

Nadir Alvarez

List of Publications by Year in descending order

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123
papers

5,222
citations

87888

38
h-index

110387

64
g-index

130
all docs

130
docs citations

130
times ranked

7031
citing authors

#	ARTICLE	IF	CITATIONS
1	Hybridization fuelled diversification in <i>Spialia</i> butterflies. <i>Molecular Ecology</i> , 2022, , .	3.9	6
2	The untapped potential of macrofossils in ancient plant DNA research. <i>New Phytologist</i> , 2022, 235, 391-401.	7.3	7
3	Congruent evolutionary responses of European steppe biota to late Quaternary climate change. <i>Nature Communications</i> , 2022, 13, 1921.	12.8	11
4	Poor performance of DNA barcoding and the impact of RAD loci filtering on the species delimitation of an Iberian ant-eating spider. <i>Molecular Phylogenetics and Evolution</i> , 2021, 154, 106997.	2.7	17
5	HyRAD-X Exome Capture Museomics Unravels Giant Ground Beetle Evolution. <i>Genome Biology and Evolution</i> , 2021, 13, .	2.5	13
6	Spatial and temporal heterogeneity in pollinator communities maintains within-species floral odour variation. <i>Oikos</i> , 2021, 130, 1487-1499.	2.7	12
7	The taxonomic impediment: a shortage of taxonomists, not the lack of technical approaches. <i>Zoological Journal of the Linnean Society</i> , 2021, 193, 381-387.	2.3	135
8	The origins and spread of domestic horses from the Western Eurasian steppes. <i>Nature</i> , 2021, 598, 634-640.	27.8	142
9	Climate drives community-wide divergence within species over a limited spatial scale: evidence from an oceanic island. <i>Ecology Letters</i> , 2020, 23, 305-315.	6.4	28
10	Linking seascape with landscape genetics: Oceanic currents favour colonization across the Galápagos Islands by a coastal plant. <i>Journal of Biogeography</i> , 2020, 47, 2622-2633.	3.0	9
11	Rapid colour shift by reproductive character displacement in <i>Cupido</i> butterflies. <i>Molecular Ecology</i> , 2020, 29, 4942-4955.	3.9	10
12	Oviposition choice and larval development of the seed beetle <i>Callosobruchus maculatus</i> (F.) (Coleoptera: Chrysomelidae: Bruchinae) on three cowpea varieties. <i>Journal of Stored Products Research</i> , 2020, 86, 101578.	2.6	7
13	Museomics identifies genetic erosion in two butterfly species across the 20th century in Finland. <i>Molecular Ecology Resources</i> , 2020, 20, 1191-1205.	4.8	34
14	DiscoSnp-RAD: de novo detection of small variants for RAD-Seq population genomics. <i>PeerJ</i> , 2020, 8, e9291.	2.0	5
15	Biogeography and Ecological Diversification of a Mayfly Clade in New Guinea. <i>Frontiers in Ecology and Evolution</i> , 2019, 7, .	2.2	13
16	A mirage of cryptic species: Genomics uncover striking mitonuclear discordance in the butterfly <i>Thymelicus sylvestris</i> . <i>Molecular Ecology</i> , 2019, 28, 3857-3868.	3.9	75
17	Genomic signatures accompanying the dietary shift to phytophagy in polyphagan beetles. <i>Genome Biology</i> , 2019, 20, 98.	8.8	27
18	Bacterial communities within <i>Phengaris</i> (Maculinea) alcon caterpillars are shifted following transition from solitary living to social parasitism of <i>Myrmica</i> ant colonies. <i>Ecology and Evolution</i> , 2019, 9, 4452-4464.	1.9	10

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19	Spatial and temporal genetic dynamics of the grasshopper <i>Oedaleus decorus</i> revealed by museum genomics. <i>Ecology and Evolution</i> , 2018, 8, 1480-1495.	1.9	13
20	Preserving genetic connectivity in the European Alps protected area network. <i>Biological Conservation</i> , 2018, 218, 99-109.	4.1	16
21	Biotic drivers of river and floodplain geomorphology – New molecular methods for assessing present-day and past biota. <i>Earth Surface Processes and Landforms</i> , 2018, 43, 333-338.	2.5	7
22	Long-term in situ persistence of biodiversity in tropical sky islands revealed by landscape genomics. <i>Molecular Ecology</i> , 2018, 27, 432-448.	3.9	39
23	Phylogeography and population genomics of a lotic water beetle across a complex tropical landscape. <i>Molecular Ecology</i> , 2018, 27, 3346-3356.	3.9	12
24	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.	2.9	21
25	Evidence for mega-landslides as drivers of island colonization. <i>Journal of Biogeography</i> , 2017, 44, 1053-1064.	3.0	20
26	Climatic niche evolution is faster in sympatric than allopatric lineages of the butterfly genus <i>Pyrgus</i> . <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2017, 284, 20170208.	2.6	21
27	Assessing the potential of RAD-sequencing to resolve phylogenetic relationships within species radiations: The fly genus <i>Chiastocheta</i> (Diptera: Anthomyiidae) as a case study. <i>Molecular Phylogenetics and Evolution</i> , 2017, 114, 189-198.	2.7	18
28	HyRAD, a versatile method combining exome capture and RAD sequencing to extract genomic information from ancient DNA. <i>Methods in Ecology and Evolution</i> , 2017, 8, 1374-1388.	5.2	52
29	When different contact zones tell different stories: putative ring species in the <i>Megachile concinna</i> species complex (Hymenoptera: Megachilidae). <i>Biological Journal of the Linnean Society</i> , 2017, 121, 815-832.	1.6	18
30	Genomics of extreme ecological specialists: multiple convergent evolution but no genetic divergence between ecotypes of <i>Maculinea alcon</i> butterflies. <i>Scientific Reports</i> , 2017, 7, 13752.	3.3	13
31	Global phylogeography of the insect pest <i>Callosobruchus maculatus</i> (Coleoptera: Bruchinae) relates to the history of its main host, <i>Vigna unguiculata</i> . <i>Journal of Biogeography</i> , 2017, 44, 2515-2526.	3.0	24
32	Elevation in tropical sky islands as the common driver in structuring genes and communities of freshwater organisms. <i>Scientific Reports</i> , 2017, 7, 16089.	3.3	14
33	Large-scale phylogenetic analysis of <i>Amorphophallus</i> (Araceae) derived from nuclear and plastid sequences reveals new subgeneric delineation. <i>Botanical Journal of the Linnean Society</i> , 2017, 184, 32-45.	1.6	27
34	Genetic consequences of Quaternary climatic oscillations in the Himalayas: <i>Primula tibetica</i> as a case study based on restriction site-associated DNA sequencing. <i>New Phytologist</i> , 2017, 213, 1500-1512.	7.3	119
35	High Rate of Protein Coding Sequence Evolution and Species Diversification in the Lycaenids. <i>Frontiers in Ecology and Evolution</i> , 2017, 5, .	2.2	6
36	Is hybridization driving the evolution of climatic niche in <i>Alyssum montanum</i> . <i>American Journal of Botany</i> , 2016, 103, 1348-1357.	1.7	43

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37	Insights into the genetic structure of the cowpea pest <i>Callosobruchus maculatus</i> in Africa. <i>Journal of Pest Science</i> , 2016, 89, 449-458.	3.7	10
38	Differential phenotypic and genetic expression of defence compounds in a plant–herbivore interaction along elevation. <i>Royal Society Open Science</i> , 2016, 3, 160226.	2.4	14
39	Past climate-driven range shifts and population genetic diversity in arctic plants. <i>Journal of Biogeography</i> , 2016, 43, 461-470.	3.0	48
40	Clustering Genes of Common Evolutionary History. <i>Molecular Biology and Evolution</i> , 2016, 33, 1590-1605.	8.9	51
41	Combining conservative and variable markers to infer the evolutionary history of <i>Prunus</i> subgen. <i>Amygdalus</i> s.l. under domestication. <i>Genetic Resources and Crop Evolution</i> , 2016, 63, 221-234.	1.6	14
42	Hybridization Capture Using RAD Probes (hyRAD), a New Tool for Performing Genomic Analyses on Collection Specimens. <i>PLoS ONE</i> , 2016, 11, e0151651.	2.5	121
43	Genome fingerprinting confirms the species status of the Loosetrifles <i>Lysimachia punctata</i> and <i>L. verticillata</i> (Primulaceae). <i>Plant Ecology and Evolution</i> , 2016, 149, 335-338.	0.7	0
44	Asymmetrical nature of the <i>Trollius</i> – <i>C. hiastocheta</i> interaction: insights into the evolution of nursery pollination systems. <i>Ecology and Evolution</i> , 2015, 5, 4766-4777.	1.9	14
45	Fifty years after Ehrlich and Raven, is there support for plant–insect coevolution as a major driver of species diversification?. <i>Entomologia Experimentalis Et Applicata</i> , 2015, 157, 98-112.	1.4	65
46	Uncovering Cryptic Parasitoid Diversity in <i>Horismenus</i> (Chalcidoidea, Eulophidae). <i>PLoS ONE</i> , 2015, 10, e0136063.	2.5	17
47	Decoupled post-glacial history in mutualistic plant–insect interactions: insights from the yellow loosestrife (<i>Lysimachia vulgaris</i>) and its associated oil-collecting bees (<i>Macropis europaea</i>) <i>Tj ETQ</i> , 2015, 1, 0.784314	1.4	13
48	Restriction site-associated DNA sequencing, genotyping error estimation and <i>de novo</i> assembly optimization for population genetic inference. <i>Molecular Ecology Resources</i> , 2015, 15, 28-41.	4.8	345
49	Gene Duplication, Population Genomics, and Species-Level Differentiation within a Tropical Mountain Shrub. <i>Genome Biology and Evolution</i> , 2014, 6, 2611-2624.	2.5	25
50	Comparative phylogeography of mutualists and the effect of the host on the genetic structure of its partners. <i>Biological Journal of the Linnean Society</i> , 2014, 113, 1021-1035.	1.6	23
51	Proto-South-East Asia as a trigger of early angiosperm diversification. <i>Botanical Journal of the Linnean Society</i> , 2014, 174, 326-333.	1.6	18
52	Wheat alleles introgress into selfing wild relatives: empirical estimates from approximate Bayesian computation in <i>Aegilops triuncialis</i> . <i>Molecular Ecology</i> , 2014, 23, 5089-5101.	3.9	11
53	Evolutionary history of almond tree domestication in the Mediterranean basin. <i>Molecular Ecology</i> , 2013, 22, 1092-1104.	3.9	55
54	Identifying genetic signatures of selection in a non-model species, alpine gentian (<i>Gentiana nivalis</i> L.), using a landscape genetic approach. <i>Conservation Genetics</i> , 2013, 14, 467-481.	1.5	65

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55	Morphological, ecological and genetic aspects associated with endemism in the <i>ly</i> <i>O</i> rhcid group. <i>Molecular Ecology</i> , 2013, 22, 1431-1446.	3.9	20
56	Phylogenetic alpha and beta diversities of butterfly communities correlate with climate in the western Swiss Alps. <i>Ecography</i> , 2013, 36, 541-550.	4.5	48
57	The abrupt climate change at the Eocene–Oligocene boundary and the emergence of South-East Asia triggered the spread of sapindaceous lineages. <i>Annals of Botany</i> , 2013, 112, 151-160.	2.9	46
58	Nature, Evolution and Characterisation of Rhizospheric Chemical Exudates Affecting Root Herbivores. <i>Advances in Insect Physiology</i> , 2013, , 97-157.	2.7	25
59	A deep digâ€ˆhindsight on Holocene vegetation composition from ancient environmental <i>DNA</i> . <i>Molecular Ecology</i> , 2013, 22, 3433-3436.	3.9	11
60	Genetic diversity in widespread species is not congruent with species richness in alpine plant communities. <i>Ecology Letters</i> , 2012, 15, 1439-1448.	6.4	135
61	Molecular substitution rate increases in myrmecophilous lycaenid butterflies (Lepidoptera). <i>Zoologica Scripta</i> , 2012, 41, 651-658.	1.7	17
62	Ecological and historical drivers of diversification in the fly genus <i>Chiastocheta</i> Pokorný. <i>Molecular Phylogenetics and Evolution</i> , 2012, 63, 466-474.	2.7	12
63	Contrasting diffusion of Quaternary gene pools across Europe: The case of the arctic–alpine <i>Gentiana nivalis</i> L. (Gentianaceae). <i>Flora: Morphology, Distribution, Functional Ecology of Plants</i> , 2012, 207, 408-413.	1.2	18
64	Climate oscillations and species interactions: large-scale congruence but regional differences in the phylogeographic structures of an alpine plant and its monophagous insect. <i>Journal of Biogeography</i> , 2012, 39, 1487-1498.	3.0	16
65	Gene flow among wild and domesticated almond species: insights from chloroplast and nuclear markers. <i>Evolutionary Applications</i> , 2012, 5, 317-329.	3.1	65
66	RECONSTRUCTING THE ORIGINS OF HIGH-ALPINE NICHES AND CUSHION LIFE FORM IN THE GENUS <i>ANDROSACE</i> S.L. (PRIMULACEAE). <i>Evolution; International Journal of Organic Evolution</i> , 2012, 66, 1255-1268.	2.3	69
67	Forecasting changes in population genetic structure of alpine plants in response to global warming. <i>Molecular Ecology</i> , 2012, 21, 2354-2368.	3.9	127
68	Broad-scale adaptive genetic variation in alpine plants is driven by temperature and precipitation. <i>Molecular Ecology</i> , 2012, 21, 3729-3738.	3.9	161
69	Predicting present and future intra-specific genetic structure through niche hindcasting across 24 millennia. <i>Ecology Letters</i> , 2012, 15, 649-657.	6.4	79
70	Phylogenetic relationships in the subfamily Psychodinae (Diptera, Psychodidae). <i>Zoologica Scripta</i> , 2012, 41, 489-498.	1.7	12
71	Investigating the relationship between pollination strategies and the size-advantage model in zoophilous plants using the reproductive biology of <i>Arum cylindraceum</i> and other European <i>Arum</i> species as case studies. <i>Arthropod-Plant Interactions</i> , 2012, 6, 35-44.	1.1	16
72	Automated Scoring of AFLPs Using RawGeno v 2.0, a Free R CRAN Library. <i>Methods in Molecular Biology</i> , 2012, 888, 155-175.	0.9	38

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73	Pollinators as drivers of plant distribution and assemblage into communities. , 2011, , 392-413.		8
74	Comparative Phylogeography in a Specific and Obligate Pollination Antagonism. PLoS ONE, 2011, 6, e28662.	2.5	23
75	An evaluation of new parsimony-based versus parametric inference methods in biogeography: a case study using the globally distributed plant family Sapindaceae. Journal of Biogeography, 2011, 38, 531-550.	3.0	171
76	Break zones in the distributions of alleles and species in alpine plants. Journal of Biogeography, 2011, 38, 772-782.	3.0	77
77	Insights into the biogeographical history of the Lower Guinea Forest Domain: evidence for the role of refugia in the intraspecific differentiation of <i>Aucoumea klaineana</i> . Molecular Ecology, 2011, 20, 131-142.	3.9	32
78	Discordances between phylogenetic and morphological patterns in alpine leaf beetles attest to an intricate biogeographic history of lineages in postglacial Europe. Molecular Ecology, 2011, 20, 2442-2463.	3.9	20
79	Variation in the proportion of flower visitors of <i>Arum maculatum</i> along its distributional range in relation with community-based climatic niche analyses. Oikos, 2011, 120, 728-734.	2.7	25
80	Cleaning wrasse species vary with respect to dependency on the mutualism and behavioural adaptations in interactions. Animal Behaviour, 2011, 82, 1067-1074.	1.9	27
81	Ecological niche overlap in sister species: how do oil-collecting bees <i>Macropis europaea</i> and <i>Macropis fulvipes</i> (Hymenoptera: Melittidae) avoid hybridization and competition?. Apidologie, 2011, 42, 579-595.	2.0	9
82	Does a shift in host plants trigger speciation in the Alpine leaf beetle <i>Oreina speciosissima</i> (Coleoptera.) Tj ETQq0 0,0 r gBT /Overlock 10	3.25	9
83	Comparative Performance of Supertree Algorithms in Large Data Sets Using the Soapberry Family (Sapindaceae) as a Case Study. Systematic Biology, 2011, 60, 32-44.	5.6	25
84	The phylogeography of an alpine leaf beetle: Divergence within <i>Oreina elongata</i> spans several ice ages. Molecular Phylogenetics and Evolution, 2010, 57, 703-709.	2.7	28
85	Origin and expansion of the allotetraploid <i>Aegilops taenensis</i> , a wild relative of wheat. New Phytologist, 2010, 187, 1170-1180.	7.3	58
86	Population genetic structure of two primary parasitoids of <i>Spodoptera frugiperda</i> (Lepidoptera), <i>Chelonus insularis</i> and <i>Campoletis sonorensis</i> (Hymenoptera): to what extent is the host plant important?. Molecular Ecology, 2010, 19, 2168-2179.	3.9	12
87	New insights into the phylogenetics and biogeography of <i>Arum</i> (Araceae): unravelling its evolutionary history. Botanical Journal of the Linnean Society, 2010, 163, 14-32.	1.6	33
88	Phylogeny of subtribe Gentianinae (Gentianaceae): Biogeographic inferences despite limitations in temporal calibration points. Taxon, 2010, 59, 1701-1711.	0.7	44
89	Phylogeny and circumscription of Sapindaceae revisited: molecular sequence data, morphology and biogeography support recognition of a new family, Xanthoceraceae. Plant Ecology and Evolution, 2010, 143, 148-159.	0.7	77
90	Phylogeography of <i>Chelonus insularis</i> (Hymenoptera: Braconidae) and <i>Campoletis sonorensis</i> (Hymenoptera: Ichneumonidae), Two Primary Neotropical Parasitoids of the Fall Armyworm (Lepidoptera: Noctuidae). Annals of the Entomological Society of America, 2010, 103, 742-749.	2.5	11

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91	The effect of host plant and isolation on the genetic structure of phytophagous insects: A preliminary study on a bruchid beetle. <i>European Journal of Entomology</i> , 2010, 107, 299-304.	1.2	5
92	Do Sebaciales commonly associate with plant roots as endophytes?. <i>Mycological Research</i> , 2009, 113, 1062-1069.	2.5	125
93	Evaluating the impact of scoring parameters on the structure of intra-specific genetic variation using RawGeno, an R package for automating AFLP scoring. <i>BMC Bioinformatics</i> , 2009, 10, 33.	2.6	144
94	Malagasy <i>Dracaena</i> Vand. ex L. (Ruscaceae): an investigation of discrepancies between morphological features and spatial genetic structure at a small evolutionary scale. <i>Plant Systematics and Evolution</i> , 2009, 280, 15-28.	0.9	6
95	Alpine "subalpine species richness of the Romanian Carpathians and the current conservation status of rare species. <i>Biodiversity and Conservation</i> , 2009, 18, 1441-1458.	2.6	21
96	Genetic structure and evolution of Alpine polyploid complexes: <i>Ranunculus kuepferi</i> (<i>Ranunculaceae</i>) as a case study. <i>Molecular Ecology</i> , 2009, 18, 3730-3744.	3.9	71
97	Concordant genetic breaks, identified by combining clustering and tessellation methods, in two co-distributed alpine plant species. <i>Molecular Ecology</i> , 2009, 18, 4495-4507.	3.9	34
98	History or ecology? Substrate type as a major driver of patial genetic structure in Alpine plants. <i>Ecology Letters</i> , 2009, 12, 632-640.	6.4	167
99	Effects of species traits on the genetic diversity of high-mountains plants: a multi-species study across the Alps and the Carpathians. <i>Global Ecology and Biogeography</i> , 2009, 18, 78-87.	5.8	62
100	Plastid and nuclear DNA markers reveal intricate relationships at subfamilial and tribal levels in the soapberry family (<i>Sapindaceae</i>). <i>Molecular Phylogenetics and Evolution</i> , 2009, 51, 238-258.	2.7	131
101	Isolation and characterization of polymorphic microsatellite loci in two primary parasitoids of the noctuid <i>Spodoptera frugiperda</i> : <i>Chelonus insularis</i> and <i>Campoletis sonorensis</i> (<i>Hymenoptera</i>). <i>Molecular Ecology Resources</i> , 2009, 9, 171-173.	4.8	4
102	Identification of seven species of hymenopteran parasitoids of <i>Spodoptera frugiperda</i> , using polymerase chain reaction amplification and restriction enzyme digestion. <i>Agricultural and Forest Entomology</i> , 2008, 10, 129-136.	1.3	33
103	Historical divergence vs. contemporary gene flow: evolutionary history of the calcicole <i>Ranunculus alpestris</i> group (<i>Ranunculaceae</i>) in the European Alps and the Carpathians. <i>Molecular Ecology</i> , 2008, 17, 4263-4275.	3.9	98
104	Assessing the phylogenetic usefulness of a previously neglected morphological structure through elliptic Fourier analyses: a case study in <i>Bruchus</i> seed-beetles (Coleoptera: Chrysomelidae: Tj ETQq0 0 0 rgrBT /Overlook 10 Tf 5	3.9	98
105	SIMIL: an <code>r</code> (CRAN) scripts collection for computing genetic structure similarities based on <code>structure</code> 2 outputs. <i>Molecular Ecology Resources</i> , 2008, 8, 757-762.	4.8	6
106	A new individual-based spatial approach for identifying genetic discontinuities in natural populations. <i>Molecular Ecology</i> , 2007, 16, 2031-2043.	3.9	72
107	Genetic consequences of Pleistocene range shifts: contrast between the Arctic, the Alps and the East African mountains. <i>Molecular Ecology</i> , 2007, 16, 2542-2559.	3.9	183
108	Genetic structure of <i>Hypochaeris uniflora</i> (<i>Asteraceae</i>) suggests vicariance in the Carpathians and rapid post-glacial colonization of the Alps from an eastern Alpine refugium. <i>Journal of Biogeography</i> , 2007, 34, 2100-2114.	3.0	90

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109	ANTHROPOGENIC EFFECTS ON POPULATION GENETICS OF PHYTOPHAGOUS INSECTS ASSOCIATED WITH DOMESTICATED PLANTS. <i>Evolution; International Journal of Organic Evolution</i> , 2007, 61, 2986-2996.	2.3	22
110	Genetic and Environmental Sources of Variation in the Autogenous Chemical Defense of a Leaf Beetle. <i>Journal of Chemical Ecology</i> , 2007, 33, 2011-2024.	1.8	7
111	Phylogeographic support for horizontal gene transfer involving sympatric bruchid species. <i>Biology Direct</i> , 2006, 1, 21.	4.6	12
112	Phylogenetic relationships in the Neotropical bruchid genus <i>Acanthoscelides</i> (Bruchinae, Bruchidae.) <i>Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5</i>	1.4	24
113	Ecological distribution and niche segregation of sibling species: the case of bean beetles, <i>Acanthoscelides obtectus</i> Say and <i>A. obvelatus</i> Bridwell. <i>Ecological Entomology</i> , 2006, 31, 582-590.	2.2	15
114	The Effect of Fibrin Glue on Autogenous and Alloplastic Bone Grafts in Rat Calvarial Defects. <i>Journal of Oral and Maxillofacial Surgery</i> , 2006, 64, 63-64.	1.2	36
115	Evolutionary history and patterns of differentiation among European <i>Maniola</i> butterflies (Lepidoptera: Satyrinae). <i>European Journal of Entomology</i> , 2006, 103, 613-618.	1.2	8
116	Sibling species of bean bruchids: a morphological and phylogenetic study of <i>Acanthoscelides obtectus</i> Say and <i>Acanthoscelides obvelatus</i> Bridwell. <i>Journal of Zoological Systematics and Evolutionary Research</i> , 2005, 43, 29-37.	1.4	23
117	Ancient and recent evolutionary history of the bruchid beetle, <i>Acanthoscelides obtectus</i> Say, a cosmopolitan pest of beans. <i>Molecular Ecology</i> , 2005, 14, 1015-1024.	3.9	53
118	Parallels in the evolution of the two largest New and Old World seed-beetle genera (Coleoptera.) <i>Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 3</i>	3.9	40
119	Farmers' practices, metapopulation dynamics, and conservation of agricultural biodiversity on-farm: a case study of sorghum among the Duupa in sub-sahelian Cameroon. <i>Biological Conservation</i> , 2005, 121, 533-543.	4.1	75
120	Microsatellite markers in a complex of <i>Horismenus</i> sp. (Hymenoptera: Eulophidae), parasitoids of bruchid beetles. <i>Molecular Ecology Notes</i> , 2004, 4, 707-709.	1.7	5
121	Isolation and characterization of polymorphic microsatellite loci in <i>Acanthoscelides obtectus</i> Say (Coleoptera: Bruchidae). <i>Molecular Ecology Notes</i> , 2004, 4, 683-685.	1.7	4
122	Isolation and characterization of polymorphic microsatellite markers in <i>Zabrotes subfasciatus</i> Boheman (Coleoptera: Bruchidae). <i>Molecular Ecology Notes</i> , 2004, 4, 752-754.	1.7	5
123	Isolation and characterization of polymorphic microsatellite loci in <i>Acanthoscelides obvelatus</i> Bridwell (Coleoptera: Bruchidae). <i>Molecular Ecology Notes</i> , 2002, 3, 12-14.	1.7	8