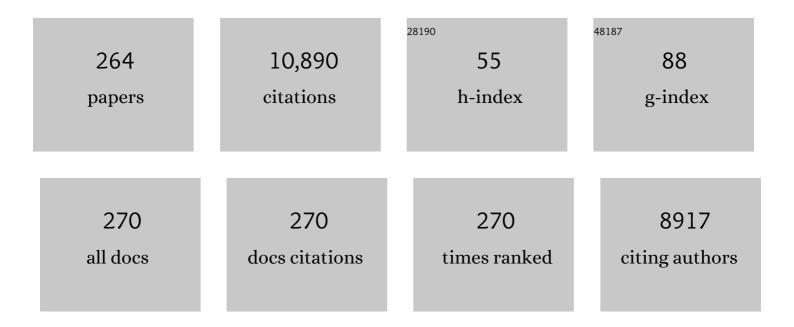
List of Publications by Year in descending order

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#	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.	1.6	1
2	Global biodiversity patterns of marine forests of brown macroalgae. Global Ecology and Biogeography, 2022, 31, 636-648.	2.7	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.	1.7	1
4	Major Expansion of Marine Forests in a Warmer Arctic. Frontiers in Marine Science, 2022, 9, .	1.2	16
5	Ocean currents shape the genetic structure of a kelp in southwestern Africa. Journal of Biogeography, 2022, 49, 822-835.	1.4	9
6	eDNA metabarcoding for diet analyses of green sea turtles (Chelonia mydas). Marine Biology, 2022, 169, 1.	0.7	14
7	The genus Cystoseira s.l. (Ochrophyta, Fucales, Sargassaceae) in the Black Sea: morphological variability and molecular taxonomy of Congolaria barbata and endemic Ericaria crinita f. bosphorica comb. nov.<:/em>:<:/strong>:. Phytotaxa. 2021. 480. 1-21.	0.1	15
8	Bottom Trawling Threatens Future Climate Refugia of Rhodoliths Globally. Frontiers in Marine Science, 2021, 7, .	1.2	27
9	Phylogeographic Analysis Suggests a Recent Population Bottleneck in the Rare Red Sea Tridacna squamosina. Frontiers in Marine Science, 2021, 8, .	1.2	Ο
10	Mates Matter: Gametophyte Kinship Recognition and Inbreeding in the Giant Kelp, <i>Macrocystispyrifera</i> (Laminariales, Phaeophyceae). Journal of Phycology, 2021, 57, 711-725.	1.0	16
11	Genomes Vary in Size and Spatial Patterns Within Chimeric Blades of Porphyra spp Frontiers in Marine Science, 2021, 8, .	1.2	5
12	New Records of Fish Species from the Coast of Luanda, Angola. Thalassas, 2021, 37, 803-811.	0.1	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.	1.5	12
14	Charting a course for genetic diversity in the UN Decade of Ocean Science. Evolutionary Applications, 2021, 14, 1497-1518.	1.5	19
15	Spatiotemporal patterns of phenology of the alien Phaeophyceae Sargassum muticum on the Atlantic coast of Morocco. Scientia Marina, 2021, 85, 103-111.	0.3	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.	1.4	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 ¹ . Journal of Phycology, 2021, 57, 1681-1698.	1.0	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.	1.5	10

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19	Predicted regime shift in the seagrass ecosystem of the Gulf of Arguin driven by climate change. Global Ecology and Conservation, 2021, 32, e01890.	1.0	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.	1.5	2
21	Microbial Surface Biofilm Responds to the Growth-Reproduction-Senescence Cycle of the Dominant Coral Reef Macroalgae Sargassum spp Life, 2021, 11, 1199.	1.1	0
22	Potential Biodiversity Connectivity in the Network of Marine Protected Areas in Western Africa. Frontiers in Marine Science, 2021, 8, .	1.2	9
23	Warming Threatens to Propel the Expansion of the Exotic Seagrass Halophila stipulacea. Frontiers in Marine Science, 2021, 8, .	1.2	13
24	Environmental drivers of rhodolith beds and epiphytes community along the South Western Atlantic coast. Marine Environmental Research, 2020, 154, 104827.	1.1	38
25	How experimental physiology and ecological niche modelling can inform the management of marine bioinvasions?. Science of the Total Environment, 2020, 700, 134692.	3.9	10
26	Brazil oil spill response: Protect rhodolith beds. Science, 2020, 367, 156-156.	6.0	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.	1.2	12
28	Linking Ecology to Genetics to Better Understand Adaptation and Evolution: A Review in Marine Macrophytes. Frontiers in Marine Science, 2020, 7, .	1.2	14
29	High Coral Bycatch in Bottom-Set Gillnet Coastal Fisheries Reveals Rich Coral Habitats in Southern Portugal. Frontiers in Marine Science, 2020, 7, .	1.2	14
30	Phylogeny and Evolution of the Brown Algae. Critical Reviews in Plant Sciences, 2020, 39, 281-321.	2.7	82
31	Spatial patterns of microbial communities across surface waters of the Great Barrier Reef. Communications Biology, 2020, 3, 442.	2.0	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.	1.6	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.	0.9	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.	2.7	3
35	Thermal traits for reproduction and recruitment differ between Arctic and Atlantic kelp Laminaria digitata. PLoS ONE, 2020, 15, e0235388.	1.1	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.	0.9	7

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37	Golden carbon of Sargassum forests revealed as an opportunity for climate change mitigation. Science of the Total Environment, 2020, 729, 138745.	3.9	68
38	A fine-tuned global distribution dataset of marine forests. Scientific Data, 2020, 7, 119.	2.4	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, .	1.2	8
40	Microbiome dynamics in the tissue and mucus of acroporid corals differ in relation to host and environmental parameters. PeerJ, 2020, 8, e9644.	0.9	30
41	Genetic Affinities and Biogeography of Putative Levantine-Endemic Seaweed Treptacantha rayssiae (Ramon) M.Mulas, J.Neiva & Ādsrael, comb. nov. (Phaeophyceae). Cryptogamie, Algologie, 2020, 41, .	0.3	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.	0.9	32
43	Sex-dependent and -independent transcriptional changes during haploid phase gametogenesis in the sugar kelp Saccharina latissima. PLoS ONE, 2019, 14, e0219723.	1.1	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.	0.8	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.	1.9	30
46	Toward a Coordinated Global Observing System for Seagrasses and Marine Macroalgae. Frontiers in Marine Science, 2019, 6, .	1.2	123
47	Open Coast Seagrass Restoration. Can We Do It? Large Scale Seagrass Transplants. Frontiers in Marine Science, 2019, 6, .	1.2	50
48	Increased evolutionary rates and conserved transcriptional response following allopolyploidization in brown algae. Evolution; International Journal of Organic Evolution, 2019, 73, 59-72.	1.1	11
49	Gene pool and connectivity patterns of <i>Pinna nobilis</i> in the Balearic Islands (Spain, Western) Tj ETQq1 Marine and Freshwater Ecosystems, 2019, 29, 175-188.	1 0.784314 ı 0.9	gBT /Overloci 9
50	Canopy microclimate modification in central and marginal populations of a marine macroalga. Marine Biodiversity, 2019, 49, 415-424.	0.3	23
51	Sexual reproduction vs. clonal propagation in the recovery of a seagrass meadow after an extreme weather event. Scientia Marina, 2019, 83, 357.	0.3	15
52	Genetic diversity increases with depth in red gorgonian populations of the Mediterranean Sea and the Atlantic Ocean. PeerJ, 2019, 7, e6794.	0.9	10
53	Genetic and oceanographic tools reveal high population connectivity and diversity in the endangered pen shell Pinna nobilis. Scientific Reports, 2018, 8, 4770.	1.6	31
54	Glacial vicariance drives phylogeographic diversification in the amphi-boreal kelp Saccharina latissima. Scientific Reports, 2018, 8, 1112.	1.6	61

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55	Bioâ€ORACLE v2.0: Extending marine data layers for bioclimatic modelling. Global Ecology and Biogeography, 2018, 27, 277-284.	2.7	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.	0.4	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.	0.9	5
58	Predicted extinction of unique genetic diversity in marine forests of Cystoseira spp Marine Environmental Research, 2018, 138, 119-128.	1.1	43
59	Projected climate changes threaten ancient refugia of kelp forests in the North Atlantic. Global Change Biology, 2018, 24, e55-e66.	4.2	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, .	1.2	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.	0.4	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.	1.1	57
63	Individual-based genetic analyses support asexual hydrochory dispersal in Zostera noltei. PLoS ONE, 2018, 13, e0199275.	1.1	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.	1.1	28
65	Harnessing positive species interactions as a tool against climate-driven loss of coastal biodiversity. PLoS Biology, 2018, 16, e2006852.	2.6	91
66	The introduction of <i>Sargassum muticum</i> modifies epifaunal patterns in a Moroccan seagrass meadow. Marine Ecology, 2018, 39, e12507.	0.4	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.	1.6	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, .	1.2	149
69	Dramatic loss of seagrass habitat under projected climate change in the Mediterranean Sea. Global Change Biology, 2018, 24, 4919-4928.	4.2	140
70	Kelps' Long-Distance Dispersal: Role of Ecological/Oceanographic Processes and Implications to Marine Forest Conservation. Diversity, 2018, 10, 11.	0.7	34
71	Postglacial expansion of the Arctic keystone copepod Calanus glacialis. Marine Biodiversity, 2018, 48, 1027-1035.	0.3	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.	1.4	44

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73	Re-assessing the origins of the invasive mussel Mytilus galloprovincialis in southern Africa. Marine and Freshwater Research, 2018, 69, 607.	0.7	22
74	Polyploid lineages in the genus Porphyra. Scientific Reports, 2018, 8, 8696.	1.6	21
75	Seaweed Loads Cause Stronger Bacterial Community Shifts in Coastal Lagoon Sediments Than Nutrient Loads. Frontiers in Microbiology, 2018, 9, 3283.	1.5	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.	0.9	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.	0.3	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.	2.0	42
80	Population dynamics of temperate kelp forests near their low-latitude limit. Aquatic Botany, 2017, 139, 8-18.	0.8	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.	1.6	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, .	0.6	43
84	Regional Genetic Structure in the Aquatic Macrophyte Ruppia cirrhosa Suggests Dispersal by Waterbirds. Estuaries and Coasts, 2017, 40, 1705-1716.	1.0	16
85	Accounting for uncertainty in predictions of a marine species: Integrating population genetics to verify past distributions. Ecological Modelling, 2017, 359, 229-239.	1.2	19
86	Larval development and allometric growth of the blackâ€faced blenny <i>Tripterygion delaisi</i> . Journal of Fish Biology, 2017, 90, 2239-2254.	0.7	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.	1.6	87
88	Development and characterization of twelve microsatellite markers for Porphyra linearis Greville. Genetica, 2017, 145, 127-130.	0.5	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.	2.0	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.	1.6	9

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91	Palaeoclimatic conditions in the Mediterranean explain genetic diversity of Posidonia oceanica seagrass meadows. Scientific Reports, 2017, 7, 2732.	1.6	29
92	First record of Ruppia maritima in West Africa supported by morphological description and phylogenetic classification. Botanica Marina, 2017, 60, .	0.6	6
93	High genetic differentiation of red gorgonian populations from the Atlantic Ocean and the Mediterranean Sea. Marine Biology Research, 2017, 13, 854-861.	0.3	2
94	First description of seagrass distribution and abundance in São Tomé and PrÃncipe. Aquatic Botany, 2017, 142, 48-52.	0.8	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.	1.5	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.	0.5	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, .	0.2	1
99	Species Specificity of Bacteria Associated to the Brown Seaweeds Lobophora (Dictyotales,) Tj ETQq1 1 0.784314 i Frontiers in Microbiology, 2016, 7, 316.	rgBT /Ove 1.5	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.	1.5	72
101	Genetic Diversity and Local Connectivity in the Mediterranean Red Gorgonian Coral after Mass Mortality Events. PLoS ONE, 2016, 11, e0150590.	1.1	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.	1.4	92
103	Temporal windows of reproductive opportunity reinforce species barriers in a marine broadcast spawning assemblage. Scientific Reports, 2016, 6, 29198.	1.6	17
104	Overlooked habitat of a vulnerable gorgonian revealed in the Mediterranean and Eastern Atlantic by ecological niche modelling. Scientific Reports, 2016, 6, 36460.	1.6	35
105	Pan-Arctic population of the keystone copepod Calanus glacialis. Polar Biology, 2016, 39, 2311-2318.	0.5	16
106	Deep reefs are climatic refugia for genetic diversity of marine forests. Journal of Biogeography, 2016, 43, 833-844.	1.4	84
107	Novel polymorphic microsatellite loci for a new target species, the sea cucumber Holothuria mammata. Biochemical Systematics and Ecology, 2016, 66, 109-113.	0.6	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.	1.9	42

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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.	0.7	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.	1.0	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.	0.9	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.	1.0	51
114	Do hatchery-reared sea urchins pose a threat to genetic diversity in wild populations?. Heredity, 2016, 116, 378-383.	1.2	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.	1.1	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.	0.9	37
117	Characterization of 12 polymorphic microsatellite markers in the sugar kelp Saccharina latissima. Journal of Applied Phycology, 2016, 28, 3071-3074.	1.5	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.	1.1	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.	1.7	15
121	Comparison of small remotely operated vehicles and diver-operated video of circalittoral benthos. Hydrobiologia, 2016, 766, 247-260.	1.0	30
122	A Well-Kept Treasure at Depth: Precious Red Coral Rediscovered in Atlantic Deep Coral Gardens (SW) Tj ETQq0 0	0 ₁₉ BT /C	verlock 10 Tf
123	High Interannual Variability in Connectivity and Genetic Pool of a Temperate Clingfish Matches Oceanographic Transport Predictions. PLoS ONE, 2016, 11, e0165881.	1.1	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.	0.9	10
125	Tradeâ€offs between lifeâ€history traits at rangeâ€edge and central locations. Journal of Phycology, 2015, 51, 808-818.	1.0	16
126	Seascape drivers of <i><scp>M</scp>acrocystis pyrifera</i> population genetic structure in the	2.0	55

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

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127	Oceanographic Conditions Limit the Spread of a Marine Invader along Southern African Shores. PLoS ONE, 2015, 10, e0128124.	1.1	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.	1.0	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.	0.4	16
130	Genetic diversity and biogeographical patterns of Caulerpa prolifera across the Mediterranean and Mediterranean/Atlantic transition zone. Marine Biology, 2015, 162, 557-569.	0.7	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.	1.7	33
132	A transcriptome resource for Antarctic krill (Euphausia superba Dana) exposed to short-term stress. Marine Genomics, 2015, 23, 45-47.	0.4	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.	1.3	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.	0.9	7
135	Metatranscriptomes reveal functional variation in diatom communities from the Antarctic Peninsula. ISME Journal, 2015, 9, 2275-2289.	4.4	55
136	A transcriptome resource for the copepod Calanus glacialis across a range of culture temperatures. Marine Genomics, 2015, 23, 27-29.	0.4	10
137	Polymorphic microsatellite markers in the brown seaweed Fucus vesiculosus. BMC Research Notes, 2015, 8, 73.	0.6	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.	1.1	2
140	European seaweeds under pressure: Consequences for communities and ecosystem functioning. Journal of Sea Research, 2015, 98, 91-108.	0.6	155
141	Response of kelps from different latitudes to consecutive heat shock. Journal of Experimental Marine Biology and Ecology, 2015, 463, 57-62.	0.7	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.	0.4	24
143	Highly polymorphic microsatellite markers for the Mediterranean endemic fan mussel Pinna nobilis. Mediterranean Marine Science, 2015, 16, 31.	0.6	13
144	Genes Left Behind: Climate Change Threatens Cryptic Genetic Diversity in the Canopy-Forming Seaweed Bifurcaria bifurcata. PLoS ONE, 2015, 10, e0131530.	1.1	52

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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.	1.1	24
146	Microsatellite markers for the Arctic copepod Calanus glacialis and cross-amplification with C. finmarchicus. Conservation Genetics Resources, 2014, 6, 1003-1005.	0.4	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.	0.9	12
148	Reproductive strategies and isolationâ€byâ€demography in a marine clonal plant along an eutrophication gradient. Molecular Ecology, 2014, 23, 5698-5711.	2.0	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.	0.8	34
150	Genetic Divergence for the Amphibian Pleurodeles waltl in Southwest Portugal: Dispersal Barriers Shaping Geographic Patterns. Journal of Herpetology, 2014, 48, 38.	0.2	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.	1.4	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.	2.0	77
153	Reprint of "Seagrasses in Portugal: A most endangered marine habitat― Aquatic Botany, 2014, 115, 3-13.	0.8	10
154	Wider sampling reveals a nonâ€sister relationship for geographically contiguous lineages of a marine mussel. Ecology and Evolution, 2014, 4, 2070-2081.	0.8	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.	0.7	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.	0.4	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.	1.0	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.	0.6	11
159	Microsatellite markers developed through pyrosequencing allow clonal discrimination in the invasive alga Caulerpa taxifolia. Conservation Genetics Resources, 2013, 5, 667-669.	0.4	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.	0.4	2
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