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1284	Population genetic dynamics of three-spined sticklebacks (Gasterosteus aculeatus) in anthropogenic altered habitats. 2012 , 2, 1122-43	8
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1261	The first microsatellite markers for little terns (Sternula albifrons). 2012 , 4, 447-450	
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1259	Isolation and characterization of 20 microsatellite loci in Neverita didyma (Raing 1798). 2012 , 4, 479-481	4
1258	Characterization of 17 microsatellite loci for use in population genetic and mating system studies of the endangered North American passerine, loggerhead shrike (Lanius ludovicianus). 2012 , 4, 503-506	3
1257	Development and characterization of microsatellite markers for the seagrass Halodule wrightii. 2012 , 4, 511-513	6
1256	Development and characterization of tetranucleotide microsatellite loci for the American alligator (Alligator mississippiensis). 2012 , 4, 567-570	4
1255	Development of eight microsatellite loci from the endangered huemul (Hippocamelus bisulcus) and cross-species amplification in six other ungulate species. 2012 , 4, 571-573	1
1254	Isolation and characterization of ten microsatellite loci of endangered Anderson crocodile newt, Echinotriton andersoni. 2012 , 4, 595-598	4
1253	Characterization of blue tit Cyanistes caeruleus microsatellite loci, and assessment of sequence homology among species to predict cross-species utility and chromosome locations. 2012 , 4, 605-612	
1252	Isolation and characterization of microsatellite markers for the marine black nerite Nerita atramentosa: tools for assessment and design of marine protected areas. 2012 , 4, 625-627	4
1251	Isolation and characterisation of microsatellite markers from the South Island robin (Petroica australis). 2012 , 4, 633-636	6

1250	Microsatellites development for rock carp (Procypris rabaudi) and cross-species testing in Onychostoma sima and Spinibarbus sinensis. 2012 , 4, 657-660	1
1249	A reliable method for individual identification and gender determination of wild leopards (Panthera pardus fusca) using non-invasive samples. 2012 , 4, 665-667	14
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1243	Identification and evaluation of novel di- and tetranucleotide microsatellite markers from the brown bear (Ursus arctos). 2012 , 4, 737-741	5
1242	Isolation and characterization of eleven novel polymorphic microsatellite loci in the spiny softshell turtle (Apalone spinifera). 2012 , 4, 759-761	2
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1239	Isolation and characterization of microsatellite loci in the Reevesell butterfly lizard Leiolepis reevesii (Agamidae). 2012 , 4, 791-794	6
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1236	Population structure and genetic diversity of the orchid bee Eufriesea violacea (Hymenoptera, Apidae, Euglossini) from Atlantic Forest remnants in southern and southeastern Brazil. 2012 , 43, 392-402	17
1235	High genetic differentiation in populations of the rare alpine plant species Campanula thyrsoides on a small mountain. 2012 , 122, 23-34	14
1234	Colony structure and reproductive sharing among queens in a tropical paper wasp, Polistes olivaceus. 2012 , 59, 401-410	3
1233	Dramatic shifts in the gene pool of a managed population of an endangered species may be exacerbated by high genetic load. 2012 , 13, 349-358	21

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1224	Low genetic diversity and evidence of population structure among subspecies of Nerodia harteri, a threatened water snake endemic to Texas. 2012 , 13, 977-986	6
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1221	Fine-scale effects of habitat loss and fragmentation despite large-scale gene flow for some regionally declining woodland bird species. 2012 , 27, 813-827	56
1220	Genetic Diversity and Characterization of a Core Collection of Malus Germplasm Using Simple Sequence Repeats (SSRs). 2012 , 30, 827-837	44
1219	Isolation and characterisation of nine microsatellite markers in the boring giant clam (Tridacna crocea) and cross-amplification in five other tridacnid species. 2012 , 42, 285-287	3
1218	Development of eleven polymorphic microsatellite loci for the sea snake Emydocephalus annulatus (Elapidae: Hydrophiinae) and cross-species amplification for seven species in the sister genus Aipysurus. 2012 , 4, 11-14	5
1217	Isolation and PCR-multiplex genotyping of 18 novel microsatellite markers for the threatened southern pygmy perch (Nannoperca australis). 2012 , 4, 15-17	7
1216	Characterization of 11 microsatellite loci for the brown smooth-hound shark, Mustelus henlei (Triakidae), discovered with next-generation sequencing. 2012 , 4, 23-25	17
1215	Development of tetranucleotide microsatellite loci and a non-invasive DNA sampling method for Texas horned lizards (Phrynosoma cornutum). 2012 , 4, 43-45	4

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1207	Characterization of 12 microsatellite loci in the waterfall damselfly (Paraphlebia zoe) for use in population genetic applications. 2012 , 4, 175-177	1
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1006	Isolation of microsatellite loci for the predatory fish Hoplias microlepis (Characiformes: Erythrinidae) from a highly impacted river system in western Ecuador. 2013 , 5, 437-439	4
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1004	Isolation and characterization of EST-based microsatellite markers for Scatophagus argus based on transcriptome analysis. 2013 , 5, 483-485	6
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1000	Development of 35 novel microsatellite markers for the two-band anemonefish Amphiprion bicinctus. 2013 , 5, 515-518	5
999	Large-scale isolation of Eastern spiny mouse Acomys dimidiatus microsatellite loci through GS-FLX 454 titanium sequencing. 2013 , 5, 519-524	1

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986	Characterisation of microsatellite markers for the granite endemic Kunzea pulchella (Lindl.) A. S. George (Myrtaceae) identified using next generation sequencing. 2013 , 5, 129-131	2
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984	Development and characterization of 20 polymorphic microsatellite markers for the Texas hornshell, Popenaias popeii (Bivalvia: Unionidae), through next-generation sequencing. 2013 , 5, 195-198	5
983	Microsatellite markers for vulnerable Australian aridzone Acacias. 2013 , 5, 199-201	7
982	Isolation and characterization of nine microsatellite loci in a Malagasy endemic rodent, Eliurus carletoni (Rodentia: Nesomyinae). 2013 , 5, 203-205	
981	Isolation of nuclear microsatellite loci in the African tree Scorodophloeus zenkeri (Fabaceae). 2013 , 5, 219-221	1

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977	Development of microsatellite loci for the western slimy salamander (Plethodon albagula) using 454 sequencing. 2013 , 5, 267-270	6
976	Isolation and characterization of 11 microsatellite loci in Schizothorax wangchiachii (Fang). 2013 , 5, 271-273	
975	Assignment test method using hypervariable markers for blue marlin (Makaira nigricans) stock identification. 2013 , 5, 293-297	7
974	Genetic structure, relationships and admixture with wild relatives in native pig breeds from Iberia and its islands. 2013 , 45, 18	25
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967	Association between loss-of-function mutations in the filaggrin gene and self-reported food allergy and alcohol sensitivity. 2013 , 161, 234-42	19
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957	Stock Origins of Subadult and Adult Atlantic Sturgeon, Acipenser oxyrinchus, in a Non-natal Estuary, Long Island Sound. 2013 , 36, 257-267	23
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932	Contemporary genetic structure reflects historical drainage isolation in an Australian snapping turtle, Elseya albagula. 2013 , 169, 200-214	9
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917	Population Genetics and Bat Rabies: A Case Study of Eptesicus serotinus in Poland. 2013 , 15, 35-56	13
916	Urban population genetics of slum-dwelling rats (Rattus norvegicus) in Salvador, Brazil. 2013 , 22, 5056-70	39
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910	Limited genetic differentiation between acoustically divergent populations of urban and rural silvereyes (Zosterops lateralis). 2013 , 27, 381-391	7
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900	Patterns of genetic differentiation at MHC class I genes and microsatellites identify conservation units in the giant panda. 2013 , 13, 227	22
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244	Fine-scale social and genetic structure of common bottlenose dolphins (Tursiops truncatus) in the Barataria Basin, Louisiana, USA.	
243	Contrasting mtDNA and microsatellite data of great reed warbler Acrocephalus arundinaceus breeding populations on a small geographic scale.	

242	Genetic variability and population structure of the Japanese sea cucumber, Apostichopus japonicus Selenka, 1867 revealed by microsatellites in Peter the Great Gulf, Sea of Japan. 2022 , 52,	0
241	Rapid evolution of selfing syndrome traits in Viola arvensis revealed by resurrection ecology.	1
240	Traces of past reintroduction in genetic diversity: The case of the Balkan chamois (Mammalia, Artiodactyla). 1116, 57-70	0
239	Genetic Diversity and Fine-Scale Spatial Genetic Structure of the Endangered Shrub Birch (Betula humilis Schrk.) Populations in Protected and Unprotected Areas. 2022 , 14, 684	
238	Genetic Analyses Reveal High Connectivity among Populations of the Honduran White Bat Ectophylla alba in the Caribbean Lowlands of Central Eastern Costa Rica. 2022 , 24,	
237	Collection of native Theobroma cacao L. accessions from the Ecuadorian Amazon highlights a hotspot of cocoa diversity.	O
236	New Microsatellite Markers for Genetic Studies on Sergentomyia schwetzi (Diptera: Psychodidae): A Suspected Vector of Leishmania infantum (Trypanosomatida: Trypanosomatidae) in the Canine Leishmaniasis Focus of Mont-Rolland, Senegal.	О
235	Genetic Analysis and Status of Brown Bear Sub-Populations in Three National Parks of Greece Functioning as Strongholds for the Species Conservation. 2022 , 13, 1388	
234	Recent declines in genetic diversity with limited dispersal among coastal cactus wren populations in San Diego County, California.	
233	Early-succession secondary forests following agropastoral abandonment are key winter habitats for the conservation of a priority bird in the European Alps.	
232	A new set of microsatellite markers for Phoxinus lumaireul senso lato , P. marsilii and P. krkae for population and molecular taxonomic studies.	
231	Evidence of an additional center of apple domestication in Iran, with contributions from the Caucasian crab apple Malus orientalis Uglitzk. to the cultivated apple gene pool.	
230	Contemporary genetic structure of Xantus's Hummingbird (Basilinna xantusii) in the Baja California peninsula.	
229	Invasive genetic rescue: Dispersal following repeated culling reinforces the genetic diversity of an invasive mammal.	
228	Ancient DNA reveals phenological diversity of Coast Salish herring harvests over multiple centuries. 2022 , 12,	
227	Nuclear genetic diversity and structure of Anastrepha ludens wild populations evidenced by microsatellite markers. 10,	
226	A conservation genetics perspective on supportive breeding: A case study of the common nase (Chondrostoma nasus).	1
225	Genetic diversity of Pinus kesiya in the Central Highlands of Vietnam.	O

224	Comparison of genetic variation between rare and common congeners of Dipodomys with estimates of contemporary and historical effective population size. 2022 , 17, e0274554	O
223	Genetic Evaluation in Natural Populations of the Threatened Conifer Amentotaxus argotaenia (Hance) Pilg. (Taxaceae) Using Microsatellites. 2022 , 13, 1452	o
222	A multifaceted index of population health to detect risk-prone populations and underlying stressors in wildlife. 2022 , 274, 109706	0
221	Female demographic changes contribute to the maintenance of social stability within a primate multilevel society. 2022 , 192, 101-108	o
220	From poops to planning: A broad non-invasive genetic survey of large mammals from the Indian Himalayan Region. 2022 , 853, 158679	0
219	Population structure of Caspian Kutum (Rutilus frisii, Nordmann, 1840) in the southern coast of Caspian Sea using Genome-wide single nucleotide polymorphism markers. 2023 , 257, 106499	O
218	Genetic Diversity and Fine-Scale Spatial Genetic Structure of Unmanaged Old-Growth Versus Managed Second-Growth Populations of Scots Pine (Pinus Sylvestris L.) in Lithuania: Effects of Systematic Tending.	O
217	Inferred family structure and dispersal patterns of a critically endangered species, Pinna nobilis, using molecular analyses: implications for conservation.	o
216	Genomic Data Reveals Profound Genetic Structure and Multiple Glacial Refugia in Lonicera oblata (Caprifoliaceae), a Threatened Montane Shrub Endemic to North China. 13,	0
215	Palaeoclimatic changes resulted in range expansion and subsequent divergence in brown fur seals, Arctocephalus pusillus. 2022 , 18,	O
214	Genetic variability of indigenous (Quercus robur L.) and late flushing oak (Quercus robur L. subsp. slavonica (G͡yer) MĒyĒ) in adult stands compared with their natural regeneration.	0
213	Characterising the molecular diversity of ash (Fraxinus excelsior L.) at its western marginal range in Europe [phylogeographic insights and implications for conservation in Ireland. 2022 , 18,	o
212	Genome architecture and diverged selection shaping pattern of genomic differentiation in wild barley.	0
211	Fine-scale genetic structure and wolbachia infection of aedes albopictus (Diptera: Culicidae) in Nanjing city, China. 13,	o
210	Development and characterization of microsatellite markers for Chaeturichthys stigmatias (Actinopterygii: Gobiiformes: Gobiidae) based on restriction site-associated DNA sequencing (RAD-seq). 2022 , 52, 229-237	0
209	Phylogeographic structure of common sage (Salvia officinalis L.) reveals microrefugia throughout the Balkans and colonizations of the Apennines. 2022 , 12,	o
208	Systematic selections and forensic application evaluations of 111 individual identification SNPs in the Chinese Inner Mongolia Manchu group. 13,	0
207	Population history, genetic variation, and conservation status of European white elm (Ulmus laevis Pall.) in Poland. 2022 , 79,	o

206	Microsatellite DNA analysis revealed increased homozygosity and recent decline of the population size in the European flounder (Platichthys flesus) from the southern Baltic Sea.	0
205	The genetic diversity and structure in the European polecat were not affected by the introduction of the American mink in Poland. 2022 , 17, e0266161	O
204	The effect of habitat and climatic on microsatellite diversity and allele length variation. 10,	Ο
203	Monitoring genetic diversity with new indicators applied to an alpine freshwater top predator.	О
202	New strategies for characterizing genetic structure in wide-ranging, continuously distributed species: A Greater Sage-grouse case study. 2022 , 17, e0274189	1
201	Assessment of connectivity patterns of the marbled crab Pachygrapsus marmoratus in the Adriatic and Ionian seas through combination of genetic data and Lagrangian simulations. 9,	O
200	Species and population genomic differentiation in Pocillopora corals (Cnidaria, Hexacorallia). 2022 , 150, 247-262	0
199	The invasive alga Gracilaria vermiculophylla in the native northwest Pacific under ocean warming: Southern genetic consequence and northern range expansion. 9,	1
198	Using genetic tools to inform conservation of fragmented populations of Asian elephants (Elephas maximus) across their range in China.	0
197	Evaluation of genetic diversity of Himalayan balsam (Impatiens glandulifera Royle) populations using microsatellites. 2022 , 109, 259-268	O
196	Giant kelp genetic monitoring before and after disturbance reveals stable genetic diversity in Southern California. 9,	0
195	A baseline for the genetic stock identification of Atlantic herring, Clupea harengus, in ICES Divisions 6.a, 7.bl. 2022 , 9,	O
194	High levels of genetic diversity and population structure in the Mediterranean seagrass Posidonia oceanica at its easternmost distribution limit.	0
193	Phylogeny and Population Genetic Structure of Minke Whales Worldwide: A Review of Recent Studies.	O
192	Demographic resilience of brook trout populations subjected to experimental size-selective harvesting.	0
191	New microsatellite markers for the shallow coral Madracis auretenra from the Caribbean. 2022 , 17, e0274895	0
190	Rapid SNP genotyping, sex identification, and hybrid-detection in threatened bull trout.	0
189	The shrunk genetic diversity of coral populations in North-Central Patagonia calls for management and conservation plans for marine resources. 2022 , 12,	O

188	Phylogeographic breaks and how to find them: Separating vicariance from isolation by distance in a lizard with restricted dispersal.	0
187	The extent of hybridisation between largemouth bass and Florida bass across two river systems in South Africa. 1-13	o
186	Isolation and characterization of 50 microsatellite loci for two shearwater species, Ardenna pacifica and Puffinus bailloni.	O
185	Multiple mating by both sexes in an invasive insect species, Aethina tumida (Coleoptera: Nitidulidae).	o
184	A comprehensive assessment of insecticide resistance mutations in source and immigrant populations of the diamondback moth Plutella xylostella (L.).	0
183	Unraveling the hierarchical genetic structure of tea green leafhopper, Matsumurasca onukii , in East Asia based on SSRs and SNPs. 2022 , 12,	O
182	Climate change threatens unique genetic diversity within the Balkan biodiversity hotspot T he case of the endangered stone crayfish. 2022 , 39, e02301	1
181	Isolation, characterisation and cross-species transferability of microsatellite markers for Riella macrocarpa (P.Allorge) Puche et al. (Riellaceae, Sphaerocarpales). 2022 , 44, 140-145	О
180	Comparative phylogeography of two commensal rat species (Rattus tanezumi and Rattus norvegicus) in China: Insights from mitochondrial DNA, microsatellite, and 2b-RAD data. 2022 , 12,	0
179	Current Genetic Structure Analysis of Leopard Cats Reveals a Weak Disparity Trend in Subpopulations in Beijing, China. 2022 , 11, 1478	О
178	EST-SSR marker-based investigation on genetic diversity and genetic structure of Juglans mandshurica Maxim. in Shandong Province of China.	О
177	Probing for depth-gradient diversification in the riverine-spawning Dolly Varden: Insights from a recently discovered ecomorph assemblage.	О
176	Life-history stage and the population genetics of the tiger mosquito Aedes albopictus at a fine spatial scale.	O
175	Vector competence of lambda-cyhalothrin resistant Aedes aegypti strains for dengue-2, Zika and chikungunya viruses in Colombia. 2022 , 17, e0276493	О
174	Larval Retention and Homing Behaviour Shape the Genetic Structure of the Bullet Tuna (Auxis rochei) in the Mediterranean Sea. 2022 , 7, 300	1
173	Climate-Driven Adaptive Differentiation in Melia azedarach: Evidence from a Common Garden Experiment. 2022 , 13, 1924	О
172	Variation in the local grey mullet populations (Mugil cephalus) on the western Pacific fringe.	О
171	The nuanced nature of mesic refugia in arid landscapes: a tale of two peas.	О

170	Microsatellite Loci of the Atlantic Horseshoe Crab (Limulus polyphemus) Reveal Inter-Localities Genetic Diversity in the Coastal Waters of the Eastern and Northern Yucatan Peninsula.	O
169	Genetic analysis of the exploited snow crab (Chionoecetes opilio) in the Barents Seapossibilities of origin.	O
168	Genetic Diversity and Population Structure of an Arctic Tertiary Relict Tree Endemic to China (Sassafras tzumu) Revealed by Novel Nuclear Microsatellite (nSSR) Markers. 2022 , 11, 2706	O
167	Genetic Diversity and Population Genetic Structure of Aedes albopictus in the Yangtze River Basin, China. 2022 , 13, 1950	O
166	Phylogeography and population genetics of a headwater-stream adapted crayfish, Cambarus pristinus (Decapoda: Cambaridae), from the Cumberland Plateau in Tennessee.	O
165	Genetic population structure of cisco, Coregonus artedi, in the Laurentian Great Lakes. 2022,	O
164	Genetic diversity of whitefly species of the Bemisia tabaci Gennadius (Hemiptera: Aleyrodidae) species complex, associated with vegetable crops in CE de devoire. 2022 , 17, e0276993	O
163	Genetic population structure of the critically endangered stellate sturgeon (Acipenser stellatus) in the Black Sea basin: Implications for conservation.	1
162	Development of novel microsatellite markers for population differentiation and detection of natural selection in wild populations of Butter catfish, Ompok bimaculatus (Bloch, 1794).	O
161	Non-random mating between nesting sites of Hawaiian hawksbill turtles: demographic discontinuity within a small isolated population.	O
160	Natural Populations of Astrocaryum aculeatum Meyer in Amazonia: Genetic Diversity and Conservation. 2022 , 11, 2957	O
159	Genetic patterns of Magnolia in the Lesser Antilles: Stepwise colonisation leading to highly inbred island β opulations \square	O
158	Genetic Diversity in 10 Populations of Domestic Turkeys by Using Microsatellites Markers. 2022 , 102311	O
157	No association between habitat, autogeny and genetics in Moroccan Culex pipiens populations. 2022 , 15,	O
156	Culture of Seriola dumerili in a marine ecosystem: Insights from genetic and morphometric fish traits and implications of escape events. 2022 , 278, 108115	O
155	Analysis of genetic diversity and population structure of the ridgetail white prawn Exopalaemon carinicauda in China. 2022 , 27, 101369	1
154	Genetic differentiation in <i>Palaemon septemtrionalis</i> between populations of two rivers examined by newly developed short tandem repeat markers. 2022 , 51, 123-127	0
153	Country-wide genetic monitoring over 21 years reveals lag in genetic recovery despite spatial connectivity in an expanding carnivore (Eurasian otter, Lutra lutra) population.	O

152	Integrating morphological and genetic limits in the taxonomic delimitation of the Cuban taxa of Magnolia subsect. Talauma (Magnoliaceae). 213, 35-66	0
151	Mating Systems of Single Families and Population Genetic Diversity of Endangered Ormosia hosiei in South China. 2022 , 13, 2117	O
150	The role of habitat features in patterns of population connectivity of two Mediterranean amphibians in arid landscapes of central Iberia.	О
149	Reproductive biology and population structure of the endangered shrub Grevillea bedggoodiana (Proteaceae).	О
148	Genetic Diversity of Montenegrin Local Sheep Breeds Based on Microsatellite Markers. 2022 , 12, 3029	0
147	Genetic diversity, differentiation and phylogeography of the Stenocereus griseus (Haw.) Buxb. species complex (Cactaceae).	О
146	Genetic divergence at species boundaries of the dolphinfish (Coryphaena hippurus) in the Tropical Eastern Pacific. 10, e14389	О
145	Population genetic structure of wild Malayan tapirs (Tapirus indicus) in Peninsular Malaysia revealed by nine cross-species microsatellite markers. 2022 , 40, e02321	О
144	Population genetic structure of Aedes albopictus (Diptera: Culicidae) in China inferred by microsatellite and association with knockdown resistance mutations.	О
143	Early Signs of the Effects of Forest Fragmentation on the Genetic Diversity and Structure of the Threatened Ecuadorian Tree Ocotea rotundata (Lauraceae). 2022 , 13, 1940	О
142	Effect of population size and selection on Toll-like receptor diversity in populations of Galpagos mockingbirds.	0
141	Population genetic structure, parasite infection and somatic condition of pumpkinseed Lepomis gibbosus (Actinopterygii: Centrarchidae) in the Oder river basin.	О
140	Regional environmental and climatic concerns on preserving native gene pools of a least concern species: Brown trout lineages in Mediterranean streams. 2023 , 862, 160739	О
139	Genetic relationships among emu populations in Japanese farms based on mitochondrial and microsatellite DNA polymorphisms. 2022 , 93,	O
138	Genetic structure and diversity of the capercaillie (Tetrao urogallus) population in Belarus in the context of de-lineation of two subspecies: major and pleskei. 2022 , 2022, 113-128	0
137	Trematode Clone Abundance Distributions: An Eco-Evolutionary Link between Parasite Transmission and Parasite Mating Systems. 2022 , 108,	О
136	Do common dispersal influences inform a large lizard landscape-scale gene flow?. 2022 , 36, 987-1006	О
135	First evidence of population genetic structure of the deep-water blackmouth catshark Galeus melastomus Rafinesque, 1810. 9,	O

134	Genetic polymorphisms and affinity analyses of the Chinese Shaanxi Han population utilising a multiplex Y-STR set. 1-12	O
133	Population genetic structure of Brydell whales (Balaenoptera brydei) on the central and western North Pacific feeding grounds.	1
132	Olfactory Sensitivity Is Associated with Body Mass Index and Polymorphism in the Voltage-Gated Potassium Channels Kv1.3. 2022 , 14, 4986	2
131	Genetic and epigenetic variation separately contribute to range expansion and local metalliferous habitat adaptation during invasions of Chenopodium ambrosioides into China.	O
130	Population genetics of an invasive mosquito vector, Aedes albopictus in the Northeastern USA. 78, 99-127	О
129	Genetic study of the American horseshoe crab throughout its Mexican distribution. Conservation and management implications.	O
128	The phylogeography of some soil-feeding termites shaped by the Andes.	О
127	Population structure and genetic diversity in wild dotted gizzard shad (Konosirus punctatus) revealed by microsatellite markers. 9,	O
126	Differentiation of Arctic Charr Salvelinus alpinus Complex (Salmonidae) in Lakes Lama and Kapchuk (Taimyr) Based on Genetic Analysis, External Morphology, and Otolith Shape.	О
125	Genetic diversity and spatial structures of snow leopards (Panthera uncia) reveal proxies of connectivity across Mongolia and northwestern China.	O
124	Conservation Genetics of the Asian Giant Soft-Shelled Turtle (Pelochelys cantorii) with Novel Microsatellite Multiplexes. 2022 , 12, 3459	O
123	The importance of well protected forests for the conservation genetics of West African colobine monkeys.	О
122	Population Structure and Morphological Pattern of the Black-Spotted Pond Frog (Pelophylax nigromaculatus) Inhabiting Watershed Areas of the Geum River in South Korea. 2022 , 14, 16530	1
121	Isolation and characterization of 21 microsatellite loci for the sea cucumber Holothuria (Halodeima) atra (Echinodermata, Holothuroidea) reveal low asexual propagation through time in Reunion Island (southwestern Indian Ocean).	О
120	Individual Identification of Panthera pardus saxicolor Using Non-Invasive Sampling and Molecular Techniques in Iran: A Case Study in Parvar Protected Area. 2022 , 48,	O
119	Genetic Diversity and Population Structure Derived from Body Remains of the Endangered Flightless Longhorn Beetle Iberodorcadion fuliginator in Grassland Fragments in Central Europe. 2023 , 15, 16	O
118	Analysis of SOD1 and C9orf72 mutations in patients with amyotrophic lateral sclerosis in Antioquia, Colombia. 2022 , 42, 623-632	0
117	Effects of Anthropogenic Habitat Fragmentation on the Genetic Connectivity of the Threatened and Endemic Campylorhynchus yucatanicus (Aves, Trogloditydae) in the Yucatan Peninsula, Mexico. 2022 , 14, 1108	O

116	vcfpop: Performing population genetics analyses for autopolyploids and aneuploids based on next-generation sequencing data sets.	Ο
115	Application of genetic and Spatially Explicit Capture-Recapture analyses to design adaptive feral cat control in a large inhabited island. 79, 51-85	Ο
114	Population structure of mud flounder Paralichthys orbignyanus from the south-western Atlantic Ocean.	0
113	New insights on patterns of genetic admixture and phylogeographic history in Iberian high mountain populations of midwife toads. 2022 , 17, e0277298	Ο
112	Genetic Variability and Admixture Zones in the Italian Populations of Turkey Oak (Quercus cerris L.). 2023 , 13, 18	О
111	Genetic Variability of Small Horse Populations from Greek Islands.	O
110	Genetic characterization of Ladakhi donkeys using microsatellite markers. 2019, 89,	1
109	Genetic diversity analysis and parentage verification of Taishu horses using 31 microsatellites. 2022 , 33, 63-69	O
108	Evolutionary history of Castanea sativa Mill. in the Caucasus driven by Middle and Late Pleistocene paleoenvironmental changes.	О
107	Population Subdivision in the Gopher Frog (Rana capito) across the Fragmented Longleaf Pine-Wiregrass Savanna of the Southeastern USA. 2023 , 15, 93	O
106	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.	O
105	Density-habitat relationships of white-tailed deer (Odocoileus virginianus) in Finland. 2023 , 13,	O
104	Water transfer determines the regional spread dynamics of non-native fish species. 2023, 100135	О
103	Performance evaluation of an in-house panel containing 59 autosomal InDels for forensic identification in Chinese Hui and Mongolian groups. 2023 , 115, 110552	O
102	Development of novel microsatellite markers for population differentiation and detection of natural selection in wild populations of butter catfish, Ompok bimaculatus (Bloch, 1794).	O
101	Airborne versus soilborne inoculum: white mould, where do you come from?.	O
100	Phylogeographic and population genetic structure of hound-like native dogs of the Mediterranean Basin. 2023 ,	0
99	Genetic Divergence and Connectivity among Gene Pools of Polyprion americanus. 2023 , 13, 302	O

98	(Not) far from home: No sex bias in dispersal, but limited genetic patch size, in an endangered species, the Spotted Turtle (Clemmys guttata). 2023 , 13,	O
97	Forensic efficiencies of individual identification, kinship testing and ancestral inference in three Yunnan groups based on a self-developed multiple DIP panel. 13,	O
96	Characterization of highly polymorphic microsatellite markers for the chinese monal (Lophophorus lhuysii, Galliformes) using Illumina MiSeq sequencing.	O
95	Inconsistency between morphological diversity and genetic structuring: proposal for one species of Undaria in Japan. 2023 ,	O
94	Genetic survey extension of the threatened Iberian Arnica montana L. revealed the presence of divergent plastid lineages and highly structured populations in northern Spain.	O
93	Genetic variability of Chamaerops humilis (Arecaceae) throughout its native range highlights two species movement pathways from its area of origin.	O
92	Novel insights into the reproductive strategies of wild Chinese sturgeon (Acipenser sinensis) populations based on the kinship analysis. 2023 , 100134	1
91	Development and Characterization of Microsatellite Markers for Harpadon nehereus Based on High-Throughput Sequencing and Cross-Species Amplification in Three Myctophiformes Fishes. 2023 , 22, 181-188	O
90	Genetic diversity of Formosan pangolins (Manis´pentadactyla´pentadactyla) in Taiwan based on the MHC gene.	O
89	No effects of fishery collapse on the genetic diversity of the Gulf of California Corvina, Cynoscion othonopterus (Perciformes: Sciaenidae). 2023 , 261, 106608	O
88	Genetic population dynamics of the critically endangered scalloped hammerhead shark (Sphyrna lewini) in the Eastern Tropical Pacific. 2022 , 12,	O
87	Genetic Diversity and Differences among Three F1 Families and Two Wild Populations of Genus Scylla Using Microsatellite Markers. 2023 , 8, 18	O
86	Integrative characterization of genetic and phenotypic differentiation in an ant species complex with strong hierarchical population structure and low dispersal abilities.	O
85	Genetic diversity, population genetic structure and gene flow in the rare and endangered wild plant Cypripedium macranthos revealed by genotyping-by-sequencing.	O
84	Appraising the Genetic Makeup of an Allochthonous Southern Pike Population: An Opportunity to Predict the Evolution of Introgressive Hybridization in Isolated Populations?. 2023 , 13, 380	O
83	Genetic diversity, structure, and effective population size of an endangered, endemic hoary bat, ?pe?ape?a, across the Hawaiian Islands. 11, e14365	O
82	New Microsatellite Markers for the Model Coral Species Stylophora pistillata from Eilat, the Red Sea. 2023 , 11, 244	О
81	The multilevel society of proboscis monkeys with a possible patrilineal basis.	O

80	Two Distinct Life History Strategies of Atlantic Sturgeon in the Ogeechee River, Georgia. 2023 , 15, 325	O
79	Genetic Diversity and Fine-Scale Genetic Structure of Spodoptera litura Fabricius (Lepidoptera: Noctuidae) in Southern China Based on Microsatellite Markers. 2023 , 13, 560	o
78	History cooling events contributed to the endangered status of Pseudotsuga brevifolia endemic to limestone habitats. 2023 , 42, e02414	0
77	Fragmentation shapes nest density and social structure but not genetic diversity of Temnothorax crassispinus (Formicidae).	o
76	A self-developed AIM-InDel panel designated for degraded DNA analysis: Forensic feature characterization and genetic legacy investigation in the Han Chinese population. 2023 , 110620	0
75	Long-term genetic monitoring of a reintroduced Eurasian lynx population does not indicate an ongoing loss of genetic diversity. 2023 , 42, e02399	0
74	Assessment of population genetic diversity and genetic structure of the North Chinese leopard (Panthera pardus japonensis) in fragmented habitats of the Loess Plateau, China. 2023 , 42, e02416	0
73	Genetic markers associated with divergent selection against the parasite Marteilia cochillia in common cockle (Cerastoderma edule) using transcriptomics and population genomics data. 10,	0
72	Systematically exploring the performance of a self-developed Multi-InDel system in forensic identification, ancestry inference and genetic structure analysis of Chinese Manchu and Mongolian groups. 2023 , 346, 111637	0
71	Polymorphism study of novel microsatellite markers to determine population genetic structure of Coptotermes gestroi (Blattodea: Rhinotermitidae) from infested urban buildings. 2023 , 31, 101768	o
70	Population genetic characteristics of the bumble bee Bombus ardens ardens (Hymenoptera: Apidae) in South Korea using novel microsatellite markers. 2023 , 26, 102071	0
69	Instinctive mate choice with genetically different partners assists sustaining genetic diversity in giant freshwater prawn (Macrobrachium rosenbergii). 2023 , 262, 106648	o
68	Convergent evolution unites the population genetics of Protea-associated ophiostomatoid fungi. 2023 , 63, 101242	0
67	Demography of a nearshore gadid navaga, Eleginus nawaga, from the Barents Sea coast during the last glacial period.	0
66	Heterosis and genetic diversity of intraspecific hybrids crosses between two selected lines of the Pacific oyster Crassostrea gigas. 2023 , 569, 739369	0
65	Development of a diagnostic single nucleotide polymorphism (SNP) panel for identifying geographic origins of Cochliomyia hominivorax, the New World screwworm. 2023 , 315, 109884	0
64	Genetic population structure in Norway lobster (Nephrops norvegicus): management regime under panmixia.	0
63	Comparative analysis of genetic diversity in wild and restored subtropical rainforest trees: Argyrodendron trifoliolatum, Ficus watkinsiana and Diploglottis australis.	О

62	Spatio-temporal genetic structure of the striped field mouse (Apodemus agrarius) populations inhabiting national parks in South Korea: Implications for conservation and management of protected areas. 11,	Ο
61	An integrative study of species distribution modelling and conservation genetics: Magnolia in Hispaniola. 2023 , 32, 1205-1231	O
60	Population genetic analysis of the wild hard-shelled mussel, Mytilus unguiculatus (Valenciennes 1858) in South Korea using a microsatellite multiplex assay.	Ο
59	Are There Edge Effects on the Genetic Diversity of the Trap-Jaw Ant Odontomachus chelifer (Formicidae: Ponerinae) in a Neotropical Savanna Fragment? A First Assessment. 2023 , 52, 279-285	Ο
58	Genotypic detection of barriers to rat dispersal: Rattus rattus behind a peninsula predator-proof fence.	0
57	Molecular epidemiological analyses reveal extensive connectivity between Echinostoma revolutum (sensu stricto) populations across Eurasia and species richness of zoonotic echinostomatids in England. 2023 , 18, e0270672	O
56	Associations of HLA Polymorphisms with Anti-SARS-CoV-2 Spike and Neutralizing Antibody Titers in Japanese Rheumatoid Arthritis Patients Vaccinated with BNT162b2. 2023 , 11, 404	0
55	Genetic Divergence of Thai Indigenous Pigs from Three Distinct Geographic Regions Revealed by Microsatellite Marker Analysis. 2023 , 13, 625	O
54	Temporal decline of genetic differentiation among populations of western flower thrips across an invaded range.	O
53	Genetic population structure ofPseudoplatystoma corruscans(Siluriformes: Pimelodidae) and evidence of temporal variation in structure.	O
52	Genotypic Analysis of the Population Structure in Malassezia globosa and Malassezia restricta. 2023 , 9, 263	O
51	Population genetic structure of a recent insect invasion: a gall midge, Asynapta groverae (Diptera: Cecidomyiidae) in South Korea since the first outbreak in 2008. 2023 , 13,	O
50	Amazonian birds in more dynamic habitats have less population genetic structure and higher gene flow.	0
49	Genetic differences among the Interior Highlands walleye (Sander vitreus) with mitochondrial and nuclear markers indicate the need for updated stocking practices.	O
48	Population structure discovered in juveniles of Greenland halibut (Reinhardtius hippoglossoidesWalbaum, 1792).	O
47	Effective population size in a partially clonal plant is not predicted by the number of genetic individuals. 2023 , 16, 750-766	O
46	Non-synonymous variation and protein structure of candidate genes associated with selection in farm and wild populations of turbot (Scophthalmus maximus). 2023 , 13,	О
45	Genetic structure of the Japanese Robin (Larvivora akahige) endemic to East Asian islands.	O

44	Indigenous cattle of Sri Lanka: Genetic and phylogeographic relationship with Zebu of Indus valley and South Indian origin.	О
43	Malaria-driven adaptation of MHC class I in wild bonobo populations. 2023 , 14,	O
42	Roadmap to the study of gene and protein phylogeny and evolution practical guide. 2023 , 18, e0279597	O
41	Commercial sharks under scrutiny: Baseline genetic distinctiveness supports structured populations of small-spotted catsharks in the Mediterranean Sea. 10,	O
40	Fine-scale functional connectivity of two syntopic pond-breeding amphibians with contrasting life-history traits: an integrative assessment of direct and indirect estimates of dispersal.	O
39	Weak range-wide population structure in the blackfin tuna (Thunnus atlanticus) revealed by analysis of genome-wide SNPs.	O
38	An extreme environment drives local adaptation of Genista tinctoria (Fabaceae) from the Mefite (Ansanto Valley, southern Italy).	O
37	Determination of declined genetic diversity of Holstein stud bulls based on microsatellite markers. 1-7	O
36	Population genomics reveals differences in genetic structure between two endemic arboreal rodent species in threatened cloud forest habitat. 2023 , 68, 223-235	O
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