Jonathan P A Gardner

List of Publications by Year
in descending order

138 papers	3,096 citations	159358 30 h-index	223531 46 g-index
140 all docs	140 docs citations	140 times ranked	3376 citing authors

#	Article	IF	Citations
1	Mining of deep-sea seafloor massive sulfides: A review of the deposits, their benthic communities, impacts from mining, regulatory frameworks and management strategies. Ocean and Coastal Management, 2013, 84, 54-67.	2.0	248
2	Title is missing!. Biological Invasions, 2000, 2, 75-79.	1.2	115
3	Population genetic subdivision in the New Zealand greenshell mussel (Perna canaliculus) inferred from single-strand conformation polymorphism analysis of mitochondrial DNA. Molecular Ecology, 2002, 11, 1617-1628.	2.0	77
4	Internal borders for managing invasive marine species. Journal of Applied Ecology, 2009, 46, 46-54.	1.9	68
5	Human dietary exposure to heavy metals via the consumption of greenshell mussels (Perna) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 5 Environment, 2009, 407, 4348-4355.	3.9	66
6	Genetic diversity of Southern hemisphere blue mussels (Bivalvia: Mytilidae) and the identification of non-indigenous taxa. Biological Journal of the Linnean Society, 2010, 101, 898-909.	0.7	66
7	Shoreline Changes and Sediment Redistribution at Palmyra Atoll (Equatorial Pacific Ocean): 1874â€“Present. Journal of Coastal Research, 2009, 253, 711-722.	0.1	65
8	Reliance of mobile species on sensitive habitats: a case study of manta rays (Manta alfredi) and lagoons. Marine Biology, 2014, 161, 1987-1998.	0.7	65
9	Seafloor massive sulfide deposits support unique megafaunal assemblages: Implications for seabed mining and conservation. Marine Environmental Research, 2016, 115, 78-88.	1.1	65
10	Historical and size-dependent genetic variation in hybrid mussel populations. Heredity, 1988, 61, 93-105.	1.2	64
11	A molecular phylogeny of the marine mussel genus Perna (Bivalvia: Mytilidae) based on nuclear (ITS1&2) and mitochondrial (COI) DNA sequences. Molecular Phylogenetics and Evolution, 2007, 44, 685-698.	1.2	63
12	The effects of coastal and estuarine conditions on the physiology and survivorship of the mussels Mytilus edulis, M. trossulus and their hybrids. Journal of Experimental Marine Biology and Ecology, 2001, 265, 119-140.	0.7	56
13	Local endemicity and high diversity characterise high-latitude coralâ€“Symbiodinium partnerships. Coral Reefs, 2010, 29, 989-1003.	0.9	55
14	Megabenthic assemblage structure on three New Zealand seamounts: implications for seafloor massive sulfide mining. Marine Ecology - Progress Series, 2015, 523, 1-14.	0.9	55
15	Effect of vessel voyage speed on survival of biofouling organisms: implications for translocation of non-indigenous marine species. Biofouling, 2010, 26, 1-13.	0.8	54
16	Native and invasive taxa on the Pacific coast of South America: Impacts on aquaculture, traceability and biodiversity of blue mussels (Mytilus spp.). Evolutionary Applications, 2018, 11, 298-311.	1.5	54
17	Invasive blue mussels threaten regional scale genetic diversity in mainland and remote offshore locations: the need for baseline data and enhanced protection in the Southern Ocean. Global Change Biology, 2016, 22, 3182-3195.	4.2	49
18	Efficacy of acetic acid treatments in the management of marine biofouling. Aquaculture, 2007, 262, 319-332.	1.7	45

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19	Population genetic variation in the New Zealand greenshell mussel: locus-dependent conflicting signals of weak structure and high gene flow balanced against pronounced structure and high self-recruitment. Marine Biology, 2013, 160, 931-949.	0.7	44
20	Quantifying and addressing the prevalence and bias of study designs in the environmental and social sciences. Nature Communications, 2020, 11, 6377.	5.8	44
21	Applying Fishers' Ecological Knowledge to Construct Past and Future Lobster Stocks in the Juan FernÃ¡ndez Archipelago, Chile. PLoS ONE, 2010, 5, e13670.	1.1	43
22	Marine reserves increase the abundance and size of blue cod and rock lobster. Marine Ecology - Progress Series, 2008, 366, 147-158.	0.9	43
23	A primer for use of genetic tools in selecting and testing the suitability of set-aside sites protected from deep-sea seafloor massive sulfide mining activities. Ocean and Coastal Management, 2016, 122, 37-48.	2.0	42
24	The Effect of Diet on the Energy Budget of the Brown Sea Cucumber, Stichopus mollis (Hutton). Journal of the World Aquaculture Society, 2009, 40, 157-170.	1.2	41
25	Mytilus Galloprovincialis (Lmk) (Bivalvia, Mollusca): The Taxonomic Status of the Mediterranean Mussel. Ophelia, 1992, 35, 219-243.	0.3	40
26	The effect of vessel speed on the survivorship of biofouling organisms at different hull locations. Biofouling, 2010, 26, 539-553.	0.8	40
27	Effects of seston variability on the clearance rate and absorption efficiency of the mussels Aulacomya maoriana, Mytilus galloprovincialis and Perna canaliculus from New Zealand. Journal of Experimental Marine Biology and Ecology, 2002, 268, 83-101.	0.7	38
28	Macrobenthicâ€“mud relations strengthen the foundation for benthic index development: A case study from shallow, temperate New Zealand estuaries. Ecological Indicators, 2015, 58, 161-174.	2.6	37
29	Small spatial scale population genetic structure in two limpet species endemic to the Kermadec Islands, New Zealand. Marine Ecology - Progress Series, 2007, 349, 159-170.	0.9	35
30	A historical perspective of the genus Mytilus (Bivalvia: Mollusca) in New Zealand: multivariate morphometric analyses of fossil, midden and contemporary blue mussels. Biological Journal of the Linnean Society, 2004, 82, 329-344.	0.7	33
31	Successful eradication of a nonâ€indigenous marine bivalve from a subtidal softâ€sediment environment. Journal of Applied Ecology, 2011, 48, 424-431.	1.9	32
32	A comparison of genetic diversity between cultured and wild populations, and a test for genetic introgression in the New Zealand greenshell mussel Perna canaliculus (Gmelin 1791). Aquaculture, 2003, 219, 193-220.	1.7	30
33	Does differential particulate food supply explain the presence of mussels in Wellington Harbour (New Zealand) and their absence on neighbouring Cook Strait shores?. Estuarine, Coastal and Shelf Science, 2007, 72, 223-234.	0.9	30
34	Population genetic structure and connectivity of deepâ€sea stony corals (Order Scleractinia) in the New Zealand region: Implications for the conservation and management of vulnerable marine ecosystems. Evolutionary Applications, 2017, 10, 1040-1054.	1.5	30
35	Development and evaluation of microsatellite markers for identification of individual Greenshellâ„¢ mussels (Perna canaliculus) in a selective breeding programme. Aquaculture, 2008, 274, 41-48.	1.7	28
36	An RFLP assay to determine if Mytilus galloprovincialis Lmk. (Mytilidae; Bivalvia) is of Northern or Southern hemisphere origin. Molecular Ecology Resources, 2010, 10, 573-575.	2.2	28

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37	Interlineage Mytilus galloprovincialis Lmk. 1819 hybridization yields inconsistent genetic outcomes in the Southern hemisphere. Biological Invasions, 2013, 15, 1493-1506.	1.2	28
38	Bioinvasion threatens the genetic integrity of native diversity and a natural hybrid zone: smooth-shelled blue mussels (Mytilusspp.) in the Strait of Magellan. Biological Journal of the Linnean Society, 2016, 117, 574-585.	0.7	28
39	Influence of genotype and geography on shell shape and morphometric trait variation among North Atlantic blue mussel (Mytilus spp.) populations. Biological Journal of the Linnean Society, 0, 96, 875-897.	0.7	27
40	Polymorphism and vestigiality: comparative anatomy and morphology of bryozoan avicularia. Zoomorphology, 2010, 129, 195-211.	0.4	27
41	Optimising a widely-used coastal health index through quantitative ecological group classifications and associated thresholds. Ecological Indicators, 2016, 69, 595-605.	2.6	27
42	Lobsters as keystone: Only in unfished ecosystems?. Ecological Modelling, 2014, 275, 48-72.	1.2	26
43	Polymorphism and variation in modular animals: morphometric and density analyses of bryozoan avicularia. Marine Ecology - Progress Series, 2010, 399, 117-130.	0.9	26
44	Contrasting patterns of mussel abundance at neighbouring sites: does recruitment limitation explain the absence of mussels on Cook Strait (New Zealand) shores?. Journal of Experimental Marine Biology and Ecology, 2004, 312, 285-298.	0.7	24
45	Size-dependent, spatial and temporal genetic variation at a leucine aminopeptidase (LAP) locus among blue mussel (Mytilus galloprovincialis) populations along a salinity gradient. Marine Biology, 1998, 132, 275-281.	0.7	23
46	Variation in scope for growth: a test of food limitation among intertidal mussels. Hydrobiologia, 2007, 586, 373-392.	1.0	23
47	Permanent Genetic Resources added to Molecular Ecology Resources Database 1 December 2011 â€“ 31 January 2012. Molecular Ecology Resources, 2012, 12, 570-572.	2.2	23
48	Seascape genetics of the New Zealand greenshell mussel: sea surface temperature explains macrogeographic scale genetic variation. Marine Ecology - Progress Series, 2013, 477, 107-121.	0.9	23
49	Long-term effects of a toxic algal bloom on subtidal soft-sediment macroinvertebrate communities in Wellington Harbour, New Zealand. Estuarine, Coastal and Shelf Science, 2006, 67, 589-604.	0.9	22
50	Restriction of sponges to an atoll lagoon as a result of reduced environmental quality. Marine Pollution Bulletin, 2013, 66, 209-220.	2.3	22
51	Phylogeography and population genetics of Schizothorax oâ€™connori: strong subdivision in the Yarlung Tsangpo River inferred from mtDNA and microsatellite markers. Scientific Reports, 2016, 6, 29821.	1.6	22
52	Factors affecting survivorship of defouled communities and the effect of fragmentation on establishment success. Journal of Experimental Marine Biology and Ecology, 2011, 396, 233-243.	0.7	21
53	Molecular identification and expression of the Foxl2 gene during gonadal sex differentiation in northern snakehead Channa argus. Fish Physiology and Biochemistry, 2015, 41, 1419-1433.	0.9	21
54	Cryptic diversity in smooth-shelled mussels on Southern Ocean islands: connectivity, hybridisation and a marine invasion. Frontiers in Zoology, 2019, 16, 32.	0.9	21

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55	Connectivity, small islands and large distances: the Cellana strigilis limpet complex in the Southern Ocean. Molecular Ecology, 2011, 20, 3399-3413.	2.0	20
56	Developmental stability is not disrupted by extensive hybridization and introgression among populations of the marine bivalve molluscs Mytilus edulis (L.) and M. galloprovincialis (Lmk.) from south-west England. Biological Journal of the Linnean Society, 1995, 54, 71-86.	0.7	18
57	Molecular Characterization and Expression Analyses of the Complement Component C8Î±, C8Î² and C9 Genes in Yellow Catfish (Pelteobagrus fulvidraco) after the Aeromonas hydrophila Challenge. International Journal of Molecular Sciences, 2016, 17, 345.	1.8	18
58	Historical divergences associated with intermittent land bridges overshadow isolation by larval dispersal in coâ€distributed species of Tridacna giant clams. Journal of Biogeography, 2018, 45, 848-858.	1.4	18
59	Species-specific genetic variation in response to deep-sea environmental variation amongst Vulnerable Marine Ecosystem indicator taxa. Scientific Reports, 2020, 10, 2844.	1.6	18
60	Naturally low seston concentration and the net energy balance of the greenshell mussel (Perna) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 54 Research, 2001, 35, 457-468.	0.8	17
61	Absence of population genetic differentiation in the New Zealand greenshell mussel Perna canaliculus (Gmelin 1791) as assessed by allozyme variation. Journal of Experimental Marine Biology and Ecology, 2001, 258, 173-194.	0.7	17
62	Coral larvae change their settlement preference for crustose coralline algae dependent on availability of bare space. Coral Reefs, 2018, 37, 397-407.	0.9	17
63	Correlation between pelagic larval duration and realised dispersal: long-distance genetic connectivity between northern New Zealand and the Kermadec Islands archipelago. Marine Biology, 2014, 161, 297-312.	0.7	16
64	A baseline biological survey of the proposed Taputeranga Marine Reserve (Wellington, New Zealand): spatial and temporal variability along a natural environmental gradient. Aquatic Conservation: Marine and Freshwater Ecosystems, 2009, 19, 237-248.	0.9	15
65	Ocean currents predict fine-scale genetic structure and source-sink dynamics in a marine invertebrate coastal fishery. ICES Journal of Marine Science, 2019, 76, 1007-1018.	1.2	15
66	Laboratory experiments on the effects of variable suspended sediment concentrations on the ecophysiology of the porcelain crab Petrolisthes elongatus (Milne Edwards, 1837). Journal of Experimental Marine Biology and Ecology, 2007, 344, 181-192.	0.7	14
67	Near-surface mixing and pronounced deep-water stratification in a compartmentalised, human-disturbed atoll lagoon system. Coral Reefs, 2011, 30, 271-282.	0.9	14
68	Spatial patterns and regional affinities of coral communities at the Kermadec Islands Marine Reserve, New Zealandâ€"a marginalÂ high-latitude site. Marine Ecology - Progress Series, 2010, 400, 101-113.	0.9	14
69	Emerging patterns of genetic variation in the New Zealand endemic scallop Pecten novaezelandiae. Molecular Ecology, 2015, 24, 5379-5393.	2.0	13
70	Identifying environmental factors associated with the genetic structure of the New Zealand scallop: linking seascape genetics and ecophysiological tolerance. ICES Journal of Marine Science, 2016, 73, 1925-1934.	1.2	13
71	Genetic Diversity of and Differentiation among Five Populations of Blunt Snout Bream (Megalobrama) Tj ETQq1 1 0.784314 rgBT /Ov 2014, 9, e108967.	1.1	13
72	The Mytilus edulis species complex in Southwest England: Multi-locus heterozygosity, background genotype and a fitness correlate. Biochemical Systematics and Ecology, 1994, 22, 1-11.	0.6	12

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73	High levels of shared allozyme polymorphism among strongly differentiated congeneric clams of the genus Astarte (Bivalvia: Mollusca). Heredity, 1999, 82, 89-99.	1.2	12
74	Evaluation and optimisation of underwater visual census monitoring for quantifying change in rocky-reef fish abundance. Biological Conservation, 2015, 186, 326-336.	1.9	12
75	Use of high-resolution acoustic cameras to study reef shark behavioral ecology. Journal of Experimental Marine Biology and Ecology, 2016, 482, 128-133.	0.7	12
76	The use of spatially explicit genetic variation data from four deep-sea sponges to inform the protection of Vulnerable Marine Ecosystems. Scientific Reports, 2019, 9, 5482.	1.6	12
77	Dermal denticle assemblages in coral reef sediments correlate with conventional shark surveys. Methods in Ecology and Evolution, 2020, 11, 362-375.	2.2	12
78	Trioecy in the Marine Mussel Semimytilus algosus (Mollusca, Bivalvia): Stable Sex Ratios Across 22 Degrees of a Latitudinal Gradient. Frontiers in Marine Science, 2020, 7, .	1.2	12
79	Mitochondrial DNA and allozyme covariation in a hybrid mussel population. Journal of Experimental Marine Biology and Ecology, 1991, 149, 45-54.	0.7	11
80	Functional innovation through vestigialization in a modular marine invertebrate. Biological Journal of the Linnean Society, 2011, 104, 63-74.	0.7	11
81	Limitations in the Use of Archived Vent Mussel Samples to Assess Genetic Connectivity Among Seafloor Massive Sulfide Deposits: A Case Study with Implications for Environmental Management. Frontiers in Marine Science, 2015, 2, .	1.2	10
82	Combining genotypic and phenotypic variation in a geospatial framework to identify sources of mussels in northern New Zealand. Scientific Reports, 2021, 11, 8196.	1.6	10
83	Development of sensitive and specific molecular tools for the efficient detection and discrimination of potentially invasive mussel species of the genus Perna. Management of Biological Invasions, 2013, 4, 155-165.	0.5	10
84	Benthic community structure and water column characteristics at two sites in the Kermadec Islands Marine Reserve, New Zealand. New Zealand Journal of Marine and Freshwater Research, 2006, 40, 179-194.	0.8	9
85	Application of the littoral cell concept to managing a protected atoll: Palmyra Atoll National Wildlife Refuge. Ocean and Coastal Management, 2009, 52, 628-635.	2.0	9
86	A morphometric approach supporting genetic results in the taxonomy of the New Zealand limpets of the Cellana strigilis complex (Mollusca : Patellogastropoda : Nacellidae). Invertebrate Systematics, 2012, 26, 193.	0.5	9
87	Combined evidence indicates that Perna indica Kuriakose and Nair 1976 is Perna perna (Linnaeus, 1758) from the Oman region introduced into southern India more than 100Âyears ago. Biological Invasions, 2016, 18, 1375-1390.	1.2	9
88	Phylogeography of the threatened tetraploid fish, Schizothorax waltoni, in the Yarlung Tsangpo River on the southern Qinghai-Tibet Plateau: implications for conservation. Scientific Reports, 2019, 9, 2704.	1.6	9
89	Combined threats to native smooth-shelled mussels (genus Mytilus) in Australia: bioinvasions and hybridization. Zoological Journal of the Linnean Society, 2022, 194, 1194-1211.	1.0	9
90	Comparison of methodologies to quantify the effects of age and area of marine reserves on the density and size of targeted species. Aquatic Biology, 2012, 14, 185-200.	0.5	9

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91	The atmospheric lead record preserved in lagoon sediments at a remote equatorial Pacific location: Palmyra Atoll, northern Line Islands. Marine Pollution Bulletin, 2011, 62, 251-257.	2.3	8
92	Comparisons among survey methodologies to test for abundance and size of a highly targeted fish species. Journal of Fish Biology, 2013, 82, 242-262.	0.7	8
93	Massive differential site-specific and species-specific responses of temperate reef fishes to marine reserve protection. Global Ecology and Conservation, 2014, 1, 13-26.	1.0	8
94	First evidence of establishment of the rayed pearl oyster, Pinctada imbricata radiata (Leach, 1814), in the eastern Adriatic Sea. Marine Pollution Bulletin, 2017, 125, 556-560.	2.3	8
95	Changes in subtidal macroinvertebrate community structure in Wellington Harbour (New Zealand) following a largeâ€scale natural dieâ€off. New Zealand Journal of Marine and Freshwater Research, 2006, 40, 29-42.	0.8	7
96	The Kapiti Marine Reserve (New Zealand): spatial and temporal comparisons of multi-species responses after 8 years of protection. New Zealand Journal of Marine and Freshwater Research, 2012, 46, 71-89.	0.8	7
97	The genetic architecture of hybridisation between two lineages of greenshell mussels. Heredity, 2015, 114, 344-355.	1.2	7
98	Population Structure and Genetic Connectivity of Squat Lobsters (Munida Leach, 1820) Associated With Vulnerable Marine Ecosystems in the Southwest Pacific Ocean. Frontiers in Marine Science, 2020, 6, .	1.2	7
99	Analyses of DNA obtained from shells and brine-preserved meat of the giant clam Tridacna maxima from the central Pacific Ocean. Marine Ecology - Progress Series, 2012, 453, 297-301.	0.9	7
100	Recovery of a subtidal soft-sediment macroinvertebrate assemblage following experimentally induced effects of a harmful algal bloom. Marine Ecology - Progress Series, 2006, 326, 85-98.	0.9	7
101	Molecular genetic differentiation of native populations of Mediterranean blue mussels,Mytilus galloprovincialisLamarck, 1819, and the relationship with environmental variables. , 2022, 89, 755-784.		7
102	Growth and production of aLittorina littorea(L.) Population in the Bay of Fundy. Ophelia, 1987, 27, 181-195.	0.3	6
103	Geographic distribution and molecular identification of a metapopulation of blue mussels (genus) Tj ETQq1 1 0.784314 rgBT /Overloc	0.4	6
104	Host tolerance, not symbiont tolerance, determines the distribution of coral species in relation to their environment at a Central Pacific atoll. Coral Reefs, 2012, 31, 389-398.	0.9	6
105	Distant Storms as Drivers of Environmental Change at Pacific Atolls. PLoS ONE, 2014, 9, e87971.	1.1	6
106	Marine reserve establishment and on-going management costs: A case study from New Zealand. Marine Policy, 2015, 60, 216-224.	1.5	6
107	Modelling the effect of wave forces on subtidal macroalgae: A spatial evaluation of predicted disturbance for two habitat-forming species. Ecological Modelling, 2015, 313, 149-161.	1.2	6
108	Reproductive traits of the threatened freshwater musselSolenaia oleivora(Bivalvia: Unionidae) from the middle Yangtze River. Journal of Molluscan Studies, 2015, 81, 522-526.	0.4	6

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109	Using Genomics to Link Populations of an Invasive Species to Its Potential Sources. Frontiers in Ecology and Evolution, 2021, 9, .	1.1	6
110	A test for overdominance at the phosphoglucomutase-2 locus in Pacific oysters (Crassostrea gigas) from New Zealand. Aquaculture, 2005, 244, 29-39.	1.7	5
111	Development of twenty-one polymorphic tetranucleotide microsatellite loci for Schizothorax o'connori and their conservation application. Biochemical Systematics and Ecology, 2013, 51, 259-263.	0.6	5
112	Isolation and characterization of nineteen novel polymorphic microsatellite loci for the northern snakehead Channa argus. Conservation Genetics Resources, 2014, 6, 621-623.	0.4	5
113	Genetic diversity and population structure of the northern snakehead (Channa argus Channidae:) Tj ETQq1 1 0.784314 rgBT /Overloc 2018, 19, 467-480.	0.8	5
114	Morphometric variability of smooth-shelled blue mussels from the Pacific coast of South America. Biological Journal of the Linnean Society, 2018, 125, 194-209.	0.7	5
115	Blue mussels of the Mytilus edulis species complex from South America: The application of species delimitation models to DNA sequence variation. PLoS ONE, 2021, 16, e0256961.	1.1	5
116	Bottom-up control of temperate rocky intertidal community structure: evidence from a transplant experiment. Marine Ecology - Progress Series, 2013, 491, 137-151.	0.9	5
117	The complete mitochondrial genome of the deep-sea stony coralSolenosmilia variabilis(Scleractinia, Caryophylliidae) and its inter-individual variation. Mitochondrial DNA, 2016, 27, 1-2.	0.6	4
118	Development and characterisation of 12 microsatellite markers for the New Zealand endemic scallop Pecten novaezelandiae. Conservation Genetics Resources, 2014, 6, 327-328.	0.4	4
119	The complete mitochondrial genome of the deep-sea spongePoecillastra laminaris(Astrophorida,) Tj ETQq1 1 0.784314 rgBT /Overloc	0.6	4
120	Conservation management options and actions: Putative decline of coral cover at Palmyra Atoll, Northern Line Islands, as a case study. Marine Pollution Bulletin, 2014, 84, 182-190.	2.3	4
121	Three polymorphic mitochondrial DNA markers for Perna canaliculus. Animal Genetics, 2001, 32, 47-49.	0.6	3
122	Effect of storm drain discharge on the soft shore ecology of Porirua Inlet, New Zealand. New Zealand Journal of Marine and Freshwater Research, 2002, 36, 241-255.	0.8	3
123	Lobster fishery and marine reserve interactions in central New Zealand. Marine Policy, 2019, 105, 67-79.	1.5	3
124	Development and characterization of 20 polymorphic microsatellite loci in the deep sea squat lobster, Munida isos Ahyong and Poore, 2004 and cross-amplification in two congeneric species. Journal of Genetics, 2019, 98, 1.	0.4	3
125	An Indigenous-led Community Challenge to Fisheries Management in New Zealand: the Revival of Regional Scale Management Practices?. Pacific Conservation Biology, 2008, 14, 248.	0.5	2
126	Regional-scale genetic differentiation of the stony coral Desmophyllum dianthus in the southwest Pacific Ocean is consistent with regional-scale physico-chemical oceanography. Deep-Sea Research Part I: Oceanographic Research Papers, 2022, 183, 103739.	0.6	2

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127	A METHOD FOR THE INVESTIGATION OF THE SHELL STRUCTURE OF NEWLY SETTLED LIMPETS. Journal of Molluscan Studies, 1986, 52, 35-37.	0.4	1
128	No evidence for overdominance at the phosphoglucomutase-2 locus in Pacific oysters (Crassostrea) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	1.7	1
129	Development of polymorphic microsatellite markers for the pulmonate limpet Siphonaria australis. Conservation Genetics Resources, 2010, 2, 377-379.	0.4	1
130	Commentary on Palmyra atoll. Marine Pollution Bulletin, 2011, 62, 2876-2877.	2.3	1
131	Development and characterization of 20 polymorphic microsatellite loci for the Lhasa schizothoracin Schizothorax waltoni. Conservation Genetics Resources, 2014, 6, 413-415.	0.4	1
132	Isolation and characterization of twenty-one polymorphic microsatellite loci from Schizothorax oâ€™connori and cross-species amplification. Journal of Genetics, 2016, 93, 60-64.	0.4	1
133	Isolation and characterization of twenty-one polymorphic microsatellite loci from Schizothorax o'connori and cross-species amplification. Journal of Genetics, 2013, 92, e60-4.	0.4	1
134	Genetic structure and diversity of the Chilean flat oyster Ostrea chilensis (Bivalvia: Ostreidae) along its natural distribution from natural beds subject to different fishing histories. Genetics and Molecular Biology, 2022, 45, e20210214.	0.6	1
135	Characterisation of novel microsatellite markers for the surf clams Paphies subtriangulata and P. australis (Bivalvia: Mesodesmatidae). Conservation Genetics Resources, 2014, 6, 315-317.	0.4	0
136	Development and characterization of 32 SNP markers for the northern snakehead (Channa argus) using high resolution melting (HRM). Conservation Genetics Resources, 2017, 9, 631-634.	0.4	0
137	Development and characterization of ten highly polymorphic microsatellite markers for the demosponge Poecillastra laminaris (Sollas). Marine Biodiversity, 2018, 48, 1265-1267.	0.3	0
138	Inner shelf habitat surrounding the Kapiti Marine Reserve, New Zealand. , 2020, , 403-419.		0