetica, 2017, 145, 127-130.	1,1	8
89	Habitat continuity and steppingâ€stone oceanographic distances explain population genetic connectivity of the brown alga <i>Cystoseira amentacea</i> . Molecular Ecology, 2017, 26, 766-780.	3.9	66
90	Mitochondrial genomes of the key zooplankton copepods Arctic Calanus glacialis and North Atlantic Calanus finmarchicus with the longest crustacean non-coding regions. Scientific Reports, 2017, 7, 13702.	3.3	9

#	Article	IF	CITATIONS
91	Palaeoclimatic conditions in the Mediterranean explain genetic diversity of Posidonia oceanica seagrass meadows. Scientific Reports, 2017, 7, 2732.	3.3	29
92	First record of Ruppia maritima in West Africa supported by morphological description and phylogenetic classification. Botanica Marina, 2017, 60, .	1.2	6
93	High genetic differentiation of red gorgonian populations from the Atlantic Ocean and the Mediterranean Sea. Marine Biology Research, 2017, 13, 854-861.	0.7	2
94	First description of seagrass distribution and abundance in São Tomé and PrÃncipe. Aquatic Botany, 2017, 142, 48-52.	1.6	13
95	Ampelisca lusitanica (Crustacea: Amphipoda): new species for the Atlantic coast of Morocco. Marine Biodiversity Records, 2017, 10, .	1.2	3
96	A population genetics toolbox for the threatened canopy-forming brown seaweeds Cystoseira tamariscifolia and C. amentacea (Fucales, Sargassaceae). Journal of Applied Phycology, 2017, 29, 627-629.	2.8	4
97	Reproductive strategies and population genetic structure of <i>Fucus spp</i> . across a northeast Atlantic biogeographic transition. Aquatic Living Resources, 2017, 30, 16.	1.2	7
98	Distribution and Genetic Structure of Fucus distichus Linnaeus 1953 (formerly F. gardneri) within Central San Francisco Bay. San Francisco Estuary and Watershed Science, 2017, 15, .	0.4	1
99	Species Specificity of Bacteria Associated to the Brown Seaweeds Lobophora (Dictyotales,) Tj ETQq1 1 0.784314 Frontiers in Microbiology, 2016, 7, 316.	rgBT /Ove 3.5	rlock 10 Tř 5 53
100	Host and Environmental Specificity in Bacterial Communities Associated to Two Highly Invasive Marine Species (Genus Asparagopsis). Frontiers in Microbiology, 2016, 7, 559.	3.5	72
101	Genetic Diversity and Local Connectivity in the Mediterranean Red Gorgonian Coral after Mass Mortality Events. PLoS ONE, 2016, 11, e0150590.	2.5	21
102	Upwelling areas as climate change refugia for the distribution and genetic diversity of a marine macroalga. Journal of Biogeography, 2016, 43, 1595-1607.	3.0	92
103	Temporal windows of reproductive opportunity reinforce species barriers in a marine broadcast spawning assemblage. Scientific Reports, 2016, 6, 29198.	3.3	17
104	Overlooked habitat of a vulnerable gorgonian revealed in the Mediterranean and Eastern Atlantic by ecological niche modelling. Scientific Reports, 2016, 6, 36460.	3.3	35
105	Pan-Arctic population of the keystone copepod Calanus glacialis. Polar Biology, 2016, 39, 2311-2318.	1.2	16
106	Deep reefs are climatic refugia for genetic diversity of marine forests. Journal of Biogeography, 2016, 43, 833-844.	3.0	84
107	Novel polymorphic microsatellite loci for a new target species, the sea cucumber Holothuria mammata. Biochemical Systematics and Ecology, 2016, 66, 109-113.	1.3	6
108	Taking the heat: distinct vulnerability to thermal stress of central and threatened peripheral lineages of a marine macroalga. Diversity and Distributions, 2016, 22, 1060-1068.	4.1	42

#	Article	IF	CITATIONS
109	First record of seagrass in Cape Verde, eastern Atlantic. Marine Biodiversity Records, 2016, 9, .	1.2	8
110	Multilocus genetic analyses provide insight into speciation and hybridization in aquatic grasses, genus <i>Ruppia</i> . Biological Journal of the Linnean Society, 2016, 117, 177-191.	1.6	18
111	Genetic diversity of <i>Saccharina latissima</i> (Phaeophyceae) along a salinity gradient in the North Sea–Baltic Sea transition zone. Journal of Phycology, 2016, 52, 523-531.	2.3	34
112	Early life history of larvae and early juvenile Atlantic horse mackerel Trachurus trachurus off the Portuguese west coast. Fisheries Research, 2016, 183, 111-118.	1.7	10
113	Large-Scale Prediction of Seagrass Distribution Integrating Landscape Metrics and Environmental Factors: The Case of Cymodocea nodosa (Mediterranean–Atlantic). Estuaries and Coasts, 2016, 39, 123-137.	2.2	51
114	Do hatchery-reared sea urchins pose a threat to genetic diversity in wild populations?. Heredity, 2016, 116, 378-383.	2.6	17
115	Future climate change is predicted to shift long-term persistence zones in the cold-temperate kelp Laminaria hyperborea. Marine Environmental Research, 2016, 113, 174-182.	2.5	67
116	Setting preliminary biometric baselines for new target sea cucumbers species of the NE Atlantic and Mediterranean fisheries. Fisheries Research, 2016, 179, 57-66.	1.7	37
117	Characterization of 12 polymorphic microsatellite markers in the sugar kelp Saccharina latissima. Journal of Applied Phycology, 2016, 28, 3071-3074.	2.8	22
118	Limited differences in fish and benthic communities and possible cascading effects inside and outside a protected marine area in Sagres (SW Portugal). Marine Environmental Research, 2016, 114, 12-23.	2.5	16
119	Climate Oscillations, Range Shifts and Phylogeographic Patterns of North Atlantic Fucaceae. , 2016, , 279-308.		27
120	The effect of mixotrophy in the ex situ culture of the soft coral Sarcophyton cf. glaucum. Aquaculture, 2016, 452, 151-159.	3.5	15
121	Comparison of small remotely operated vehicles and diver-operated video of circalittoral benthos. Hydrobiologia, 2016, 766, 247-260.	2.0	30
122	A Well-Kept Treasure at Depth: Precious Red Coral Rediscovered in Atlantic Deep Coral Gardens (SW) Tj ETQq0 0	0 rgBT /O	werlock 10 Tf
123	High Interannual Variability in Connectivity and Genetic Pool of a Temperate Clingfish Matches Oceanographic Transport Predictions. PLoS ONE, 2016, 11, e0165881.	2.5	16
124	Lack of fine-scale genetic structure and distant mating in natural populations of Fucus vesiculosus. Marine Ecology - Progress Series, 2016, 544, 131-142.	1.9	10
125	Tradeâ€offs between lifeâ€history traits at rangeâ€edge and central locations. Journal of Phycology, 2015, 51, 808-818.	2.3	16
126	Seascape drivers of <i><scp>M</scp>acrocystis pyrifera</i> population genetic structure in the	3.9	55

northeast <scp>P</scp>acific. Molecular Ecology, 2015, 24, 4866-4885. 126

#	Article	IF	CITATIONS
127	Oceanographic Conditions Limit the Spread of a Marine Invader along Southern African Shores. PLoS ONE, 2015, 10, e0128124.	2.5	58
128	Closer to the rear edge: ecology and genetic diversity down the coreâ€edge gradient of a marine macroalga. Ecosphere, 2015, 6, 1-25.	2.2	39
129	Behind the mask: cryptic genetic diversity of <i>Mytilus galloprovincialis</i> along southern European and northern African shores. Journal of Molluscan Studies, 2015, 81, 380-387.	1.2	16
130	Genetic diversity and biogeographical patterns of Caulerpa prolifera across the Mediterranean and Mediterranean/Atlantic transition zone. Marine Biology, 2015, 162, 557-569.	1.5	9
131	Some don't like it hot: microhabitatâ€dependent thermal and water stresses in a trailing edge population. Functional Ecology, 2015, 29, 640-649.	3.6	33
132	A transcriptome resource for Antarctic krill (Euphausia superba Dana) exposed to short-term stress. Marine Genomics, 2015, 23, 45-47.	1.1	8
133	Hologenome theory supported by cooccurrence networks of species-specific bacterial communities in siphonous algae ( <i>Caulerpa</i> ). FEMS Microbiology Ecology, 2015, 91, fiv067.	2.7	55
134	Contrasting timing of life stages across latitudes – a case study of a marine forest-forming species. European Journal of Phycology, 2015, 50, 361-369.	2.0	7
135	Metatranscriptomes reveal functional variation in diatom communities from the Antarctic Peninsula. ISME Journal, 2015, 9, 2275-2289.	9.8	55
136	A transcriptome resource for the copepod Calanus glacialis across a range of culture temperatures. Marine Genomics, 2015, 23, 27-29.	1.1	10
137	Polymorphic microsatellite markers in the brown seaweed Fucus vesiculosus. BMC Research Notes, 2015, 8, 73.	1.4	2
138	Intraspecific genetic lineages of a marine mussel show behavioural divergence and spatial segregation over a tropical/subtropical biogeographic transition. BMC Evolutionary Biology, 2015, 15, 100.	3.2	24
139	Reproductive investment, synchrony and recruitment success in marine broadcast spawners: Effects of mating system and habitat (exposed shore versus estuary). Marine Environmental Research, 2015, 112, 33-39.	2.5	2
140	European seaweeds under pressure: Consequences for communities and ecosystem functioning. Journal of Sea Research, 2015, 98, 91-108.	1.6	155
141	Response of kelps from different latitudes to consecutive heat shock. Journal of Experimental Marine Biology and Ecology, 2015, 463, 57-62.	1.5	25
142	West <i>versus </i> <scp>E</scp> ast <scp>M</scp> editerranean <scp>S</scp> ea: origin and genetic differentiation of the sea cucumber <i><scp>H</scp>olothuria polii</i> . Marine Ecology, 2015, 36, 485-495.	1.1	24
143	Highly polymorphic microsatellite markers for the Mediterranean endemic fan mussel Pinna nobilis. Mediterranean Marine Science, 2015, 16, 31.	1.6	13
144	Genes Left Behind: Climate Change Threatens Cryptic Genetic Diversity in the Canopy-Forming Seaweed Bifurcaria bifurcata. PLoS ONE, 2015, 10, e0131530.	2.5	52

#	Article	IF	CITATIONS
145	Spatial and Temporal Dynamics of Fucoid Populations (Ascophyllum nodosum and Fucus serratus): A Comparison between Central and Range Edge Populations. PLoS ONE, 2014, 9, e92177.	2.5	24
146	Microsatellite markers for the Arctic copepod Calanus glacialis and cross-amplification with C. finmarchicus. Conservation Genetics Resources, 2014, 6, 1003-1005.	0.8	4
147	New highly polymorphic microsatellite markers for the aquatic angiosperm <i>Ruppia cirrhosa</i> reveal population diversity and differentiation. Genome, 2014, 57, 57-59.	2.0	12
148	Reproductive strategies and isolationâ€byâ€demography in a marine clonal plant along an eutrophication gradient. Molecular Ecology, 2014, 23, 5698-5711.	3.9	14
149	Extending the life history of a clonal aquatic plant: Dispersal potential of sexual and asexual propagules of Zostera noltii. Aquatic Botany, 2014, 113, 123-129.	1.6	34
150	Genetic Divergence for the Amphibian Pleurodeles waltl in Southwest Portugal: Dispersal Barriers Shaping Geographic Patterns. Journal of Herpetology, 2014, 48, 38.	0.5	8
151	Species distribution models and mitochondrial <scp>DNA</scp> phylogeography suggest an extensive biogeographical shift in the highâ€intertidal seaweed <i>Pelvetia canaliculata</i> . Journal of Biogeography, 2014, 41, 1137-1148.	3.0	61
152	Climateâ€driven range shifts explain the distribution of extant gene pools and predict future loss of unique lineages in a marine brown alga. Molecular Ecology, 2014, 23, 2797-2810.	3.9	77
153	Reprint of "Seagrasses in Portugal: A most endangered marine habitat― Aquatic Botany, 2014, 115, 3-13.	1.6	10
154	Wider sampling reveals a nonâ€sister relationship for geographically contiguous lineages of a marine mussel. Ecology and Evolution, 2014, 4, 2070-2081.	1.9	33
155	Biomares, a LIFE project to restore and manage the biodiversity of Prof. Luiz Saldanha Marine Park. Journal of Coastal Conservation, 2014, 18, 643-655.	1.6	14
156	Characterization of fifteen microsatellite markers for the kelp Laminaria ochroleuca and cross species amplification within the genus. Conservation Genetics Resources, 2014, 6, 949-950.	0.8	6
157	Disentangling the Influence of Mutation and Migration in Clonal Seagrasses Using the Genetic Diversity Spectrum for Microsatellites. Journal of Heredity, 2014, 105, 532-541.	2.4	28
158	Genetic signature of a recent invasion: The ragged sea hare Bursatella leachii in Mar Menor (SE Spain). Biochemical Systematics and Ecology, 2014, 54, 123-129.	1.3	11
159	Microsatellite markers developed through pyrosequencing allow clonal discrimination in the invasive alga Caulerpa taxifolia. Conservation Genetics Resources, 2013, 5, 667-669.	0.8	6
160	Characterization of 15 polymorphic microsatellite loci in the temperate reef fish Lepadogaster lepadogaster, developed using 454-sequencing. Conservation Genetics Resources, 2013, 5, 55-57.	0.8	2
161	Shift happens: trailing edge contraction associated with recent warming trends threatens a distinct genetic lineage in the marine macroalga Fucus vesiculosus. BMC Biology, 2013, 11, 6.	3.8	130
162	Broad scale agreement between intertidal habitats and adaptive traits on a basis of contrasting population genetic structure. Estuarine, Coastal and Shelf Science, 2013, 131, 140-148.	2.1	34

#	Article	IF	CITATIONS
163	Looking into the black box: simulating the role of selfâ€fertilization and mortality in the genetic structure of <i>Macrocystis pyrifera</i> . Molecular Ecology, 2013, 22, 4842-4854.	3.9	17
164	High connectivity across the fragmented chemosynthetic ecosystems of the deep <scp>A</scp> tlantic <scp>E</scp> quatorial <scp>B</scp> elt: efficient dispersal mechanisms or questionable endemism?. Molecular Ecology, 2013, 22, 4663-4680.	3.9	51
165	Connectivity, neutral theories and the assessment of species vulnerability to global change in temperate estuaries. Estuarine, Coastal and Shelf Science, 2013, 131, 52-63.	2.1	28
166	Entangled effects of allelic and clonal (genotypic) richness in the resistance and resilience of experimental populations of the seagrass Zostera noltii to diatom invasion. BMC Ecology, 2013, 13, 39.	3.0	43
167	Polar marine biology science in Portugal and Spain: Recent advances and future perspectives. Journal of Sea Research, 2013, 83, 9-29.	1.6	15
168	Seagrasses in Portugal: A most endangered marine habitat. Aquatic Botany, 2013, 104, 193-203.	1.6	79
169	Performing fish counts with a wide-angle camera, a promising approach reducing divers' limitations. Journal of Experimental Marine Biology and Ecology, 2013, 445, 93-98.	1.5	35
170	Comparison of phototrophic shell-degrading endoliths in invasive and native populations of the intertidal mussel Mytilus galloprovincialis. Biological Invasions, 2013, 15, 1253-1272.	2.4	29
171	Genetic Evidence for Polygynandry in the Black-Striped Pipefish Syngnathus abaster: A Microsatellite-Based Parentage Analysis. Journal of Heredity, 2013, 104, 791-797.	2.4	6
172	High and Distinct Range-Edge Genetic Diversity despite Local Bottlenecks. PLoS ONE, 2013, 8, e68646.	2.5	90
173	Invasion Is a Community Affair: Clandestine Followers in the Bacterial Community Associated to Green Algae, Caulerpa racemosa, Track the Invasion Source. PLoS ONE, 2013, 8, e68429.	2.5	63
174	First record of the brown mussel (Perna perna) from the European Atlantic coast. Marine Biodiversity Records, 2012, 5, .	1.2	22
175	<i>Fucus cottonii</i> (Fucales, Phaeophyceae) is not a single genetic entity but a convergent salt-marsh morphotype with multiple independent origins. European Journal of Phycology, 2012, 47, 461-468.	2.0	13
176	Spatial synchronies in the seasonal occurrence of larvae of oysters (Crassostrea gigas) and mussels (Mytilus edulis/galloprovincialis) in European coastal waters. Estuarine, Coastal and Shelf Science, 2012, 108, 52-63.	2.1	31
177	Genetic flow directionality and geographical segregation in a Cymodocea nodosa genetic diversity network. EPJ Data Science, 2012, 1, .	2.8	14
178	Fine-scale genetic breaks driven by historical range dynamics and ongoing density-barrier effects in the estuarine seaweed Fucus ceranoides L. BMC Evolutionary Biology, 2012, 12, 78.	3.2	44
179	Characterization of ten highly polymorphic microsatellite loci for the intertidal mussel Perna perna, and cross species amplification within the genus. BMC Research Notes, 2012, 5, 558.	1.4	6
180	Implications of Extreme Life Span in Clonal Organisms: Millenary Clones in Meadows of the Threatened Seagrass Posidonia oceanica. PLoS ONE, 2012, 7, e30454.	2.5	195

#	Article	IF	CITATIONS
181	Prezygotic Barriers to Hybridization in Marine Broadcast Spawners: Reproductive Timing and Mating System Variation. PLoS ONE, 2012, 7, e35978.	2.5	22
182	Panmixia in a Fragmented and Unstable Environment: The Hydrothermal Shrimp Rimicaris exoculata Disperses Extensively along the Mid-Atlantic Ridge. PLoS ONE, 2012, 7, e38521.	2.5	59
183	Love Thy Neighbour: Group Properties of Gaping Behaviour in Mussel Aggregations. PLoS ONE, 2012, 7, e47382.	2.5	57
184	Mediterranean Species of Caulerpa Are Polyploid with Smaller Genomes in the Invasive Ones. PLoS ONE, 2012, 7, e47728.	2.5	24
185	Characterization of 15 polymorphic microsatellite loci in Rimicaris exoculata, and cross-amplification in other hydrothermal-vent shrimp. Conservation Genetics Resources, 2012, 4, 81-84.	0.8	2
186	Drifting fronds and drifting alleles: range dynamics, local dispersal and habitat isolation shape the population structure of the estuarine seaweed <i>Fucus ceranoides</i> . Journal of Biogeography, 2012, 39, 1167-1178.	3.0	48
187	SELECTIVE ELIMINATION OF CHLOROPLASTIDIAL DNA FOR METAGENOMICS OF BACTERIA ASSOCIATED WITH THE GREEN ALGA <i>CAULERPA TAXIFOLIA</i> (BRYOPSIDOPHYCEAE) <sup>1</sup> . Journal of Phycology, 2012, 48, 483-490.	2.3	19
188	The role of disturbance in differential regulation of co-occurring brown algae species: Interactive effects of sediment deposition, abrasion and grazing on algae recruits. Journal of Experimental Marine Biology and Ecology, 2012, 422-423, 1-8.	1.5	14
189	Rangeâ€edge genetic diversity: locally poor extant southern patches maintain a regionally diverse hotspot in the seagrass <i>Zostera marina</i> . Molecular Ecology, 2012, 21, 1647-1657.	3.9	68
190	Recovery after trampling disturbance in a canopy-forming seaweed population. Marine Biology, 2012, 159, 697-707.	1.5	21
191	Characterization of ten highly polymorphic microsatellite loci for the intertidal mussel Perna perna, and cross species amplification within the genus. BMC Research Notes, 2012, 5, 2101791285670501.	1.4	0
192	Expressed sequence tags from heat-shocked seagrass Zostera noltii (Hornemann) from its southern distribution range. Marine Genomics, 2011, 4, 181-188.	1.1	29
193	Development and characterization of 35 single nucleotide polymorphism markers for the brown alga <i>Fucus vesiculosus</i> . European Journal of Phycology, 2011, 46, 342-351.	2.0	4
194	Recent population expansion and connectivity in the hydrothermal shrimp Rimicaris exoculata along the Mid-Atlantic Ridge. Journal of Biogeography, 2011, 38, 564-574.	3.0	57
195	Isolation by oceanographic distance explains genetic structure for <i>Macrocystis pyrifera</i> in the Santa Barbara Channel. Molecular Ecology, 2011, 20, 2543-2554.	3.9	102
196	PHENOTYPIC DIFFERENTIATION AT SOUTHERN LIMIT BORDERS: THE CASE STUDY OF TWO FUCOID MACROALGAL SPECIES WITH DIFFERENT LIFE-HISTORY TRAITS1. Journal of Phycology, 2011, 47, 451-462.	2.3	36
197	Evolution and diversification within the intertidal brown macroalgae Fucus spiralis/F. vesiculosus species complex in the North Atlantic. Molecular Phylogenetics and Evolution, 2011, 58, 283-296.	2.7	65
198	Driving south: a multi-gene phylogeny of the brown algal family Fucaceae reveals relationships and recent drivers of a marine radiation. BMC Evolutionary Biology, 2011, 11, 371.	3.2	53

#	Article	IF	CITATIONS
199	Dinucleotide microsatellite markers in the genus Caulerpa. Journal of Applied Phycology, 2011, 23, 715-719.	2.8	6
200	Travelling in time with networks: Revealing present day hybridization versus ancestral polymorphism between two species of brown algae, Fucus vesiculosus and F. spiralis. BMC Evolutionary Biology, 2011, 11, 33.	3.2	23
201	Adaptive Traits Are Maintained on Steep Selective Gradients despite Gene Flow and Hybridization in the Intertidal Zone. PLoS ONE, 2011, 6, e19402.	2.5	86
202	Evolutionary history of the seagrass genus Posidonia. Marine Ecology - Progress Series, 2011, 421, 117-130.	1.9	40
203	The possible origin of Zostera noltii in the Canary Islands and guidelines for restoration. Marine Biology, 2010, 157, 2109-2115.	1.5	17
204	An Expressed Sequence Tag Analysis of the Intertidal Brown Seaweeds Fucus serratus (L.) and F. vesiculosus (L.) (Heterokontophyta, Phaeophyceae) in Response to Abiotic Stressors. Marine Biotechnology, 2010, 12, 195-213.	2.4	77
205	Comparative Analysis of Stability—Genetic Diversity in Seagrass (Posidonia oceanica) Meadows Yields Unexpected Results. Estuaries and Coasts, 2010, 33, 878-889.	2.2	51
206	Genetic structure in the Mediterranean seagrass <i>Posidonia oceanica</i> : disentangling past vicariance events from contemporary patterns of gene flow. Molecular Ecology, 2010, 19, 557-568.	3.9	101
207	Surfing the wave on a borrowed board: range expansion and spread of introgressed organellar genomes in the seaweed <i>Fucus ceranoides</i> L Molecular Ecology, 2010, 19, 4812-4822.	3.9	61
208	Habitat continuity and geographic distance predict population genetic differentiation in giant kelp. Ecology, 2010, 91, 49-56.	3.2	81
209	Fucus vesiculosus and spiralis species complex: a nested model of local adaptation at the shore level. Marine Ecology - Progress Series, 2010, 405, 163-174.	1.9	44
210	Temperature tolerance and survival of intertidal populations of the seagrass Zostera noltii (Hornemann) in Southern Europe (Ria Formosa, Portugal). Hydrobiologia, 2009, 619, 195-201.	2.0	78
211	Highly polymorphic microsatellite loci for the Parsley frog (Pelodytes punctatus): characterization and testing for cross-species amplification. Conservation Genetics, 2009, 10, 665-668.	1.5	7
212	Development and characterization of highly polymorphic microsatellite loci for the Western Spadefoot toad, Pelobates cultripes. Conservation Genetics, 2009, 10, 993-996.	1.5	3
213	Microsatellite markers for the giant kelp Macrocystis pyrifera. Conservation Genetics, 2009, 10, 1915-1917.	1.5	16
214	Highly polymorphic microsatellite markers for the short-snouted seahorse (HippocampusÀhippocampus), including markers from a closely related species the long-snouted seahorse (HippocampusÂguttulatus). Conservation Genetics Resources, 2009, 1, 93-96.	0.8	9
215	HABITAT DIFFERENCES IN THE TIMING OF REPRODUCTION OF THE INVASIVE ALGA <i>SARGASSUM MUTICUM</i> (PHAEOPHYTA, SARGASSACEAE) OVER TIDAL AND LUNAR CYCLES <sup>1</sup> . Journal of Phycology, 2009, 45, 1-7.	2.3	17

216 Fertilization Strategies. Ecological Studies, 2009, , 149-164.

1.2 12

#	Article	IF	CITATIONS
217	Isolation of highly polymorphic microsatellite loci for a species with a large genome size: sharpâ€ribbed salamander ( <i>Pleurodeles waltl</i> ). Molecular Ecology Resources, 2009, 9, 425-428.	4.8	12
218	Permanent Genetic Resources added to Molecular Ecology Resources Database 1 May 2009–31 July 2009. Molecular Ecology Resources, 2009, 9, 1460-1466.	4.8	128
219	Effects of disturbance on marginal populations: human trampling on Ascophyllum nodosum assemblages at its southern distribution limit. Marine Ecology - Progress Series, 2009, 378, 81-92.	1.9	35
220	Genetic recolonization of mangrove: genetic diversity still increasing in the Mekong Delta 30 years after Agent Orange. Marine Ecology - Progress Series, 2009, 390, 129-135.	1.9	18
221	Egg release and settlement patterns of dioecious and hermaphroditic fucoid algae during the tidal cycle. Marine Biology, 2008, 155, 583-591.	1.5	15
222	Genetic differentiation and secondary contact zone in the seagrass <i>Cymodocea nodosa</i> across the Mediterranean–Atlantic transition region. Journal of Biogeography, 2008, 35, 1279-1294.	3.0	105
223	ECOLOGICAL GENETICS IN THE NORTH ATLANTIC: ENVIRONMENTAL GRADIENTS AND ADAPTATION AT SPECIFIC LOCI. Ecology, 2008, 89, S91-107.	3.2	124
224	Periodicity of propagule expulsion and settlement in the competing native and invasive brown seaweeds, <b><i>Cystoseira humilis</i></b> and <b><i>Sargassum muticum</i></b> (Phaeophyta). European Journal of Phycology, 2008, 43, 275-282.	2.0	34
225	Network analysis identifies weak and strong links in a metapopulation system. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 18824-18829.	7.1	152
226	THE EVOLUTION OF CICADA SONGS CONTRASTED WITH THE RELATIONSHIPS INFERRED FROM MITOCHONDRIAL DNA (INSECTA, HEMIPTERA). Bioacoustics, 2008, 18, 17-34.	1.7	10
227	Evolutionary and Ecological Trees and Networks. AIP Conference Proceedings, 2007, , .	0.4	3
228	Implications of mating system for genetic diversity of sister algal species: <b><i>Fucus spiralis</i></b> and <b><i>Fucus vesiculosus</i></b> (Heterokontophyta, Phaeophyceae). European Journal of Phycology, 2007, 42, 219-230.	2.0	47
229	Spectrum of genetic diversity and networks of clonal organisms. Journal of the Royal Society Interface, 2007, 4, 1093-1102.	3.4	72
230	Standardizing methods to address clonality in population studies. Molecular Ecology, 2007, 16, 5115-5139.	3.9	568
231	Vicariance patterns in the Mediterranean Sea: east–west cleavage and low dispersal in the endemic seagrass Posidonia oceanica. Journal of Biogeography, 2007, 34, 963-976.	3.0	159
232	Genetic sub-structure and intermediate optimal outcrossing distance in the marine angiosperm Zostera marina. Marine Biology, 2007, 152, 793-801.	1.5	35
233	Feed-backs between genetic structure and perturbation-driven decline in seagrass (Posidonia) Tj ETQq1 1 0.7843	14 rgBT / 1.5	Overlock 10 47
234	Convergent adaptation to a marginal habitat by homoploid hybrids and polyploid ecads in the seaweed	2.3	54

#	Article	IF	CITATIONS
235	Timing and success of reproductive stages in the seagrass Zostera noltii. Aquatic Botany, 2006, 85, 219-223.	1.6	25
236	GENOMIC DNA ISOLATION FROM GREEN AND BROWN ALGAE (CAULERPALES AND FUCALES) FOR MICROSATELLITE LIBRARY CONSTRUCTION1. Journal of Phycology, 2006, 42, 741-745.	2.3	60
237	When is a hybrid a hybrid? A counter-reply to Wallace etÂal Molecular Ecology, 2006, 15, 3481-3482.	3.9	4
238	Genetic structure at range edge: low diversity and high inbreeding in Southeast Asian mangrove (Avicennia marina) populations. Molecular Ecology, 2006, 15, 3515-3525.	3.9	173
239	Simple and rapid RNA extraction from freeze-dried tissue of brown algae and seagrasses. European Journal of Phycology, 2006, 41, 97-104.	2.0	60
240	Revisiting synchronous gamete release by fucoid algae in the intertidal zone: fertilization success and beyond?. Integrative and Comparative Biology, 2006, 46, 587-597.	2.0	57
241	Genetic diversity of a clonal angiosperm near its range limit: the case of Cymodocea nodosa at the Canary Islands. Marine Ecology - Progress Series, 2006, 309, 117-129.	1.9	53
242	Intriguing asexual life in marginal populations of the brown seaweed Fucus vesiculosus. Molecular Ecology, 2005, 14, 647-651.	3.9	115
243	Genetic entities and mating system in hermaphroditic Fucus spiralis and its close dioecious relative F. vesiculosus (Fucaceae, Phaeophyceae). Molecular Ecology, 2005, 14, 2033-2046.	3.9	74
244	Within-population spatial genetic structure, neighbourhood size and clonal subrange in the seagrass Cymodocea nodosa. Molecular Ecology, 2005, 14, 2669-2681.	3.9	123
245	GENETIC ISOLATION BETWEEN THREE CLOSELY RELATED TAXA: FUCUS VESICULOSUS, F. SPIRALIS, AND F. CERANOIDES (PHAOPHYCEAE)1. Journal of Phycology, 2005, 41, 900-905.	2.3	40
246	Analysis of sexual phenotype and prezygotic fertility in natural populations ofFucus spiralis, F. vesiculosus(Fucaceae, Phaeophyceae) and their putative hybrids. European Journal of Phycology, 2005, 40, 397-407.	2.0	33
247	Assessing Genetic Diversity in Clonal Organisms: Low Diversity or Low Resolution? Combining Power and Cost Efficiency in Selecting Markers. Journal of Heredity, 2005, 96, 434-440.	2.4	156
248	North Atlantic phylogeography and large-scale population differentiation of the seagrass Zostera marina L Molecular Ecology, 2004, 13, 1923-1941.	3.9	277
249	Characterization of microsatellite loci in the dwarf eelgrass Zostera noltii (Zosteraceae) and cross-reactivity with Z. japonica. Molecular Ecology Notes, 2004, 4, 497-499.	1.7	25
250	Blue- and green-light signals for gamete release in the brown alga, Silvetia compressa. Oecologia, 2004, 138, 193-201.	2.0	22
251	Isolation and cross-species amplification of microsatellite loci from the fucoid seaweeds Fucus vesiculosus, F. serratus and Ascophyllum nodosum (Heterokontophyta, Fucaceae). Molecular Ecology Notes, 2003, 3, 180-182.	1.7	61
252	New microsatellite markers for the endemic Mediterranean seagrass Posidonia oceanica. Molecular Ecology Notes, 2003, 3, 253-255.	1.7	35

#	Article	IF	CITATIONS
253	Isolation and characterization of microsatellite markers for the seagrassCymodocea nodosa. Molecular Ecology Notes, 2003, 3, 397-399.	1.7	14
254	Polymorphic microsatellite DNA markers in the mangrove tree Avicennia alba. Molecular Ecology Notes, 2003, 3, 544-546.	1.7	8
255	Spatial patterns of groundfish assemblages on the continental shelf of Portugal. ICES Journal of Marine Science, 2001, 58, 633-647.	2.5	70
256	Suppression subtractive hybridization for studying gene expression during aerial exposure and desiccation in fucoid algae. European Journal of Phycology, 2001, 36, 359-366.	2.0	20
257	REPRODUCTIVE SUCCESS OF FUCUS VESICULOSUS (PHAEOPHYCEAE) IN THE BALTIC SEA. Journal of Phycology, 1999, 35, 254-269.	2.3	90
258	EVOLUTION OF THE FUCACEAE (PHAEOPHYCEAE) INFERRED FROM nrDNA-ITS. Journal of Phycology, 1999, 35, 382-394.	2.3	141
259	CONTROL OF GAMETE RELEASE IN FUCOID ALGAE: SENSING HYDRODYNAMIC CONDITIONS VIA CARBON ACQUISITION. Ecology, 1998, 79, 1725-1739.	3.2	89
260	Successful external fertilization in turbulent environments Proceedings of the National Academy of Sciences of the United States of America, 1996, 93, 5286-5290.	7.1	145
261	Distributional success of the marine seaweedFucus vesiculosus L. in the brackish Baltic Sea correlates with osmotic capabilities of Baltic gametes. Oecologia, 1996, 107, 1-12.	2.0	106
262	Deep-water macroalgae from the Canary Islands: new records and biogeographical relationships. Helgolâ^šÂ§nder Meeresuntersuchungen, 1993, 47, 125-143.	0.2	15
263	Microscopic life stages of Arctic kelp differ in their resilience and reproductive output in response to Arctic seasonality. European Journal of Phycology, 0, , 1-15.	2.0	4
264	Seagrass Connectivity on the West Coast of Africa Supports the Hypothesis of Grazer-Mediated Seed Dispersal. Frontiers in Marine Science, 0, 9, .	2.5	3