# Nadir Alvarez

List of Publications by Year
in descending order

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Version: 2024-02-01

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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 | Restriction siteâ€associated <scp>DNA</scp> sequencing, genotyping error estimation and de novo assembly optimization for population genetic inference. Molecular Ecology Resources, 2015, 15, 28-41. | 4.8 | 345 |
| 2 | Genetic consequences of Pleistocene range shifts: contrast between the Arctic, the Alps and the East African mountains. Molecular Ecology, 2007, 16, 2542-2559. | 3.9 | 183 |
| 3 | 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 |
| 4 | History or ecology? Substrate type as a major driver of patial genetic structure in Alpine plants. Ecology Letters, 2009, 12, 632-640. | 6.4 | 167 |
| 5 | Broadâ€scale adaptive genetic variation in alpine plants is driven by temperature and precipitation. Molecular Ecology, 2012, 21, 3729-3738. | 3.9 | 161 |
| 6 | Evaluating the impact of scoring parameters on the structure of intra-specific genetic variation using RawGeno, an R package for automating AFLP scoring. BMC Bioinformatics, 2009, 10, 33. | 2.6 | 144 |
| 7 | The origins and spread of domestic horses from the Western Eurasian steppes. Nature, 2021, 598, 634-640. | 27.8 | 142 |
| 8 | Genetic diversity in widespread species is not congruent with species richness in alpine plant communities. Ecology Letters, 2012, 15, 1439-1448. | 6.4 | 135 |
| 9 | The taxonomic impediment: a shortage of taxonomists, not the lack of technical approaches. Zoological Journal of the Linnean Society, 2021, 193, 381-387. | 2.3 | 135 |
| 10 | Plastid and nuclear DNA markers reveal intricate relationships at subfamilial and tribal levels in the soapberry family (Sapindaceae). Molecular Phylogenetics and Evolution, 2009, 51, 238-258. | 2.7 | 131 |
| 11 | Forecasting changes in population genetic structure of alpine plants in response to global warming. Molecular Ecology, 2012, 21, 2354-2368. | 3.9 | 127 |
| 12 | Do Sebacinales commonly associate with plant roots as endophytes?. Mycological Research, 2009, 113, 1062-1069. | 2.5 | 125 |
| 13 | Hybridization Capture Using RAD Probes (hyRAD), a New Tool for Performing Genomic Analyses on Collection Specimens. PLoS ONE, 2016, 11, e0151651. | 2.5 | 121 |
| 14 | Genetic consequences of Quaternary climatic oscillations in the Himalayas: Primula tibetica as a case study based on restriction siteâ€associated DNA sequencing. New Phytologist, 2017, 213, 1500-1512. | 7.3 | 119 |
| 15 | Historical divergence vs. contemporary gene flow: evolutionary history of the calcicole Ranunculus alpestris group (Ranunculaceae) in the European Alps and the Carpathians. Molecular Ecology, 2008, 17, 4263-4275. | 3.9 | 98 |
| 16 | Genetic structure of Hypochaeris uniflora (Asteraceae) suggests vicariance in the Carpathians and rapid postâ€glacial colonization of the Alps from an eastern Alpine refugium. Journal of Biogeography, 2007, 34, 2100-2114. | 3.0 | 90 |
| 17 | Predicting present and future intraâ€specific genetic structure through niche hindcasting across 24 millennia. Ecology Letters, 2012, 15, 649-657. | 6.4 | 79 |
| 18 | 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 |

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|---|---|---|---|
| 19 | Break zones in the distributions of alleles and species in alpine plants. Journal of Biogeography, 2011, 38, 772-782. | 3.0 | 77 |
| 20 | Farmers' practices, metapopulation dynamics, and conservation of agricultural biodiversity on-farm: a case study of sorghum among the Duupa in sub-sahelian Cameroon. Biological Conservation, 2005, 121, 533-543. | 4.1 | 75 |
| 21 | A mirage of cryptic species: Genomics uncover striking mitonuclear discordance in the butterfly Thymelicus sylvestris. Molecular Ecology, 2019, 28, 3857-3868. | 3.9 | 75 |
| 22 | A new individual-based spatial approach for identifying genetic discontinuities in natural populations. Molecular Ecology, 2007, 16, 2031-2043. | 3.9 | 72 |
| 23 | Genetic structure and evolution of Alpine polyploid complexes: Ranunculus kuepferi (Ranunculaceae) as a case study. Molecular Ecology, 2009, 18, 3730-3744. | 3.9 | 71 |
| 24 | RECONSTRUCTING THE ORIGINS OF HIGH-ALPINE NICHES AND CUSHION LIFE FORM IN THE GENUS ANDROSACE S.L. (PRIMULACEAE). Evolution; International Journal of Organic Evolution, 2012, 66, 1255-1268. | 2.3 | 69 |
| 25 | Gene flow among wild and domesticated almond species: insights from chloroplast and nuclear markers. Evolutionary Applications, 2012, 5, 317-329. | 3.1 | 65 |
| 26 | Identifying genetic signatures of selection in a non-model species, alpine gentian (Gentiana nivalis L.), using a landscape genetic approach. Conservation Genetics, 2013, 14, 467-481. | 1.5 | 65 |
| 27 | Fifty years after Ehrlich and Raven, is there support for plantâ€“insect coevolution as a major driver of species diversification?. Entomologia Experimentalis Et Applicata, 2015, 157, 98-112. | 1.4 | 65 |
| 28 | Effects of species traits on the genetic diversity of highâ€mountain plants: a multiâ€species study across the Alps and the Carpathians. Global Ecology and Biogeography, 2009, 18, 78-87. | 5.8 | 62 |
| 29 | Origin and expansion of the allotetraploid Aegilopsâ€ƒgeniculata, a wild relative of wheat. New Phytologist, 2010, 187, 1170-1180. | 7.3 | 58 |
| 30 | Evolutionary history of almond tree domestication in the <scp>M</scp>editerranean basin. Molecular Ecology, 2013, 22, 1092-1104. | 3.9 | 55 |
| 31 | Ancient and recent evolutionary history of the bruchid beetle, Acanthoscelides obtectus Say, a cosmopolitan pest of beans. Molecular Ecology, 2005, 14, 1015-1024. | 3.9 | 53 |
| 32 | Hy<scp>RAD</scp>â€, a versatile method combining exome capture and <scp>RAD</scp> sequencing to extract genomic information from ancient <scp>DNA</scp>. Methods in Ecology and Evolution, 2017, 8, 1374-1388. | 5.2 | 52 |
| 33 | Clustering Genes of Common Evolutionary History. Molecular Biology and Evolution, 2016, 33, 1590-1605. | 8.9 | 51 |
| 34 | Phylogenetic alpha and beta diversities of butterfly communities correlate with climate in the western Swiss Alps. Ecography, 2013, 36, 541-550. | 4.5 | 48 |
| 35 | Past climateâ€driven range shifts and population genetic diversity in arctic plants. Journal of Biogeography, 2016, 43, 461-470. | 3.0 | 48 |
| 36 | The abrupt climate change at the Eoceneâ€“Oligocene boundary and the emergence of South-East Asia triggered the spread of sapindaceous lineages. Annals of Botany, 2013, 112, 151-160. | 2.9 | 46 |

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|---|---|---|---|
| 37 | Phylogeny of subtribe Gentianinae (Gentianaceae): Biogeographic inferences despite limitations in temporal calibration points. Taxon, 2010, 59, 1701-1711. | 0.7 | 44 |
| 38 | Is hybridization driving the evolution of climatic niche in Alyssum montanum. American Journal of Botany, 2016, 103, 1348-1357. | 1.7 | 43 |
| 39 | Parallels in the evolution of the two largest New and Old World seed-beetle genera (Coleoptera,) Tj ETQq1 1 0.784314 rgBT /Overlock | 3.9 | 40 |
| 40 | Longâ€term inÂsitu persistence of biodiversity in tropical sky islands revealed by landscape genomics. Molecular Ecology, 2018, 27, 432-448. | 3.9 | 39 |
| 41 | Automated Scoring of AFLPs Using RawGeno v 2.0, a Free R CRAN Library. Methods in Molecular Biology, 2012, 888, 155-175. | 0.9 | 38 |
| 42 | The Effect of Fibrin Glue on Autogenous and Alloplastic Bone Grafts in Rat Calvarial Defects. Journal of Oral and Maxillofacial Surgery, 2006, 64, 63-64. | 1.2 | 36 |
| 43 | Concordant genetic breaks, identified by combining clustering and tessellation methods, in two coâ€distributed alpine plant species. Molecular Ecology, 2009, 18, 4495-4507. | 3.9 | 34 |
| 44 | Museomics identifies genetic erosion in two butterfly species across the 20th century in Finland. Molecular Ecology Resources, 2020, 20, 1191-1205. | 4.8 | 34 |
| 45 | Identification of seven species of hymenopteran parasitoids of Spodoptera frugiperda, using polymerase chain reaction amplification and restriction enzyme digestion. Agricultural and Forest Entomology, 2008, 10, 129-136. | 1.3 | 33 |
| 46 | New insights into the phylogenetics and biogeography of Arum (Araceae): unravelling its evolutionary history. Botanical Journal of the Linnean Society, 2010, 163, 14-32. | 1.6 | 33 |
| 47 | Insights into the biogeographical history of the Lower Guinea Forest Domain: evidence for the role of refugia in the intraspecific differentiation of Aucoumea klaineana. Molecular Ecology, 2011, 20, 131-142. | 3.9 | 32 |
| 48 | The phylogeography of an alpine leaf beetle: Divergence within Oreina elongata spans several ice ages. Molecular Phylogenetics and Evolution, 2010, 57, 703-709. | 2.7 | 28 |
| 49 | Climate drives communityâ€wide divergence within species over a limited spatial scale: evidence from an oceanic island. Ecology Letters, 2020, 23, 305-315. | 6.4 | 28 |
| 50 | 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 |
| 51 | Large-scale phylogenetic analysis of Amorphophallus (Araceae) derived from nuclear and plastid sequences reveals new subgeneric delineation. Botanical Journal of the Linnean Society, 2017, 184, 32-45. | 1.6 | 27 |
| 52 | Genomic signatures accompanying the dietary shift to phytophagy in polyphagan beetles. Genome Biology, 2019, 20, 98. | 8.8 | 27 |
| 53 | Variation in the proportion of flower visitors of Arum maculatum along its distributional range in relation with communityâ€based climatic niche analyses. Oikos, 2011, 120, 728-734. | 2.7 | 25 |
| 54 | 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 |

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|---|---|---|---|
| 55 | Nature, Evolution and Characterisation of Rhizospheric Chemical Exudates Affecting Root Herbivores. Advances in Insect Physiology, 2013, , 97-157. | 2.7 | 25 |
| 56 | Gene Duplication, Population Genomics, and Species-Level Differentiation within a Tropical Mountain Shrub. Genome Biology and Evolution, 2014, 6, 2611-2624. | 2.5 | 25 |
| 57 | Phylogenetic relationships in the Neotropical bruchid genus Acanthoscelides (Bruchinae, Bruchidae,) Tj ETQq1 1 0.784314 rgBT /Over | 1.4 | 24 |
| 58 | Global phylogeography of the insect pest Callosobruchus maculatus (Coleoptera: Bruchinae) relates to the history of its main host, Vigna unguiculata. Journal of Biogeography, 2017, 44, 2515-2526. | 3.0 | 24 |
| 59 | Sibling species of bean bruchids: a morphological and phylogenetic study of Acanthoscelides obtectus Say and Acanthoscelides obvelatus Bridwell. Journal of Zoological Systematics and Evolutionary Research, 2005, 43, 29-37. | 1.4 | 23 |
| 60 | Comparative Phylogeography in a Specific and Obligate Pollination Antagonism. PLoS ONE, 2011, 6, e28662. | 2.5 | 23 |
| 61 | Comparative phylogeography of mutualists and the effect of the host on the genetic structure of its partners. Biological Journal of the Linnean Society, 2014, 113, 1021-1035. | 1.6 | 23 |
| 62 | ANTHROPOGENIC EFFECTS ON POPULATION GENETICS OF PHYTOPHAGOUS INSECTS ASSOCIATED WITH DOMESTICATED PLANTS. Evolution; International Journal of Organic Evolution, 2007, 61, 2986-2996. | 2.3 | 22 |
| 63 | Alpineâ€“subalpine species richness of the Romanian Carpathians and the current conservation status of rare species. Biodiversity and Conservation, 2009, 18, 1441-1458. | 2.6 | 21 |
| 64 | Climatic niche evolution is faster in sympatric than allopatric lineages of the butterfly genus Pyrgus. Proceedings of the Royal Society B: Biological Sciences, 2017, 284, 20170208. | 2.6 | 21 |
| 65 | A tale of two forests: ongoing aridification drives population decline and genetic diversity loss at continental scale in Afro-Macaronesian evergreen-forest archipelago endemics. Annals of Botany, 2018, 122, 1005-1017. | 2.9 | 21 |
| 66 | 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 |
| 67 | Morphological, ecological and genetic aspects associated with endemism in the <scp>F</scp>ly <scp>O</scp>rchid group. Molecular Ecology, 2013, 22, 1431-1446. | 3.9 | 20 |
| 68 | Evidence for megaâ€landslides as drivers of island colonization. Journal of Biogeography, 2017, 44, 1053-1064. | 3.0 | 20 |
| 69 | Contrasting diffusion of Quaternary gene pools across Europe: The case of the arcticâ€“alpine Gentiana nivalis L. (Gentianaceae). Flora: Morphology, Distribution, Functional Ecology of Plants, 2012, 207, 408-413. | 1.2 | 18 |
| 70 | Proto-South-East Asia as a trigger of early angiosperm diversification. Botanical Journal of the Linnean Society, 2014, 174, 326-333. | 1.6 | 18 |
| 71 | Assessing the potential of RAD-sequencing to resolve phylogenetic relationships within species radiations: The fly genus Chiastocheta (Diptera: Anthomyiidae) as a case study. Molecular Phylogenetics and Evolution, 2017, 114, 189-198. | 2.7 | 18 |
| 72 | When different contact zones tell different stories: putative ring species in the Megachile concinna species complex (Hymenoptera: Megachilidae). Biological Journal of the Linnean Society, 2017, 121, 815-832. | 1.6 | 18 |

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|---|---|---|---|
| 73 | Molecular substitution rate increases in myrmecophilous lycaenid butterflies (Lepidoptera). Zoologica Scripta, 2012, 41, 651-658. | 1.7 | 17 |
| 74 | Uncovering Cryptic Parasitoid Diversity in Horismenus (Chalcidoidea, Eulophidae). PLoS ONE, 2015, 10, e0136063. | 2.5 | 17 |
| 75 | Poor performance of DNA barcoding and the impact of RAD loci filtering on the species delimitation of an Iberian ant-eating spider. Molecular Phylogenetics and Evolution, 2021, 154, 106997. | 2.7 | 17 |
| 76 | Climate oscillations and species interactions: largeâ€scale congruence but regional differences in the phylogeographic structures of an alpine plant and its monophagous insect. Journal of Biogeography, 2012, 39, 1487-1498. | 3.0 | 16 |
| 77 | Investigating the relationship between pollination strategies and the size-advantage model in zoophilous plants using the reproductive biology of Arum cylindraceum and other European Arum species as case studies. Arthropod-Plant Interactions, 2012, 6, 35-44. | 1.1 | 16 |
| 78 | Preserving genetic connectivity in the European Alps protected area network. Biological Conservation, 2018, 218, 99-109. | 4.1 | 16 |
| 79 | Ecological distribution and niche segregation of sibling species: the case of bean beetles, Acanthoscelides obtectus Say and A. obvelatus Bridwell. Ecological Entomology, 2006, 31, 582-590. | 2.2 | 15 |
| 80 | Asymmetrical nature of the <scp>T</scp>rolliusâ€“<scp>C</scp>hiastocheta interaction: insights into the evolution of nursery pollination systems. Ecology and Evolution, 2015, 5, 4766-4777. | 1.9 | 14 |
| 81 | Differential phenotypic and genetic expression of defence compounds in a plantâ€“herbivore interaction along elevation. Royal Society Open Science, 2016, 3, 160226. | 2.4 | 14 |
| 82 | Combining conservative and variable markers to infer the evolutionary history of Prunus subgen. Amygdalus s.l. under domestication. Genetic Resources and Crop Evolution, 2016, 63, 221-234. | 1.6 | 14 |
| 83 | Elevation in tropical sky islands as the common driver in structuring genes and communities of freshwater organisms. Scientific Reports, 2017, 7, 16089. | 3.3 | 14 |
| 84 | Decoupled postâ€glacial history in mutualistic plantâ€“insect interactions: insights from the yellow loosestrife (LysimachiaÂvulgaris) and its associated oilâ€collecting bees (MacropisÂeuropaea) Tj ETQq0 0 0 rgBT /Overloc | 3.0 | 13 |
| 85 | Genomics of extreme ecological specialists: multiple convergent evolution but no genetic divergence between ecotypes of Maculinea alcon butterflies. Scientific Reports, 2017, 7, 13752. | 3.3 | 13 |
| 86 | Spatial and temporal genetic dynamics of the grasshopper Oedaleus decorus revealed by museum genomics. Ecology and Evolution, 2018, 8, 1480-1495. | 1.9 | 13 |
| 87 | Biogeography and Ecological Diversification of a Mayfly Clade in New Guinea. Frontiers in Ecology and Evolution, 2019, 7, . | 2.2 | 13 |
| 88 | HyRAD-X Exome Capture Museomics Unravels Giant Ground Beetle Evolution. Genome Biology and Evolution, 2021, 13, . | 2.5 | 13 |
| 89 | Phylogeographic support for horizontal gene transfer involving sympatric bruchid species. Biology Direct, 2006, 1, 21. | 4.6 | 12 |
| 90 | Assessing the phylogenetic usefulness of a previously neglected morphological structure through elliptic Fourier analyses: a case study in Bruchus seedâ€beetles (Coleoptera: Chrysomelidae:) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5 | 3.3 | 12 |

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|---|---|---|---|
| 91 | Population genetic structure of two primary parasitoids of Spodoptera frugiperda (Lepidoptera), Chelonus insularis and Campoletis sonorensis (Hymenoptera): to what extent is the host plant important?. Molecular Ecology, 2010, 19, 2168-2179. | 3.9 | 12 |
| 92 | Ecological and historical drivers of diversification in the fly genus Chiastocheta Pokorny. Molecular Phylogenetics and Evolution, 2012, 63, 466-474. | 2.7 | 12 |
| 93 | Phylogenetic relationships in the subfamily Psychodinae (Diptera, Psychodidae). Zoologica Scripta, 2012, 41, 489-498. | 1.7 | 12 |
| 94 | Phylogeography and population genomics of a lotic water beetle across a complex tropical landscape. Molecular Ecology, 2018, 27, 3346-3356. | 3.9 | 12 |
| 95 | Spatial and temporal heterogeneity in pollinator communities maintains withinâ€species floral odour variation. Oikos, 2021, 130, 1487-1499. | 2.7 | 12 |
| 96 | Phylogeography ofChelonus insularis(Hymenoptera: Braconidae) andCampoletis sonorensis(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 |
| 97 | A deep digâ€“â€“hindsight on Holocene vegetation composition from ancient environmental <scp>DNA</scp>. Molecular Ecology, 2013, 22, 3433-3436. | 3.9 | 11 |
| 98 | Wheat alleles introgress into selfing wild relatives: empirical estimates from approximate Bayesian computation in Aegilops triuncialis. Molecular Ecology, 2014, 23, 5089-5101. | 3.9 | 11 |
| 99 | Congruent evolutionary responses of European steppe biota to late Quaternary climate change. Nature Communications, 2022, 13, 1921. | 12.8 | 11 |
| 100 | Insights into the genetic structure of the cowpea pest Callosobruchus maculatus in Africa. Journal of Pest Science, 2016, 89, 449-458. | 3.7 | 10 |
| 101 | Bacterial communities within Phengaris (Maculinea) alcon caterpillars are shifted following transition from solitary living to social parasitism of Myrmica ant colonies. Ecology and Evolution, 2019, 9, 4452-4464. | 1.9 | 10 |
| 102 | Rapid colour shift by reproductive character displacement in Cupido butterflies. Molecular Ecology, 2020, 29, 4942-4955. | 3.9 | 10 |
| 103 | Ecological niche overlap in sister species: how do oil-collecting bees Macropis europaea and Macropis fulvipes (Hymenoptera: Melittidae) avoid hybridization and competition?. Apidologie, 2011, 42, 579-595. | 2.0 | 9 |
| 104 | Does a shift in host plants trigger speciation in the Alpine leaf beetle Oreina speciosissima(Coleoptera,) Tj ETQq0 0 0 rgBT /Overlock 10 | 3.2 | 9 |
| 105 | Linking seascape with landscape genetics: Oceanic currents favour colonization across the GalÃ¡pagos Islands by a coastal plant. Journal of Biogeography, 2020, 47, 2622-2633. | 3.0 | 9 |
| 106 | Pollinators as drivers of plant distribution and assemblage into communities. , 2011, , 392-413. | | 8 |
| 107 | Isolation and characterization of polymorphic microsatellite loci in Acanthoscelides obvelatus Bridwell (Coleoptera: Bruchidae). Molecular Ecology Notes, 2002, 3, 12-14. | 1.7 | 8 |
| 108 | Evolutionary history and patterns of differentiation among European Maniola butterflies (Lepidoptera: Satyrinae). European Journal of Entomology, 2006, 103, 613-618. | 1.2 | 8 |

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|---|---|---|---|
| 109 | Genetic and Environmental Sources of Variation in the Autogenous Chemical Defense of a Leaf Beetle. Journal of Chemical Ecology, 2007, 33, 2011-2024. | 1.8 | 7 |
| 110 | Biotic drivers of river and floodplain geomorphology â€“ New molecular methods for assessing presentâ€day and past biota. Earth Surface Processes and Landforms, 2018, 43, 333-338. | 2.5 | 7 |
| 111 | Oviposition choice and larval development of the seed beetle Callosobruchus maculatus (F.) (Coleoptera: Chrysomelidae: Bruchinae) on three cowpea varieties. Journal of Stored Products Research, 2020, 86, 101578. | 2.6 | 7 |
| 112 | The untapped potential of macrofossils in ancient plant DNA research. New Phytologist, 2022, 235, 391-401. | 7.3 | 7 |
| 113 | SIMIL: an <scp>r</scp> (CRAN) scripts collection for computing genetic structure similarities based on <scp>structure</scp> 2 outputs. Molecular Ecology Resources, 2008, 8, 757-762. | 4.8 | 6 |
| 114 | Malagasy Dracaena Vand. ex L. (Ruscaceae): an investigation of discrepancies between morphological features and spatial genetic structure at a small evolutionary scale. Plant Systematics and Evolution, 2009, 280, 15-28. | 0.9 | 6 |
| 115 | High Rate of Protein Coding Sequence Evolution and Species Diversification in the Lycaenids. Frontiers in Ecology and Evolution, 2017, 5, . | 2.2 | 6 |
| 116 | Hybridization fuelled diversification in Spialia butterflies. Molecular Ecology, 2022, , . | 3.9 | 6 |
| 117 | Microsatellite markers in a complex of Horismenus sp. (Hymenoptera: Eulophidae), parasitoids of bruchid beetles. Molecular Ecology Notes, 2004, 4, 707-709. | 1.7 | 5 |
| 118 | Isolation and characterization of polymorphic microsatellite markers in Zabrotes subfasciatus Boheman (Coleoptera: Bruchidae). Molecular Ecology Notes, 2004, 4, 752-754. | 1.7 | 5 |
| 119 | The effect of host plant and isolation on the genetic structure of phytophagous insects: A preliminary study on a bruchid beetle. European Journal of Entomology, 2010, 107, 299-304. | 1.2 | 5 |
| 120 | DiscoSnp-RAD: de novo detection of small variants for RAD-Seq population genomics. PeerJ, 2020, 8, e9291. | 2.0 | 5 |
| 121 | Isolation and characterization of polymorphic microsatellite loci in Acanthoscelides obtectus Say (Coleoptera: Bruchidae). Molecular Ecology Notes, 2004, 4, 683-685. | 1.7 | 4 |
| 122 | Isolation and characterization of polymorphic microsatellite loci in two primary parasitoids of the noctuid Spodoptera frugiperda: Chelonus insularis and Campoletis sonorensis (Hymenoptera). Molecular Ecology Resources, 2009, 9, 171-173. | 4.8 | 4 |
| 123 | Genome fingerprinting confirms the species status of the Loosestrifes Lysimachia punctata and L. verticillata (Primulaceae). Plant Ecology and Evolution, 2016, 149, 335-338. | 0.7 | 0 |