

# Isabel Sanmartin

## List of Publications by Year in descending order

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79  
papers

9,140  
citations

101384

36  
h-index

64668

79  
g-index

86  
all docs

86  
docs citations

86  
times ranked

10392  
citing authors

#	ARTICLE	IF	CITATIONS
1	Mitogenomics and hidden trait models reveal the role of phoresy and host shifts in the diversification of parasitoid blister beetles (Coleoptera: Meloidae). <i>Molecular Ecology</i> , 2022, 31, 2453-2474.	2.0	5
2	<i>Cyphoderia ampulla</i> (Cyphoderiidae: Rhizaria), a tale of freshwater sailors: The causes and consequences of ecological transitions through the salinity barrier in a family of benthic protists. <i>Molecular Ecology</i> , 2022, 31, 2644-2663.	2.0	7
3	Macroevolutionary dynamics in the transition of angiosperms to aquatic environments. <i>New Phytologist</i> , 2022, 235, 344-355.	3.5	3
4	Congruent evolutionary responses of European steppe biota to late Quaternary climate change. <i>Nature Communications</i> , 2022, 13, 1921.	5.8	11
5	A road map for phylogenetic models of species trees. <i>Molecular Phylogenetics and Evolution</i> , 2022, 173, 107483.	1.2	3
6	The late Pleistocene endemism increase hypothesis and the origins of diversity in the Canary Islands Flora. <i>Journal of Biogeography</i> , 2022, 49, 1469-1480.	1.4	4
7	More than one sweet tabaiba: Disentangling the systematics of the succulent dendroid shrub <i>Euphorbia balsamifera</i> . <i>Journal of Systematics and Evolution</i> , 2021, 59, 490-503.	1.6	9
8	Performance comparison of two reduced-representation based genome-wide marker-discovery strategies in a multi-taxon phylogeographic framework. <i>Scientific Reports</i> , 2021, 11, 3978.	1.6	7
9	Skipping the Dry Diagonal: spatio-temporal evolution of <i>Croton</i> section <i>Cleodora</i> (Euphorbiaceae) in the Neotropics. <i>Botanical Journal of the Linnean Society</i> , 2021, 197, 61-84.	0.8	10
10	Shared patterns of spatial accumulation of lineages across terrestrial vertebrates. <i>Journal of Biogeography</i> , 2021, 48, 1811-1823.	1.4	3
11	Spread of a SARS-CoV-2 variant through Europe in the summer of 2020. <i>Nature</i> , 2021, 595, 707-712.	13.7	363
12	Women in biogeography. <i>Journal of Biogeography</i> , 2021, 48, 2117-2120.	1.4	4
13	Modelling the tempo and mode of lineage dispersal. <i>Trends in Ecology and Evolution</i> , 2021, 36, 1102-1112.	4.2	13
14	Ecological and geological processes impacting speciation modes drive the formation of wide-range disjunctions within tribe Putorieae (Rubiaceae). <i>Journal of Systematics and Evolution</i> , 2021, 59, 915-934.	1.6	12
15	The first wave of the COVID-19 epidemic in Spain was associated with early introductions and fast spread of a dominating genetic variant. <i>Nature Genetics</i> , 2021, 53, 1405-1414.	9.4	35
16	Biogeography Meets Niche Modeling: Inferring the Role of Deep Time Climate Change When Data Is Limited. <i>Frontiers in Ecology and Evolution</i> , 2021, 9, .	1.1	2
17	Rare and widespread: integrating Bayesian MCMC approaches, Sanger sequencing and HybSeq phylogenomics to reconstruct the origin of the enigmatic Rand Flora genus <i>Camptoloma</i> . <i>American Journal of Botany</i> , 2021, 108, 1673-1691.	0.8	6
18	Long-term isolation of European steppe outposts boosts the biome's conservation value. <i>Nature Communications</i> , 2020, 11, 1968.	5.8	34

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19	Evaluating character partitioning and molecular models in plastid phylogenomics at low taxonomic levels: A case study using <i>Amphilophium</i> (Bignoniaceae). <i>Journal of Systematics and Evolution</i> , 2020, 58, 1071-1089.	1.6	29
20	Colonization time on island settings: lessons from the Hawaiian and Canary Island floras. <i>Botanical Journal of the Linnean Society</i> , 2019, 191, 155-163.	0.8	35
21	Exploring the power of Bayesian birth-death skyline models to detect mass extinction events from phylogenies with only extant taxa. <i>Evolution; International Journal of Organic Evolution</i> , 2019, 73, 1133-1150.	1.1	12
22	Contrasting patterns of diversification between Amazonian and Atlantic forest clades of Neotropical lianas ( <i>Amphilophium</i> , Bignoniaceae) inferred from plastid genomic data. <i>Molecular Phylogenetics and Evolution</i> , 2019, 133, 92-106.	1.2	43
23	High extinction rates and non-adaptive radiation explains patterns of low diversity and extreme morphological disparity in North American blister beetles (Coleoptera, Meloidae). <i>Molecular Phylogenetics and Evolution</i> , 2019, 130, 156-168.	1.2	27
24	Reconstructing deep-time palaeoclimate legacies in the clusioid Malpighiales unveils their role in the evolution and extinction of the boreotropical flora. <i>Global Ecology and Biogeography</i> , 2018, 27, 616-628.	2.7	41
25	Conceptual and statistical problems with the $DEC+J$ model of founder-event speciation and its comparison with $DEC$ via model selection. <i>Journal of Biogeography</i> , 2018, 45, 741-749.	1.4	471
26	Testing the Role of the Red Queen and Court Jester as Drivers of the Macroevolution of Apollo Butterflies. <i>Systematic Biology</i> , 2018, 67, 940-964.	2.7	83
27	Bayesian spatio-temporal reconstruction reveals rapid diversification and Pleistocene range expansion in the widespread columnar cactus <i>Pilosocereus</i> . <i>Journal of Biogeography</i> , 2018, 46, 238.	1.4	25
28	Bridging the micro- and macroevolutionary levels in phylogenomics: $HybSeq$ solves relationships from populations to species and above. <i>New Phytologist</i> , 2018, 220, 636-650.	3.5	152
29	A tale of two forests: ongoing aridification drives population decline and genetic diversity loss at continental scale in Afro-Macaronesian evergreen-forest archipelago endemics. <i>Annals of Botany</i> , 2018, 122, 1005-1017.	1.4	21
30	Morphological Innovations and Vast Extensions of Mountain Habitats Triggered Rapid Diversification Within the Species-Rich Irano-Turanian Genus <i>Acantholimon</i> (Plumbaginaceae). <i>Frontiers in Genetics</i> , 2018, 9, 698.	1.1	22
31	Conceptual and empirical advances in Neotropical biodiversity research. <i>PeerJ</i> , 2018, 6, e5644.	0.9	107
32	Lineage-specific climatic niche drives the tempo of vicariance in the Rand Flora. <i>Journal of Biogeography</i> , 2017, 44, 911-923.	1.4	35
33	Northwest Africa as a source and refuge area of plant biodiversity: a case study on <i>Campanula kremeri</i> and <i>Campanula occidentalis</i> . <i>Journal of Biogeography</i> , 2017, 44, 2057-2068.	1.4	17
34	Opposite trends in the genus <i>Monsonia</i> (Geraniaceae): specialization in the African deserts and range expansions throughout eastern Africa. <i>Scientific Reports</i> , 2017, 7, 9872.	1.6	10
35	Geographic barriers and Pleistocene climate change shaped patterns of genetic variation in the Eastern Afrotropical biodiversity hotspot. <i>Scientific Reports</i> , 2017, 7, 45749.	1.6	58
36	Bipolar distributions in vascular plants: A review. <i>American Journal of Botany</i> , 2017, 104, 1680-1694.	0.8	26

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37	Further progress in historical biogeography. <i>Australian Systematic Botany</i> , 2017, 30, i.	0.3	4
38	Extinction in Phylogenetics and Biogeography: From Timetrees to Patterns of Biotic Assemblage. <i>Frontiers in Genetics</i> , 2016, 7, 35.	1.1	63
39	Late Cretaceous–Early Eocene origin of yams ( <i>Dioscorea</i> , Dioscoreaceae) in the Laurasian Palaeartic and their subsequent Oligocene–Miocene diversification. <i>Journal of Biogeography</i> , 2016, 43, 750-762.	1.4	93
40	Palaeo-islands as refugia and sources of genetic diversity within volcanic archipelagos: the case of the widespread endemic <i>Cyanarina canariensis</i> (Campanulaceae). <i>Molecular Ecology</i> , 2015, 24, 3944-3963.	2.0	50
41	Living on the edge: timing of Rand Flora disjunctions congruent with ongoing aridification in Africa. <i>Frontiers in Genetics</i> , 2015, 6, 154.	1.1	90
42	Islands as model systems in ecology and evolution: prospects fifty years after MacArthur–Wilson. <i>Ecology Letters</i> , 2015, 18, 200-217.	3.0	356
43	Ancient vicariance and climate-driven extinction explain continental-wide disjunctions in Africa: the case of the Rand Flora genus <i>Canarina</i> (Campanulaceae). <i>Molecular Ecology</i> , 2015, 24, 1335-1354.	2.0	58
44	Integrating Fossils, Phylogenies, and Niche Models into Biogeography to Reveal Ancient Evolutionary History: The Case of <i>Hypericum</i> (Hypericaceae). <i>Systematic Biology</i> , 2015, 64, 215-232.	2.7	111
45	Utility of low-copy nuclear markers in phylogenetic reconstruction of <i>Hypericum</i> L. (Hypericaceae). <i>Plant Systematics and Evolution</i> , 2014, 300, 1503-1514.	0.3	7
46	The explosive radiation of <i>Cheirolophus</i> (Asteraceae, Cardueae) in Macaronesia. <i>BMC Evolutionary Biology</i> , 2014, 14, 118.	3.2	47
47	Mediterranean origin and Miocene–Holocene Old World diversification of meadow fescues and ryegrasses ( <i>Festuca</i> subgenus <i>Schedonorus</i> and <i>Lolium</i> ). <i>Journal of Biogeography</i> , 2014, 41, 600-614.	1.4	35
48	Bayesian inference of phylogeny, morphology and range evolution reveals a complex evolutionary history in St. John's wort ( <i>Hypericum</i> ). <i>Molecular Phylogenetics and Evolution</i> , 2013, 67, 379-403.	1.2	68
49	Reconstructing the evolution and biogeographic history of tribe Cardueae (Compositae). <i>American Journal of Botany</i> , 2013, 100, 867-882.	0.8	137
50	Historical Biogeography: Evolution in Time and Space. <i>Evolution: Education and Outreach</i> , 2012, 5, 555-568.	0.3	66
51	DEEP UNDER THE SEA: UNRAVELING THE EVOLUTIONARY HISTORY OF THE DEEP-SEA SQUAT LOBSTER PARAMUNIDA (DECAPODA, MUNIDIDAE). <i>Evolution; International Journal of Organic Evolution</i> , 2012, 66, 1878-1896.	1.1	28
52	Paleobiology of the genus <i>Hypericum</i> (Hypericaceae): a survey of the fossil record and its palaeogeographic implications. <i>Anales Del Jardin Botanico De Madrid</i> , 2012, 69, 97-106.	0.2	18
53	Phylogenetic Methods in Biogeography. <i>Annual Review of Ecology, Evolution, and Systematics</i> , 2011, 42, 441-464.	3.8	222
54	Why are there so many plant species in the Neotropics?. <i>Taxon</i> , 2011, 60, 403-414.	0.4	438

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55	An evaluation of new parsimony-based versus parametric inference methods in biogeography: a case study using the globally distributed plant family Sapindaceae. <i>Journal of Biogeography</i> , 2011, 38, 531-550.	1.4	171
56	Origins of Biodiversityâ€”Response. <i>Science</i> , 2011, 331, 399-400.	6.0	23
57	Mass Extinction, Gradual Cooling, or Rapid Radiation? Reconstructing the Spatiotemporal Evolution of the Ancient Angiosperm Genus <i>Hedyosmum</i> (Chloranthaceae) Using Empirical and Simulated Approaches. <i>Systematic Biology</i> , 2011, 60, 596-615.	2.7	99
58	Amazonia Through Time: Andean Uplift, Climate Change, Landscape Evolution, and Biodiversity. <i>Science</i> , 2010, 330, 927-931.	6.0	1,826
59	Bayesian island biogeography in a continental setting: the Rand Flora case. <i>Biology Letters</i> , 2010, 6, 703-707.	1.0	88
60	Evolutionary Biogeography: An Integrative Approach. <i>Systematic Biology</i> , 2010, 59, 486-488.	2.7	3
61	Mediterranean diversification of the grass-feeding Anisopliinae beetles (Scarabaeidae, Rutelinae). <i>Tj ETQq1 1 0.784314 rgBT /Over</i> <i>Journal of Biogeography</i> , 2009, 36, 546-560.	1.4	26
62	Prospects and challenges for parametric models in historical biogeographical inference. <i>Journal of Biogeography</i> , 2009, 36, 1211-1220.	1.4	164
63	Plastid and nuclear DNA markers reveal intricate relationships at subfamilial and tribal levels in the soapberry family (Sapindaceae). <i>Molecular Phylogenetics and Evolution</i> , 2009, 51, 238-258.	1.2	131
64	Reconstructing the history of Campanulaceae with a Bayesian approach to molecular dating and dispersalâ€”vicariance analyses. <i>Molecular Phylogenetics and Evolution</i> , 2009, 52, 575-587.	1.2	84
65	Tracing the impact of the Andean uplift on Neotropical plant evolution. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 9749-9754.	3.3	550
66	Inferring dispersal: a Bayesian approach to phylogeny-based island biogeography, with special reference to the Canary Islands. <i>Journal of Biogeography</i> , 2008, 35, 428-449.	1.4	208
67	Accounting for Phylogenetic Uncertainty in Biogeography: A Bayesian Approach to Dispersal-Vicariance Analysis of the Thrushes (Aves: Turdus). <i>Systematic Biology</i> , 2008, 57, 257-268.	2.7	336
68	West Wind Drift revisited: testing for directional dispersal in the Southern Hemisphere using event-based tree fitting. <i>Journal of Biogeography</i> , 2007, 34, 398-416.	1.4	138
69	Biogeographic patterns of the East African coastal forest vertebrate fauna. <i>Biodiversity and Conservation</i> , 2007, 16, 883-912.	1.2	39
70	Biology and larval morphology of the genus <i>Ceramida</i> ; Baraud (Coleoptera: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 142 T	0.6	4
71	Biogeography: An Ecological and Evolutionary Approach, 7th edition. <i>Systematic Biology</i> , 2006, 55, 361-363.	2.7	5
72	Southern Hemisphere Biogeography Inferred by Event-Based Models: Plant versus Animal Patterns. <i>Systematic Biology</i> , 2004, 53, 216-243.	2.7	796

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73	Dispersal vs. vicariance in the Mediterranean: historical biogeography of the Palearctic Pachydeminae (Coleoptera, Scarabaeoidea). <i>Journal of Biogeography</i> , 2003, 30, 1883-1897.	1.4	180
74	First phylogenetic analysis of the subfamily Pachydeminae (Coleoptera, Scarabaeoidea, Melolonthidae): the Palearctic Pachydeminae*. <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2003, 41, 2-46.	0.6	26
75	Evoluci3n biogeogr3fica de los Pachydeminae pale3rticos (Coleoptera, Scarabaeoidea) mediante an3lisis de dispersi3n-vicarianza. <i>Graellsia</i> , 2003, 59, 427-441.	0.1	6
76	Patterns of animal dispersal, vicariance and diversification in the Holarctic. <i>Biological Journal of the Linnean Society</i> , 2001, 73, 345-390.	0.7	258
77	Patterns of animal dispersal, vicariance and diversification in the Holarctic. <i>Biological Journal of the Linnean Society</i> , 2001, 73, 345-390.	0.7	326
78	Separation of <i>Aspidiotes</i> species using morphometric analysis (Coleoptera: Curculionidae). <i>European Journal of Entomology</i> , 2000, 97, 85-94.	1.2	5
79	A MORPHOMETRIC APPROACH TO THE TAXONOMY OF THE GENUS <i>CERAMIDA</i> (COLEOPTERA: Tj ETQq1 1 0,784314 rgBT /Overl 0,4 8	0,4	8