

Manuel Curto

List of Publications by Year in descending order

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#	ARTICLE	IF	CITATIONS
1	Application of a SSR-GBS marker system on investigation of European Hedgehog species and their hybrid zone dynamics. <i>Ecology and Evolution</i> , 2019, 9, 2814-2832.	0.8	43
2	Application of microsatellite genotyping by sequencing (SSR-GBS) to measure genetic diversity of the East African <i>Oreochromis niloticus</i> . <i>Conservation Genetics</i> , 2019, 20, 357-372.	0.8	42
3	Metabarcoding Analyses of Gut Microbiota of Nile Tilapia (<i>Oreochromis niloticus</i>) from Lake Awassa and Lake Chamo, Ethiopia. <i>Microorganisms</i> , 2020, 8, 1040.	1.6	37
4	Molecular phylogenetics of <i>Micromeria</i> (Lamiaceae) in the Canary Islands, diversification and inter-island colonization patterns inferred from nuclear genes. <i>Molecular Phylogenetics and Evolution</i> , 2015, 89, 160-170.	1.2	31
5	Molecular genetic diversity and differentiation of Nile tilapia (<i>Oreochromis niloticus</i> , L. 1758) in East African natural and stocked populations. <i>BMC Evolutionary Biology</i> , 2020, 20, 16.	3.2	31
6	Development of phylogenetic markers from single-copy nuclear genes for multi locus, species level analyses in the mint family (Lamiaceae). <i>Molecular Phylogenetics and Evolution</i> , 2012, 63, 758-767.	1.2	29
7	Pollen availability for the Horned mason bee (<i>Osmia cornuta</i>) in regions of different land use and landscape structures. <i>Ecological Entomology</i> , 2020, 45, 525-537.	1.1	25
8	Genetic structure of <i>Micromeria</i> (Lamiaceae) in Tenerife, the imprint of geological history and hybridization on within-island diversification. <i>Ecology and Evolution</i> , 2016, 6, 3443-3460.	0.8	21
9	Genetic diversity and differentiation patterns in <i>Micromeria</i> from the Canary Islands are congruent with multiple colonization dynamics and the establishment of species syngameons. <i>BMC Evolutionary Biology</i> , 2017, 17, 198.	3.2	20
10	The Impact of Sampling Season and Catching Site (Wild and Aquaculture) on Gut Microbiota Composition and Diversity of Nile Tilapia (<i>Oreochromis niloticus</i>). <i>Biology</i> , 2021, 10, 180.	1.3	20
11	Evaluation of microsatellites of <i>Catha edulis</i> (qat; Celastraceae) identified using pyrosequencing. <i>Biochemical Systematics and Ecology</i> , 2013, 49, 1-9.	0.6	17
12	Analysis of microsatellite loci in tree of heaven (<i>Ailanthus altissima</i> (Mill.) Swingle) using SSR-GBS. <i>Tree Genetics and Genomes</i> , 2018, 14, 1.	0.6	17
13	Anthropogenic impacts on the contextual morphological diversification and adaptation of Nile tilapia (<i>Oreochromis niloticus</i> , L. 1758) in East Africa. <i>Environmental Biology of Fishes</i> , 2018, 101, 363-381.	0.4	16
14	The gut bacterial microbiome of Nile tilapia (<i>Oreochromis niloticus</i>) from lakes across an altitudinal gradient. <i>BMC Microbiology</i> , 2022, 22, 87.	1.3	16
15	Using a new RAD-sequencing approach to study the evolution of <i>Micromeria</i> in the Canary islands. <i>Molecular Phylogenetics and Evolution</i> , 2018, 119, 160-169.	1.2	15
16	Evidence for multiple introductions of an invasive wild bee species currently under rapid range expansion in Europe. <i>Bmc Ecology and Evolution</i> , 2021, 21, 17.	0.7	15
17	The influence of geological history on diversification in insular species: genetic and morphological patterns of <i>Micromeria</i> Benth. (Lamiaceae) in Tenerife (Canary archipelago). <i>Journal of Biogeography</i> , 2014, 41, 1871-1882.	1.4	12
18	Illumina midi-barcodes: quality proof and applications. <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2019, 30, 490-499.	0.7	10

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19	Population bottlenecks have shaped the genetic variation of <i>Ailanthus altissima</i> (Mill.) Swingle in an area of early introduction. <i>Forestry</i> , 2020, 93, 495-504.	1.2	10
20	A new amplicon based approach of whole mitogenome sequencing for phylogenetic and phylogeographic analysis: An example of East African white-eyes (<i>Aves</i> , <i>Zosteropidae</i>). <i>Molecular Phylogenetics and Evolution</i> , 2016, 102, 74-85.	1.2	9
21	Phylogeography of the wild and cultivated stimulant plant qat (<i>Catha edulis</i> , <i>Celastraceae</i>) in areas of historical cultivation. <i>American Journal of Botany</i> , 2017, 104, 538-549.	0.8	9
22	Application of microsatellite genotyping by amplicon sequencing for delimitation of African tilapiine species relevant for aquaculture. <i>Aquaculture</i> , 2021, 537, 736501.	1.7	8
23	Genetic characterization of fragmented populations of <i>Cinchona officinalis</i> L. (<i>Rubiaceae</i>), a threatened tree of the northern Andean cloud forests. <i>Tree Genetics and Genomes</i> , 2019, 15, 1.	0.6	7
24	Identifying geographically differentiated features of Ethiopian Nile tilapia (<i>Oreochromis niloticus</i>) morphology with machine learning. <i>PLoS ONE</i> , 2021, 16, e0249593.	1.1	6
25	Genetic diversity of Nile tilapia (<i>Oreochromis niloticus</i>) populations in Ethiopia: insights from nuclear DNA microsatellites and implications for conservation. <i>Bmc Ecology and Evolution</i> , 2021, 21, 113.	0.7	6
26	Isoflavone synthase (IFS) gene phylogeny in <i>Trifolium</i> species associated with plant isoflavone contents. <i>Plant Systematics and Evolution</i> , 2013, 299, 357-367.	0.3	5
27	Evaluating the Impact of Wildlife Shelter Management on the Genetic Diversity of <i>Erinaceus europaeus</i> and <i>E. roumanicus</i> in Their Contact Zone. <i>Animals</i> , 2020, 10, 1452.	1.0	5
28	Comparative Transcriptome Analysis of Two Root-Feeding Grape Phylloxera (<i>D. vitifoliae</i>) Lineages Feeding on a Rootstock and <i>V. vinifera</i> . <i>Insects</i> , 2020, 11, 691.	1.0	5
29	Influence of past agricultural fragmentation to the genetic structure of <i>Juniperus oxycedrus</i> in a Mediterranean landscape. <i>Tree Genetics and Genomes</i> , 2015, 11, 1.	0.6	4
30	New Microsatellite Markers for Two Sympatric Tinamou Species, the Ornate Tinamou (<i>Nothoprocta</i>) and the Overlock Tinamou (<i>Ortymis</i>). <i>Conservation Genetics</i> , 2014, 15, 1041-1048.	0.4	4
31	Knowledge Gaps in Taxonomy, Ecology, Population Distribution Drivers and Genetic Diversity of African Sandalwood (<i>Osyris lanceolata</i> Hochst. & Steud.): A Scoping Review for Conservation. <i>Plants</i> , 2021, 10, 1780.	1.6	4
32	Development and characterization of 16 microsatellite markers for <i>Micromeria</i> (<i>Lamiaceae</i>) from Tenerife (Canary Islands, Spain) using 454 sequencing. <i>Conservation Genetics Resources</i> , 2015, 7, 743-749.	0.4	3
33	Fifteen Microsatellite Markers for <i>Herbertia zebrina</i> (<i>Iridaceae</i>): An Endangered Species from South American Grasslands. <i>Applications in Plant Sciences</i> , 2017, 5, 1700035.	0.8	2
34	Investigating Shape Variation Using Generalized Procrustes Analysis and Machine Learning. <i>Applied Sciences (Switzerland)</i> , 2022, 12, 3158.	1.3	2
35	Four new mitochondrial genomes of the genus <i>Zosterops</i> (<i>Aves</i> : <i>Passeriformes</i> : <i>Zosteropidae</i>) from East Africa with a phylogenetic evaluation of the group. <i>Mitochondrial DNA Part B: Resources</i> , 2016, 1, 544-548.	0.2	1
36	Clonal Diversity, Cultivar Traits, Geographic Dispersal, and the Ethnotaxonomy of Cultivated Qat (<i>Catha edulis</i> , <i>Celastraceae</i>). <i>Economic Botany</i> , 2020, 74, 273-291.	0.8	0

#	ARTICLE	IF	CITATIONS
37	Development and characterization of microsatellite markers for two subspecies of <i>Handroanthus chrysanthus</i> . <i>Rodriguesia</i> , 0, 72, .	0.9	0