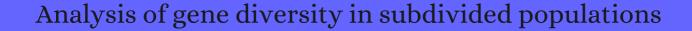
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	Genetic patterns of Magnolia in the Lesser Antilles: Stepwise colonisation leading to highly inbred	
122	Genetic patterns of Magnolia in the Lesser Antilles: Stepwise colonisation leading to highly inbred island populations. How ancient forest fragmentation and riparian connectivity generate high levels of genetic	0
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114	Genetic diversity, population structure, and clonal verification in South African avocado cultivars using single nucleotide polymorphism (SNP) markers. 2022 , 18,	0
113	Towards a genetic theory of island biogeography: Inferring processes from multidimensional community-scale data.	O
112	Detection of small sequence variations within the goat GHR gene and its effects on growth traits. 1-6	0
111	Population structure of sumac (Rhus coriaria L.) from TĒkiye based on transcriptome-developed SSR marker.	O
110	Genetic variability and pathogenicity of Fusarium verticillioides isolates from the summer-sown maize regions in China.	O
109	Multivariate analysis of chemical and genetic diversity of wild Humulus lupulus L. (hop) collected in situ in northern France. 2022 , 113508	O
108	Unexpected genetic integrity boosts hope for the conservation of the red-legged partridge (Alectoris rufa, Galliformes) in Italy. 2022 , 126056	O
107	Investigation of antimicrobial activities and molecular characterization of the species belong to Origanum, Thymus and Thymbra genera by ISSR.	1
106	Genetic divergence at species boundaries of the dolphinfish (Coryphaena hippurus) in the Tropical Eastern Pacific. 10, e14389	O
105	Diversity, genetic structure and core collection of mangaba (Hancornia speciosa) genebank.	O
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93	Hybrid purity identification using EST-SSR markers and heterosis analysis of quantitative traits of Russian wildrye. 10, e14442	1
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91	Molecular characterization and genetic diversity assessment in Mandukaparni (Centella asiatica L.) accessions.	O
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89	The assessment of genetic diversity and population structure of Verbascum gypsicola by ISSR markers for conservation purposes.	O
88	Population genetic structure of the globally introduced big-headed ant in Taiwan. 2022, 12,	О
87	Genetic diversity analysis of volunteer wheat based on microsatellite simple sequence repeats(SSR) markers.	O
86	Genetic Diversity, Population Structure and Subset Development in a Sesbania sesban Collection. 2023 , 12, 13	О
85	Diversity Analysis Using Agricultural Traits and SSR Markers of Wheat Germplasms. 2022 , 54, 345-357	O
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83	Virulence structure of wheat powdery mildew pathogen, Blumeria graminis tritici: a review.	O
82	Phylogeographic and morphological analysis ofBotrylloides nigerHerdman, 1886 from the northeastern Mediterranean Sea.	О
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66	A quantitative theory for genomic offset statistics.	Ο
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64	Genotyping by sequencing and a newly developed mRNA-GBS approach to link population genetic and transcriptome analyses reveal pattern differences between sites and treatments in red clover (Trifolium pratense L.). 10,	Ο
63	The Effect of the Genotypes of the CSN2 Gene on Test-Day Milk Yields in the Slovak Holstein Cow. 2023 , 13, 154	Ο
62	Molecular characterization of tomato (Solanum lycopersicum L.) landraces under drought stress.	Ο
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49	Not out of the Mediterranean: Atlantic populations of the gorgonian Paramuricea clavata are a separate sister species under further lineage diversification. 2023 , 13,	O
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47	Speciation and historical migration pattern interaction: examples from P. nigra and P. sylvestris phylogeography. 2023 , 142, 1-26	1
46	Genetic variability and phylogenetic relationship establishes distinctness of Kaunayen chicken of Manipur. 2017 , 87,	O
45	Analysis of Conservation Priorities and Runs of Homozygosity Patterns for Chinese Indigenous Chicken Breeds. 2023 , 13, 599	O
44	Inferring future changes in gene flow under climate change in riverscapes: a pilot case study in fluvial sculpin. 2023 , 38, 1351-1362	0
43	Genetic diversity and haplotype analysis of Leishmania tropica identified in sand fly vectors of the genera Phlebotomus and Sergentomyia using next-generation sequencing technology.	O

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41	The use of evolutionary analyses to predict functionally relevant traits in filamentous plant pathogens. 2023 , 73, 102244	O
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38	Analysis of Genetic Diversity, Population Structure and Association Mapping for Late Blight Resistance in Potato (Solanum tuberosum L.) Accessions Using SSR Markers. 2023 , 13, 294	0
37	An integrative study of species distribution modelling and conservation genetics: Magnolia in Hispaniola. 2023 , 32, 1205-1231	O
36	Genotyping of DNA pools identifies untapped landraces and genomic regions to develop next-generation varieties.	0
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34	Genetic variability and population structure of Agropyron desertorum accessions from Iran based on ISSR assay.	0
33	Genetic diversity and IUCN Red List status.	O
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31	Analysis of genetic diversity and population structure in sweetpotato using SSR markers. 2023,	O
30	Genetic Diversity of Five Galician (Northwestern Spain) Local Primitive Bovine Breeds Using Pedigree Records. 2023 , 15, 252	O
29	Quantum entropy reveals chromosomal disorder of ancestry tracts in genetic admixture.	O
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27	Amazonian birds in more dynamic habitats have less population genetic structure and higher gene flow.	O
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25	Great morphological diversity in wild apples (Malus spp.) from a region of central Spain (Guadarrama and Ayllfi mountain ranges).	O

24	The effects of irrigation canals and roads as barriers to gene flow between Japanese brown frog (Rana japonica) breeding sites at a fine scale in a Satoyama landscape. 2023 , 31, 191-202	O
23	Genetic Variation versus Morphological Variability in European Peatland Violets (Viola epipsilal). palustris Group). 2023 , 12, 362	O
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21	Molecular evidences for population differentiation and the migration from south to north of Puccinia triticina in eastern China. 2023 , 5,	O
20	The concordance of length- and sequence-based STRs used in forensic markers with guidance for practice.	O
19	Comparative population genetics of habitat-forming octocorals in two marine protected areas: eco-evolutionary and management implications.	O
18	Population Structure and Association Mapping for Agronomical and Biochemical Traits of a Large Spanish Apple Germplasm. 2023 , 12, 1249	0
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15	Development of a Core Collection for Tetraclinis articulata Using ISSR Markers and Maximization Strategy.	O
14	Association Analysis for Important Quantitative and Morphological Traits in Cultivars and Advanced Lines of Soybean (Glycine max (L.)) using Microsatellite Markers. 2022 , 14, 108-118	0
13	Populations of Saccharomyces cerevisiae in Vineyards: Biodiversity and Persistence Associated with Terroir. 2023 , 9, 292	O
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10	Indirect Estimation of Genetic Structure and Gene Flow in Some Black Poplar (Populus Nigra) Clones using Leaf Morphophysiological Traits. 2022 , 10, 117-126	0
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7	A reduced SNP panel optimised for non-invasive genetic assessment of a genetically impoverished conservation icon, the European bison.	O

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4	Genome-wide SNP analyses reveal a substantial gene flow and isolated-genetic structure of sea cucumber Holothuria leucospilota populations in Western Central Pacific. 2023 , 264, 106718	О
3	Virulence and genetic variability among Colletotrichum lindemuthianum (Sacc. & Magnus) Briosi & Magnus, Cavara isolates from common bean (Phaseolus vulgaris L.). 2023 , 37,	O
2	Phylogenetic relationships among five local sheep breeds of China. 2012 , 2012, 1-9	О
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