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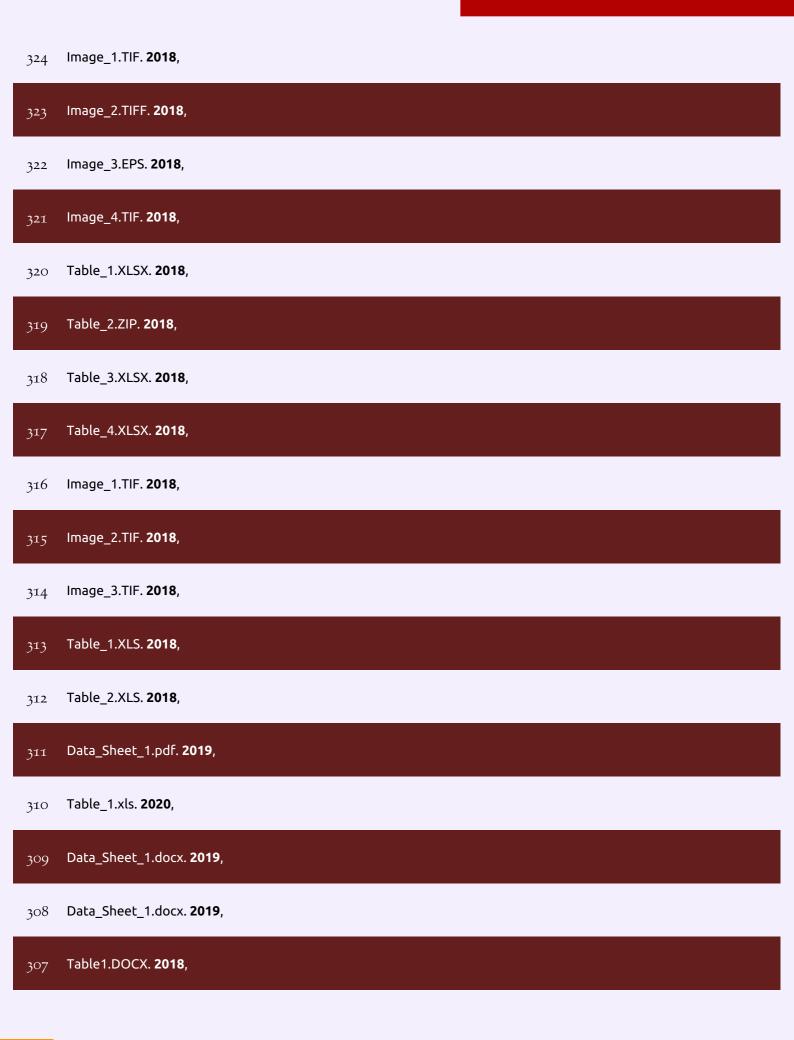
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143	Limited genetic differentiation of Mycetomoellerius mikromelanos in Parque National Soberana, Panama: Implications for queen dispersal.	O
142	Cyto-nuclear incompatibilities across a bluebell hybrid zone in northern Spain do not prevent genome-wide introgression of neutral markers.	O
141	Comparative phylogeography of a bathymetrically segregated pair of sister taxa of rockfishes (genus Sebastes): black rockfish, Sebastes melanops, and yellowtail rockfish, Sebastes flavidus.	O
140	Provenance of rhizobial symbionts is similar for invasive and non-invasive acacias introduced to California.	O
139	Molecular and quantitative genetic variation within and between populations of the declining grassland species Saxifraga granulata. 2022 , 12,	O
138	Effects of fifty years of shelterwood harvesting on genetic diversity and population structure of Oriental beech (Fagus orientalis L.) in the relict Hyrcanian forest. 2023 , 529, 120623	1
137	Regional environmental and climatic concerns on preserving native gene pools of a least concern species: Brown trout lineages in Mediterranean streams. 2023 , 862, 160739	O
136	Genetic variation and relationships among spring camelina (Camelina sativa, Brassicaceae) accessions of different origin. 2022 , 59, 86-90	0
135	Biotic and abiotic factors affecting the microbiota of Chrysomelidae inhabiting wetland vegetation.	O
134	Allopatric Lineage Divergence of the East Asian Endemic Herb Conandron ramondioides Inferred from Low-Copy Nuclear and Plastid Markers. 2022 , 23, 14932	О
133	First evidence of population genetic structure of the deep-water blackmouth catshark Galeus melastomus Rafinesque, 1810. 9,	O
132	Restoration of calcareous grasslands by natural recolonization after forest clearing and its impact on the genetic variation of three common herb species.	O
131	Genetic study of the American horseshoe crab throughout its Mexican distribution. Conservation and management implications.	O
130	Genetic diversity and structure of Bipolaris oryzae and Exserohilum rostratum populations causing brown spot of rice in Burkina Faso based on genotyping-by-sequencing. 13,	0
129	Genetic characterization of oleaginous bottle gourd (Lagenaria siceraria) germplasm from Cle d'Ivoire using agromorphological and molecular markers. 1-12	O
128	Phytochemicals and Their Correlation with Molecular Data in Micromeria and Clinopodium (Lamiaceae) Taxa. 2022 , 11, 3407	1
127	Table Mountain Pine (Pinus pungens): Genetic Diversity and Conservation of an Imperiled Conifer.	O

126	Rear-edge daylily populations show legacies of habitat fragmentation due to the Holocene climate warming.	O
125	Three mitochondrial lineages and no Atlantic-Mediterranean barrier for the bogue Boops boops across its widespread distribution. 2022 , 12,	O
124	Genetic diversity of Mimosa tenuiflora (Willd.) Poir.: an intensively exploited wood tree in the Brazilian tropical semi-arid vegetation.	0
123	Panmictic stock structure of milkfish (Chanos´chanos, Forsskl1775) from Indian waters determined using mtDNA marker. 2023 , 102,	O
122	Population genetic structure of the globally introduced big-headed ant in Taiwan. 2022, 12,	0
121	Commerson dolphin population structure: evidence for female phylopatry and male dispersal. 2022 , 12,	O
120	Fencing Can Alter Gene Flow of Asian Elephant Populations within Protected Areas. 2022, 2, 709-725	1
119	Genetic Variation and Population Structure of Oryza sativa Accessions in the AfricaRice Collection and Development of the AfricaRice O. sativa Core Collection.	O
118	Multiple introductions, polyploidy and mixed reproductive strategies are linked to genetic diversity and structure in the most widespread invasive plant across Southern Ocean archipelagos.	0
117	Has insecticidal pressure influenced Spodoptera litura (Fabricius, 1775) population genetic structure and genetic diversity in India?.	О
116	The Quality of Sequence Data Affects Biodiversity and Conservation Perspectives in the Neotropical Damselfly Megaloprepus caerulatus. 2022 , 14, 1056	0
115	Genetic Diversity of Black Amaranth (Amaranthus quitensis Kunth) Landraces of Ecuadorian Highlands: Association Genotypes (Iolor Morphotypes. 2023 , 13, 34	O
114	Population structure and genetic connectivity of the scalloped hammerhead shark (Sphyrna lewini) across nursery grounds from the Eastern Tropical Pacific: Implications for management and conservation. 2022 , 17, e0264879	0
113	Legacies of domestication, Neolithic diffusion and trade between Indian subcontinent and Island Southeast Asia shape maternal genetic diversity of Andaman cattle. 2022 , 17, e0278681	O
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111	Stepping up to genome scan allows stock differentiation in the worldwide distributed blue shark Prionace glauca.	1
110	SSR Genotypes of the Puccinia triticina in 15 Provinces of China Indicate Regional Migration in One Season from East to West and South to North. 2022 , 12, 3068	O
109	New insights on patterns of genetic admixture and phylogeographic history in Iberian high mountain populations of midwife toads. 2022 , 17, e0277298	О

108	Spatial and Temporal Patterns of Genetic Diversity and Structure in Danish Populations of the Alcon Blue Butterfly Phengaris alcon (Denis & Schifferm Ler). 2022 , 14, 1098	О
107	Riverscape community genomics: A comparative analytical approach to identify common drivers of spatial structure.	О
106	Phylogenetic Analysis and Genetic Structure of Schlegel Japanese Gecko (Gekko japonicus) from China Based on Mitochondrial DNA Sequences. 2023 , 14, 18	О
105	Genotypic variation, phylogeography, unified species concept, and the grey zonelbf taxonomic uncertainty in kliuka: recognition of Kunzea ericoides (A.Rich.) Joy Thomps. sens. lat. (Myrtaceae). 1-30	O
104	Population Subdivision in the Gopher Frog (Rana capito) across the Fragmented Longleaf Pine-Wiregrass Savanna of the Southeastern USA. 2023 , 15, 93	O
103	Multidrug-Resistant ESBL-Producing E. coli in Clinical Samples from the UK. 2023 , 12, 169	1
102	High-Resolution Population Genetic Structure of Tawny Crazy Ant (Nylanderia fulva Mayr: Hymenoptera: Formicidae) from the Origin in South America and Introduced Regions of the United States.	0
101	Range-wide population genomics of the spongy moth, Lymantria dispar (Erebidae): Implications for biosurveillance, subspecies classification and phylogeography of a destructive moth.	1
100	Galacto-Oligosaccharides Increase the Abundance of Beneficial Probiotic Bacteria and Improve Gut Architecture and Goblet Cell Expression in Poorly Performing Piglets, but Not Performance. 2023 , 13, 230	О
99	Small-scale genetic structure of coral populations in Palau based on whole mitochondrial genomes: Implications for future coral resilience.	O
98	Comparative Analysis of the Genetic Diversity of Faba Bean (Vicia faba L.). 2023, 15, 1016	1
97	Genetic variation of Amaranthus retroflexus L. and Chenopodium album L. (Amaranthaceae) suggests multiple independent introductions into Iran. 13,	O
96	Mitochondrial haplotypes reveal low diversity and restricted connectivity in the critically endangered batoid population of a Marine Protected Area.	0
95	Analysis of the Genetic Diversity of Two Rhopalosiphum Species from China and Europe Based on Nuclear and Mitochondrial Genes. 2023 , 14, 57	0
94	Population genetics of the African snakehead fish Parachanna obscura along West Africa's water networks: Implications for sustainable management and conservation. 2023 , 13,	О
93	Effects of Wild Forest Fires on Genetic Diversity and Population Structure of a Boreal Conifer, White Spruce (Picea glauca (Moench) Voss): Implications for Genetic Resource Management and Adaptive Potential under Climate Change. 2023 , 14, 157	O
92	Past climate changes and geographical barriers: the evolutionary history of a subtropical highland grassland species of Solanaceae, Petunia altiplana.	0
91	Outlier analyses and genome-wide association study identify glgC and ERD6-like 4 as candidate genes for foliar water-soluble carbohydrate accumulation in Trifolium repens. 13,	O

90	Another Chapter in the History of the European Invasion by the Western Conifer Seed Bug, Leptoglossus occidentalis: The Iberian Peninsula. 2023 , 15, 64	0
89	Population Structure and Genetic Diversity of Colletotrichum gloeosporioides on Citrus in China. 2023 , 13, 184	O
88	Are populations less genetically diverse and more differentiated at the species range edges? Analysis of the quantitative trait and molecular variation in wild oat Avena sterilis. 2022 , 1-9	O
87	QTL for yield per plant under water deficit and well-watered conditions and drought susceptibility index in soybean (Glycine max (L.) Merr.). 2023 , 37, 92-103	O
86	Coalescent-Based Species Delimitation in Herbaceous Bamboos (Bambusoideae, Olyreae) from Eastern Brazil: Implications for Taxonomy and Conservation in a Group with Weak Morphological Divergence Coupled with Low Genetic Diversity. 2023 , 12, 107	O
85	Genetic diversity among turf grasses by ISSR markers. 2017 , 87,	O
84	Morphological characterization and genetic diversity of a new microsporidium, Neoflabelliforma dubium n. sp. from the adipose tissue of Diaphanosoma dubium (Crustacea: Sididae). 13,	O
83	Searching for Novel Oat Crown Rust Resistance in Diploid Oat Avena strigosa Schreb. Reveals the Complexity and Heterogeneity of the Analyzed Genebank Accessions. 2023 , 13, 296	O
82	Population genetic research on the Spadenose shark, Scoliodon laticaudus (Chondrichthyes, Carcharhinidae), reveals the presence of two significantly differentiated clades along the Indian coast. 2023 , 53,	0
81	Temporal and Spatial Variation in the Population Structure of Spanish Fusarium circinatum Infecting Pine Stands. 2023 , 9, 159	O
80	No loss of genetic divergence in Torymus sinensis following its release for Dryocosmus kuriphilus control in Europe.	0
79	ISSR-assisted spatial genetic structure, population admixture, and biodiversity estimates across locally adopted saffron ecotypes from 18 different provenances of Iran. 2023 , 100467	O
78	Not out of the Mediterranean: Atlantic populations of the gorgonian Paramuricea clavata are a separate sister species under further lineage diversification. 2023 , 13,	O
77	Pathometagenomics reveals susceptibility to intestinal infection by Morganella to be mediated by the blood group-related B4galnt2 gene in wild mice. 2023 , 15,	O
76	The comparison of polymorphism among Avena species revealed by retrotransposon-based DNA markers and soluble carbohydrates in seeds.	0
75	Circumpolar sampling reveals high genetic connectivity of Antarctic toothfish across their spatial distribution.	O
74	Genetic Diversity of Jinshaia sinensis (Cypriniformes, Balitoridae) Distributed Upstream of the Yangtze River. 2023 , 8, 75	O
73	Bacterial Contamination of the Surgical Site at the Time of Elective Caesarean Section in Belgian Blue Cows P art 2: Identified by 16Sr DNA Amplicon Sequencing. 2023 , 10, 94	O

72	Analyzing Autopolyploid Genetic Data Using GenoDive. 2023, 261-277	0
71	Speciation and historical migration pattern interaction: examples from P. nigra and P. sylvestris phylogeography. 2023 , 142, 1-26	1
70	Investigation of B-atp6-orfH79 distributing in Chinese populations of Oryza rufipogon and analysis of its chimeric structure. 2023 , 23,	0
69	Rivers as corridors for gene flow of the neotropical bamboo Guadua velutina. 2023 , 186, 103632	O
68	Discovery and Genetic Characterization of Single Cohort Adult Colonies With Male Aggregations, and Preliminary Evidence for Lekking in a Malagasy Kite Spider (Isoxya, Gasteracanthinae). 2023 , 7,	O
67	Genetic variation and cryptic lineage among the sergestid shrimp Acetes americanus (Decapoda). 11, e14751	O
66	Chloroplast DNA reveals genetic population structure in Sinomenium acutum in subtropical China. 2023 ,	0
65	Individualized network analysis: A novel approach to investigate tau PET using graph theory in the Alzheimer disease continuum. 17,	O
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63	Population structure and genetic diversity of tobacco mild green mosaic virus variants in Western Anatolia of Turkey. 2023 , 125, 102008	O
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61	Fine scale genetics reveals the subtle negative effects of roads on an endangered bat. 2023 , 869, 161705	O
60	Genetic population structure in Norway lobster (Nephrops norvegicus): management regime under panmixia.	0
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58	Untangling the Hymenaea stigonocarpa complex (Fabaceae) using population genetics and morphology.	О
57	Intraspecific differentiation in genetic structure in Panicum coloratum L.: importance for germplasm conservation and breeding.	O
56	Genetic Structure and Phylogeography of Commercial Mytilus unguiculatus in China Based on Mitochondrial COI and Cytb Sequences. 2023 , 8, 89	0
55	Strong Genetic Structure and Limited Gene Flow among Populations of the Tropical Seagrass Thalassia hemprichii in the Philippines. 2023 , 11, 356	O

54	Galacto-oligosaccharides fed during gestation increase Rotavirus A specific antibodies in sow colostrum, modulate the microbiome, and reduce infectivity in neonatal piglets in a commercial farm setting. 10,	Ο
53	Structure and Functional Properties of Bacterial Communities in Surface Sediments of the Recently Declared Nutrient-Saturated Lake Villarrica in Southern Chile.	Ο
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51	The blowfly Chrysomya latifrons inhabits fragmented rainforests, but shows no population structure. 2023 , 201, 703-719	0
50	Neutral and adaptive genetic diversity in plants: An overview. 11,	0
49	Molecular biology tools used to study hominin evolution. 2023 , 219-279	O
48	Genetic and Metabolite Variability among Commercial Varieties and Advanced Lines of Vicia faba L 2023 , 12, 908	0
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46	Thermal heterogeneity is an important factor for maintaining the genetic differentiation pattern of the pelagic barnacle Lepas anatifera in the northwest Pacific. 2023 , 13,	0
45	Genomic insights into the NPGS intermediate wheatgrass germplasm collection.	O
45	Genomic insights into the NPGS intermediate wheatgrass germplasm collection. Weak range-wide population structure in the blackfin tuna (Thunnus atlanticus) revealed by analysis of genome-wide SNPs.	0
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34	A Comparison of Entropic Diversity and Variance in the Study of Population Structure. 2023 , 25, 492	O
33	Genetic diversity of longjack (Eurycoma longifolia Jack) distributed in Thua Thien Hue, Vietnam.	O
32	Genetic Evidence for Indo-Western Pacific Olive Ridley Sea Turtles in Mexican Waters. 2023 , 15, 430	O
31	Landscape configurations determining the genetic structure of the Yellow-Spotted Amazon River Turtle (Podocnemis unifilis) in Brazilian Amazonia.	O
30	Genetic diversity assessment of the indigenous goat population of Benin using microsatellite markers. 14,	О
29	On the concepts and measures of diversity in the genomics era. 2023 , 33, 100278	O
28	Flushing Temporarily Improves Microbiological Water Quality for Buildings Supplied with Chloraminated Surface Water but Has Little Effect for Groundwater Supplies. 2023 , 57, 5453-5463	O
27	Floral and genetic divergence across environmental gradients is moderated by inter-population gene flow in Platanthera dilatata (Orchidaceae). 11,	O
26	To all the gar I loved before: range-wide population genetic structure in Alligator gar.	О
25	Peeping into Mitochondrial Diversity of Andaman Goats: Unveils Possibility of Maritime Transport with Diversified Geographic Signaling. 2023 , 14, 784	O
24	Genotypic characterization of Phytophthora infestans populations in Bangladesh.	O
23	Morphological and Molecular Characterization of Micromeria croatica (Lamiaceae), an Endemic and Potentially Valuable Horticultural Species of the Dinaric Alps. 2023 , 9, 418	O
22	Update on the genetic diversity and population structure of Echinococcus granulosus in Gansu Province, Tibet Autonomous Region, and Xinjiang Uygur Autonomous Region, Western China, inferred from mitochondrial cox1, nad1, and nad5 sequences. 2023 , 122, 1107-1126	О
21	Genetic Diversity and Population Structure of Ugandan Soybean (Glycine max L.) Germplasm Based on DArTseq.	1
20	Molecular and Technological Characterization of Saccharomyces´cerevisiae from Sourdough. 2023 , 9, 329	О
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11	A genogeographic study of the Kyrgyz mountain merino via microsatellite markers. 2023 , 27, 162-168	0
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8	Sponge diversification in marine lakes: Implications for phylogeography and population genomic studies on sponges. 2023 , 13,	0
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5	Population genetics comparison of Lilioceris cheni (Coleoptera: Chrysomelidae) colonies released onto Dioscorea bulbifera in Southeastern U.S.A 1-19	0
4	Conservation and sustainable development of coastal species of horticultural importance: insights from genetic and environmental patterns at spatio-temporal scale.	O
3	Mitochondrial Heteroplasmy and PCR Amplification Bias Lead to Wrong Species Delimitation with High Confidence in the South American and Antarctic Marine Bivalve Aequiyoldia eightsii Species Complex. 2023 , 14, 935	O
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1	Attack of the clones: Population genetics reveals clonality of Colletotrichum lupini , the causal agent of lupin anthracnose.	0