## Sarah Samadi

## List of Publications by Year in descending order

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109137 161609 3,315 79 35 54 h-index citations g-index papers 82 82 82 4438 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Largeâ€scale species delimitation method for hyperdiverse groups. Molecular Ecology, 2012, 21, 2671-2691.	2.0	259
2	Species are hypotheses: avoid connectivity assessments based on pillars of sand. Molecular Ecology, 2015, 24, 525-544.	2.0	197
3	Use of RAD sequencing for delimiting species. Heredity, 2015, 114, 450-459.	1.2	163
4	Seamount endemism questioned by the geographic distribution and population genetic structure of marine invertebrates. Marine Biology, 2006, 149, 1463-1475.	0.7	162
5	Phase determination from direct sequencing of length-variable DNA regions. Molecular Ecology Notes, 2006, 6, 627-630.	1.7	123
6	Importance of Assessing Population Genetic Structure before Eradication of Invasive Species: Examples from Insular Norway Rat Populations. Conservation Biology, 2005, 19, 1509-1518.	2.4	112
7	Symbioses between deep-sea mussels (Mytilidae: Bathymodiolinae) and chemosynthetic bacteria: diversity, function and evolution. Comptes Rendus - Biologies, 2009, 332, 298-310.	0.1	98
8	Several deep-sea mussels and their associated symbionts are able to live both on wood and on whale falls. Proceedings of the Royal Society B: Biological Sciences, 2009, 276, 177-185.	1.2	81
9	Microsatellite and morphological analysis of population structure in the parthenogenetic freshwater snail Melanoides tuberculata: insights into the creation of clonal variability. Molecular Ecology, 1999, 8, 1141-1153.	2.0	79
10	New insights into diversity and evolution of deep-sea Mytilidae (Mollusca: Bivalvia). Molecular Phylogenetics and Evolution, 2010, 57, 71-83.	1.2	72
11	Permanent Genetic Resources added to Molecular Ecology Resources Database 1 April 2010 – 31 May 2010. Molecular Ecology Resources, 2010, 10, 1098-1105.	2.2	71
12	Starting to unravel the toxoglossan knot: Molecular phylogeny of the "turrids―(Neogastropoda:) Tj ETQq0 (	0 0 rgBT /(	Overlock 10 Tf
13	The contrasted evolutionary fates of deepâ€sea chemosynthetic mussels (Bivalvia, Bathymodiolinae). Ecology and Evolution, 2013, 3, 4748-4766.	0.8	69
14	Establishing Causes of Eradication Failure Based on Genetics: Case Study of Ship Rat Eradication in Ste. Anne Archipelago. Conservation Biology, 2007, 21, 719-730.	2.4	68
15	The tree, the network, and the species. Biological Journal of the Linnean Society, 2006, 89, 509-521.	0.7	66
16	Molecular phylogeny in mytilids supports the wooden steps to deep-sea vents hypothesis. Comptes Rendus - Biologies, 2007, 330, 446-456.	0.1	64
17	Island colonization and founder effects: the invasion of the Guadeloupe islands by ship rats (Rattus) Tj ETQq $1\ 1\ 0$	0.784314	rgBT/Overloc
18	MORPHOLOGICAL STUDIES OF LYMNAEID SNAILS FROM THE HUMAN FASCIOLIASIS ENDEMIC ZONE OF BOLIVIA. Journal of Molluscan Studies, 2000, 66, 31-44.	0.4	58

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19	A gleam in the dark: Phylogenetic species delimitation in the confusing spring-snail genus Bythinella Moquin-Tandon, 1856 (Gastropoda: Rissooidea: Amnicolidae). Molecular Phylogenetics and Evolution, 2007, 45, 927-941.	1.2	57
20	New taxonomy and old collections: integrating DNA barcoding into the collection curation process. Molecular Ecology Resources, 2012, 12, 396-402.	2.2	57
21	The Coral Sea. Advances in Marine Biology, 2013, 66, 213-290.	0.7	51
22	Is the Species Flock Concept Operational? The Antarctic Shelf Case. PLoS ONE, 2013, 8, e68787.	1.1	51
23	Identifying gastropod spawn from DNA barcodes: possible but not yet practicable. Molecular Ecology Resources, 2009, 9, 1311-1321.	2.2	50
24	Deep-Sea Origin and In-Situ Diversification of Chrysogorgiid Octocorals. PLoS ONE, 2012, 7, e38357.	1.1	50
25	An integrative approach to species delimitation in Benthomangelia (Mollusca: Conoidea). Biological Journal of the Linnean Society, 2009, 96, 696-708.	0.7	49
26	Barcoding type specimens helps to identify synonyms and an unnamed new species in Eumunida Smith, 1883 (Decapoda: Eumunididae). Invertebrate Systematics, 2011, 25, 322.	0.5	48
27	Genetic structure of the saxicole Pitcairnia geyskesii (Bromeliaceae) on inselbergs in French Guiana. American Journal of Botany, 2001, 88, 861-868.	0.8	46
28	Genetic structure of the xerophilous bromeliad <i>Pitcairnia geyskesii</i> on inselbergs in French Guiana – a test of the forest refuge hypothesis. Ecography, 2010, 33, 175-184.	2.1	46
29	An improved taxonomic sampling is a necessary but not sufficient condition for resolving inter-families relationships in Caridean decapods. Genetica, 2015, 143, 195-205.	0.5	45
30	Development of coral and zooxanthella-specific microsatellites in three species of Pocillopora (Cnidaria, Scleractinia) from French Polynesia. Molecular Ecology Notes, 2004, 4, 206-208.	1.7	42
31	Wood-based diet and gut microflora of a galatheid crab associated with Pacific deep-sea wood falls. Marine Biology, 2009, 156, 2421-2439.	0.7	41
32	Molecular and ultrastructural characterization of two ascomycetes found on sunken wood off Vanuatu Islands in the deep Pacific Ocean. Mycological Research, 2009, 113, 1351-1364.	2.5	38
33	Biogeography of the deep-sea galatheid squat lobsters of the Pacific Ocean. Deep-Sea Research Part I: Oceanographic Research Papers, 2010, 57, 228-238.	0.6	38
34	Genetic variation in a network of natural and reintroduced populations of Griffon vulture (Gyps) Tj ETQq0 0 0 rgBT	Oyerlock	≀ 10 Tf 50 14
35	Hidden diversity and endemism on seamounts: focus on poorly dispersive neogastropods. Biological Journal of the Linnean Society, 0, 100, 420-438.	0.7	37
36	A dual process perspective on advances in cognitive science and alcohol use disorder. Clinical Psychology Review, 2019, 69, 83-96.	6.0	37

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37	Speciation patterns in gastropods with longâ€lived larvae from deepâ€sea seamounts. Molecular Ecology, 2012, 21, 4828-4853.	2.0	36
38	Local variation within marinas: Effects of pollutants and implications for invasive species. Marine Pollution Bulletin, 2018, 133, 96-106.	2.3	35
39	Evolution in the deep sea: a combined analysis of the earliest diverging living chitons (Mollusca :) Tj ETQq $1\ 1\ 0.78$	4314 rgBT 0.5	   Overlock
40	Species Delimitation In The Genus Bythinella (Mollusca: Caenogastropoda: Rissooidea): A First Attempt Combining Molecular And Morphometrical Data. Malacologia, 2007, 49, 293-311.	0.2	32
41	Introduction and spread of Thiara granifera (Lamarck, 1822) in Martinique, French West Indies. Biodiversity and Conservation, 1998, 7, 1277-1290.	1.2	31
42	Exploration of the Deep-Sea Fauna of Papua New Guinea. Oceanography, 2012, 25, .	0.5	26
43	Eight new mitogenomes for exploring the phylogeny and classification of Vetigastropoda. Journal of Molluscan Studies, 2016, 82, 534-541.	0.4	26
44	DNA barcoding and molecular systematics of the benthic andÂdemersal organisms of the CEAMARC survey. Polar Science, 2011, 5, 298-312.	0.5	25
45	Next generation sequencing for characterizing biodiversity: promises and challenges. Genetica, 2015, 143, 133-138.	0.5	22
46	Polymorphic microsatellites for the study of fragmented populations of Pitcairnia geyskesii L. B. Smith (Bromeliaceae), a specific saxicolous species of inselbergs in French Guiana. Molecular Ecology Notes, 2003, 3, 221-223.	1.7	21
47	Molluskan species richness and endemism on New Caledonian seamounts: Are they enhanced compared to adjacent slopes?. Deep-Sea Research Part I: Oceanographic Research Papers, 2011, 58, 637-646.	0.6	21
48	An optimised protocol for barcoding museum collections of decapod crustaceans: a case-study for a 10 - 40-years-old collection. Invertebrate Systematics, 2012, 26, 592.	0.5	21
49	Incorporation of deep-sea and small-sized species provides new insights into gastropods phylogeny. Molecular Phylogenetics and Evolution, 2019, 135, 136-147.	1.2	21
50	Density and variability of dinucleotide microsatellites in the parthenogenetic polyploid snail Melanoides tuberculata. Molecular Ecology, 1998, 7, 1233-1236.	2.0	19
51	Rapid morphological changes, admixture and invasive success in populations of Ring-necked parakeets (Psittacula krameri) established in Europe. Biological Invasions, 2016, 18, 1581-1598.	1.2	18
52	Genetic structure and functioning of alien ship rat populations from a Corsican micro-insular complex. Biological Invasions, 2009, 11, 473-482.	1.2	17
53	Title is missing!. Biodiversity and Conservation, 2001, 10, 911-928.	1.2	16
54	Deep-sea benthic communities in the largest oceanic desert are structured by the presence of polymetallic crust. Scientific Reports, 2019, 9, 6977.	1.6	15

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55	First stage zoeal descriptions of five Galatheoidea species from Western Pacific (Crustacea: Decapoda:) Tj ETQq1	10.78431 0.2	4 15gBT /Ove
56	Species from Darwin onward. Integrative Zoology, 2010, 5, 187-197.	1.3	14
57	Integrative Biology of Idas iwaotakii (Habe, 1958), a â€~Model Species' Associated with Sunken Organic Substrates. PLoS ONE, 2013, 8, e69680.	1.1	14
58	Species: towards new, well-grounded practices. Biological Journal of the Linnean Society, 0, 97, 217-222.	0.7	13
59	Patchiness of deepâ€sea communities in Papua New Guinea and potential susceptibility to anthropogenic disturbances illustrated by seep organisms. Marine Ecology, 2015, 36, 109-132.	0.4	12
60	One for each ocean: revision of the Bursa granularis (Röding, 1798) species complex (Gastropoda:) Tj ETQq0 0 0	rgBT /Over	rlock 10 Tf 5
61	Diet and gut microorganisms of <i>Munidopsis </i> squat lobsters associated with natural woods and mesh-enclosed substrates in the deep South Pacific. Marine Biology Research, 2012, 8, 28-47.	0.3	11
62	An interâ€ocean comparison of coral endemism on seamounts: the case of <i>Chrysogorgia</i> . Journal of Biogeography, 2015, 42, 1907-1918.	1.4	10
63	Invasion history and demographic processes associated with rapid morphological changes in the Redâ€whiskered bulbul established on tropical islands. Molecular Ecology, 2016, 25, 5359-5376.	2.0	10
64	Rhodopsin gene evolution in early teleost fishes. PLoS ONE, 2018, 13, e0206918.	1.1	10
65	Isolation and characterization of eight microsatellite loci for the study of gene flow between Testudo marginata and Testudo weissingeri (Testudines: Testudinidae). Molecular Ecology Notes, 2006, 6, 1096-1098.	1.7	7
66	Effects of landscape features and demographic history on the genetic structure of Testudo marginata populations in the southern Peloponnese and Sardinia. Biological Journal of the Linnean Society, 2012, 105, 591-606.	0.7	7
67	The influence of mutation, selection and reproductive systems on microsatellite variability: a simulation approach. Genetical Research, 1998, 71, 213-222.	0.3	5
68	Characterization of eight polymorphic microsatellites in the shrew Crocidura suaveolens and its application to the study of insular populations of the French Atlantic coast. Molecular Ecology Notes, 2004, 4, 426-428.	1.7	5
69	Polymorphic microsatellites for the study of Aconitum napellus L. (Ranunculaceae), a rare species in France. Molecular Ecology Notes, 2005, 5, 358-360.	1.7	5
70	Characterization of seven polymorphic microsatellites for the study of two Ranunculaceae: Ranunculus nodiflorus L., a rare endangered species and Ranunculus flammula L., a common closely related species. Molecular Ecology Notes, 2005, 5, 827-829.	1.7	4
71	Elopomorpha (Teleostei) as a New Model Fish Group for Evolutionary Biology and Comparative Genomics., 2015,, 329-344.		3
72	The crisis in taxonomy. Revue D'Anthropologie Des Connaissances, 2013, 7, .	0.1	3

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73	When Imagery and Physical Sampling Work Together: Toward an Integrative Methodology of Deep-Sea Image-Based Megafauna Identification. Frontiers in Marine Science, 2021, 8, .	1.2	3
74	Formalising Evolutionary Theory. , 2015, , 229-246.		2
75	Untangling species identity in gastropods with polymorphic shells in the genus Bolma Risso, 1826 (Mollusca, Vetigastropoda). European Journal of Taxonomy, 2017, , .	0.6	2
76	VARIATION OF SHELL SHAPE IN THE CLONAL SNAIL MELANOIDES TUBERCULATA AND ITS CONSEQUENCES FOR THE INTERPRETATION OF FOSSIL SERIES. Evolution; International Journal of Organic Evolution, 2000, 54, 492.	1.1	1
77	Evolutionary origins of hydrothermal vents metazoans. BIO Web of Conferences, 2015, 4, 00007.	0.1	1
78	Assembly of the mitochondrial genome of the hydrothermal vent crab <i>Segonzacia mesatlantica</i> i>and detection of potential nuclear pseudogenes. Mitochondrial DNA Part B: Resources, 2017, 2, 291-293.	0.2	1
79	Chapitre 6. La taxonomie et les collections d'histoire naturelle à l'heure de la sixième extinction. , 2014, , 155.		1