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- 331 Teaching writing skills in online paleontology and evolution courses. 7, ○
- 330 Chromosome-level genome assembly and resequencing of camphor tree (*Cinnamomum camphora*) provides insight into phylogeny and diversification of terpenoid and triglyceride biosynthesis of *Cinnamomum*. ○
- 329 Isometric spiracular scaling in scarab beetles: Implications for diffusive and advective oxygen transport. 11, 1
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- 327 Discovering candidate SNPs for resilience breeding of red clover. 13, ○
- 326 The population-level impact of *Enterococcus faecalis* genetics on intestinal colonisation and extraintestinal infection. ○
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- 315 Molecular phylogeny of the tropical wandering spiders (Araneae, Ctenidae) and the evolution of eye conformation in the RTA clade. 1
- 314 A global analysis of viviparity in squamates highlights its prevalence in cold climates. 0
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- 305 Species delimitation polyphasic approach reveals *Meyerella similis* sp. nov.: a new species of "small green balls" within the *Chlorella*-clade (Trebouxiophyceae, Chlorophyta). 0
- 304 Correlation between seabirds and fisheries varies by species at fine-scale pattern. 0
- 303 Common evolutionary origin of acoustic communication in choanate vertebrates. **2022**, 13, 4
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- 301 Transcriptome-based phylogeny of the semi-aquatic bugs (Hemiptera: Heteroptera: Gerromorpha) reveals patterns of lineage expansion in a series of new adaptive zones. 0
- 300 Critical pollination chemistry: Specific sesquiterpene floral volatiles in carrot inhibit honey bee feeding. 0
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- 296 Stuck on top of a mountain: Consequences of dispersal limitations for alpine diversity. ○
- 295 Oceanographic gradients explain changes in the biological traits of nesting seabird assemblages across the south-eastern Pacific. 9, ○
- 294 Geographic Range Overlap Rather than Phylogenetic Distance Explains Rabies Virus Transmission among Closely Related Bat Species. **2022**, 14, 2399 ○
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- 284 Population structure of threatened caribou in western Canada inferred from genome-wide SNP data. ○
- 283 A Chromosome-Scale Genome Assembly of *Mitragyna speciosa* (Kratom) and the Assessment of Its Genetic Diversity in Thailand. **2022**, 11, 1492 1
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- 274 Why is the beautyberry so colourful? Evolution, biogeography, and diversification of fruit colours in *Callicarpa* (Lamiaceae). 2022, 2
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- 242 Molecular diversity and biogeography of benthic microeukaryotes in temperate seagrass (*Zostera japonica*) systems of northern China. **2022**, 41, 115-125 ○
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- 239 Interspecific differences and ecological correlations of ultraviolet radiation tolerance in low- and high-altitude fishes. 9, ○
- 238 Whole-genome sequencing analysis of wild house mice (<i>Mus musculus</i>) captured in Madagascar. **2022**, ○
- 237 Aquaculture rearing systems induce no legacy effects in Atlantic cod larvae or their rearing water bacterial communities. **2022**, 12, ○
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- 171 Contrasting Phylogeographic Patterns of Mitochondrial and Genome-Wide Variation in the Groundwater Amphipod *Crangonyx islandicus* That Survived the Ice Age in Iceland. **2023**, 15, 88 ○
- 170 Adaptive evolution of carnivory in *Nepenthes* pitcher plants: a comparative transcriptomics and proteomics perspective. ○
- 169 Phylogenomics reveals pervasive ancient introgression in the radiation of macaques. ○
- 168 A nuclear target sequence capture probe set for phylogeny reconstruction of the charismatic plant family Bignoniaceae. 13, ○
- 167 Molecular Features and Antimicrobial Susceptibilities of *Streptococcus equi* ssp. *equi* Isolates from Strangles Cases in Indonesia. **2023**, 10, 49 ○
- 166 Competitiveness and symbiotic efficiency in alfalfa of *Rhizobium favelukesii* ORY1 strain in which homologous genes of peptidases HrrP and SapA that negatively affect symbiosis were identified. 4, ○
- 165 The Evolution of Life History Traits and Their Thermal Plasticity in *Daphnia*. **2023**, 2, 55-74 ○
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- 162 Spatial genetic structure of European wild boar, with inferences on late-Pleistocene and Holocene demographic history. ○
- 161 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, ○
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