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- 250 Immune Monitoring of Paediatric Patients Infected with *Rickettsia rickettsii*, *Ehrlichia canis* and Coinfected. **2022**, 11, 1351 ○
- 249 Correlation between microbial communities and key odourants in fermented capsicum inoculated with *Pediococcus pentosaceus* and *Cyberlindnera rhodanensis*. ○
- 248 Plant growth promoting and antimicrobial functionalities of rhizosphere *Pseudomonas* species in flooded rice soils of coastal Odisha. ○
- 247 Distribution and Identification of *Ulva aragoensis* (Ulvaceae, Chlorophyta), a Constituent Species of Green Tides in the Southern Yellow Sea, Based on Molecular Data. **2022**, 10, 1767 ○
- 246 Insights into the structure and composition of mineralized hard cocoons constructed by the oriental moth, *Monema* (*Cnidocampa*) *flavescens* Walker. **2022**, 103878 ○
- 245 Identification and Characterization of the HD-Zip Gene Family and Dimerization Analysis of HB7 and HB12 in *Brassica napus* L.. **2022**, 13, 2139 ○
- 244 Development of a Novel HTLV-1 Protease: Human Fc α Recombinant Fusion Molecule in the CHO Eukaryotic Expression System. ○
- 243 Responses of ABCB and ABCC transporters to the toxic dinoflagellate *Prorocentrum lima* in the mussel *Perna viridis*. **2023**, 254, 106368 1
- 242 Molecular cloning and characterization of NLRX1 from orange-spotted grouper (*Epinephelus coioides*) as a mediator of ROS production through its LRR domains. **2023**, 565, 739146 ○
- 241 Pocket structures of surface protrusions shared among serologically distinct nervous necrosis viruses (NNVs) were predicted in silico to bind to sialylated N-glycans, a host cellular receptor. **2023**, 565, 739157 ○
- 240 Phylogenetic analysis of the genus *Miscanthus* and its relative genera (Poaceae) in China. **2023**, 192, 116113 ○

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- 238 Characterization and expression analysis of the interferon regulatory factor (IRF) gene family in zig-zag eel (*Mastacembelus armatus*) against *Aeromonas veronii* infection. **2023**, 140, 104622 ○
- 237 Description of *Pseudogemmobacter faecipullorum* sp. nov., isolated from poultry manure. **2022**, 369, ○
- 236 Discovery, Genomic Sequence Characterization and Phylogenetic Analysis of Novel RNA Viruses in the Turfgrass Pathogenic *Colletotrichum* spp. in Japan. **2022**, 14, 2572 ○
- 235 Molecular characterization and phylogenetic relationships in Asiatic *Vigna* using ITS and cpDNA loci. ○
- 234 Application of microalgae *Chlamydomonas applanata* M9V and *Chlorella vulgaris* S3 for wheat growth promotion and as urea alternatives. 13, ○
- 233 Heterologous expression of MiMLDP, a major lipid droplet protein of the microalga *Myrmecea incisa*, in *Saccharomyces cerevisiae* promotes more triacylglycerol storage principally by enlarging lipid droplet size. ○
- 232 Brittle culm 3, encoding a cellulose synthase subunit 5, is required for cell wall biosynthesis in barley (*Hordeum vulgare* L.). 13, ○
- 231 *Fructobacillus apis* sp. nov., isolated from the gut of honeybee (*Apis mellifera*). **2022**, 72, ○
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- 229 Systemic Infection of *Mycobacterium abscessus* in a Free-Ranging Wild Eurasian Eagle Owl (*Bubo bubo*). **2022**, 58, ○
- 228 Phosphorus-Solubilizing Capacity of *Mortierella* Species Isolated from Rhizosphere Soil of a Poplar Plantation. **2022**, 10, 2361 ○
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- 223 Characterization of a novel mycotombus-like virus from the plant pathogenic fungus *Phoma matteucciicola*. ○
- 222 *Succinivibrio faecicola* sp. nov., isolated from cow faeces. **2022**, 72, ○

- 221 *Spiribacter halobius* sp. nov., a novel halophilic Gammaproteobacterium with a relatively large genome. 9, ○
- 220 *Halomonas binhaiensis* sp. nov., isolated from saline-alkali soil. **2022**, 72, ○
- 219 Proposal to reclassify *Aquiflexum aquatile* into a novel genus as *Cognataquiflexum aquatile* gen. nov., comb. nov., and description of *Cognataquiflexum nitidum* sp. nov. and *Cognataquiflexum rubidum* sp. nov., isolated from freshwater lakes on the Tibetan Plateau. **2022**, 72, ○
- 218 *Exophiala yunnanensis* and *Exophiala yuxiensis* (Chaetothyriales, Herpotrichiellaceae), two new species of soil-inhabiting *Exophiala* from Yunnan Province, China. 94, 109-124 ○
- 217 Molecular characterizations of TLR1 and TLR2 in Qihe crucian carp (*Carassius auratus*) and responses to stimulations of *Aeromonas hydrophila* and TLR ligands. ○
- 216 Genome-Wide Identification and Expression Analysis of SWEET Family Genes in Sweet Potato and Its Two Diploid Relatives. **2022**, 23, 15848 ○
- 215 *Lysobacter lactosilyticus* sp. nov., a Novel β -Galactosidase Producing Bacterial Strain Isolated from Farmland Soil Applied with Amino Acid Fertilizer. **2023**, 80, ○
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- 212 *Allium sulaimanicum*: A new *Allium* species and section from Pakistan. 13, ○
- 211 Origin and evolutionary history of *Populus* (Salicaceae): Further insights based on time divergence and biogeographic analysis. 13, ○
- 210 Genetic Diversity Evaluation and Conservation of Topmouth Culter (*Culter alburnus*) Germplasm in Five River Basins in China. **2023**, 12, 12 ○
- 209 Susceptibility of fig cultivars to *Diaporthe* canker in Iran. 1
- 208 Diversity of *Trichoderma* species associated with soil in the Zoige alpine wetland of Southwest China. **2022**, 12, 1
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- 204 Genome-wide comparison and identification of myosin gene family in *Arabidopsis thaliana* and *Helianthus annuus*. **2022**, 8, e12070 ○

- 203 A new contribution to *Megasporoporia sensu lato*: Six new species and three new combinations. **13**, 1
- 202 Identification of Peanut AhMYB44 Transcription Factors and Their Multiple Roles in Drought Stress Responses. **2022**, 11, 3522 0
- 201 Low level of DNA variation in blast resistance Pi-ta gene of wild, weedy rice and Thai jasmine rice Khao Dawk Mali 105. 0
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- 199 *Devosia oryzae* sp. nov., a novel moderately halotolerant bacterium isolated from the roots of rice plants and genome mining revealed the biosynthesis potential as plant growth promoter. 0
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- 197 *Cedecea sulfonyleivorans* sp. nov., a novel chlorimuron-ethyldegrading bacterium isolated from an herbicides-degrading consortium. **2023**, 205, 0
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- 190 *Delphinieae* flowers originated from the rewiring of interactions between duplicated and diversified floral organ identity and symmetry genes. 0
- 189 *Paraconexibacter antarcticus* sp. nov., a novel actinobacterium isolated from Antarctic tundra soil. **2022**, 72, 0
- 188 Morphological and genetic analysis of *Ceratomyxa saurida* Zhao et al. 2015 and *Ceratomyxa mai* sp. nov. (Myxozoa: Ceratomyxidae) from the East China Sea. **2023**, 73, 0
- 187 Characterization of the First Alternavirus Identified in *Fusarium avenaceum*, the Causal Agent of Potato Dry Rot. **2023**, 15, 145 0
- 186 *Caproiciproducens faecalis* sp. nov., Isolated from Cow Faeces. **2023**, 80, 0

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- 184 Genetic evidence for *Colletotrichum gloeosporioides* transmission between the invasive plant *Ageratina adenophora* and co-occurring neighbor plants. ○
- 183 Diversification of *Sinorhizobium* populations associated with *Medicago polymorpha* and *Medicago lupulina* in purple soil of China. 13, ○
- 182 *Cytospora parapleurivora* sp. nov. isolated from orchards with fruit tree decline syndrome in Ontario, Canada. **2023**, 18, e0279490 ○
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- 178 Genetic Diversity and Cryptic Species Identification of Genus *Triplophysa* from River Swat in Malakand Division, Khyber Pakhtunkhwa, Pakistan. **2023**, 2023, 1-8 ○
- 177 Novel insights into the reproductive strategies of wild Chinese sturgeon (*Acipenser sinensis*) populations based on the kinship analysis. **2023**, 100134 1
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- 169 Species delimitation in *Ceratozamia* (Zamiaceae) from Southwestern Mexico, in light of reproductive and climatic diversification. ○
- 168 First Mitogenome of Endangered *Enteromius thysi* (Actinopterygii: Cypriniformes: Cyprinidae) from Africa: Characterization and Phylogeny. **2023**, 8, 25 1

- 167 Mitochondrial DNA and Distribution Modelling Evidenced the Lost Genetic Diversity and Wild-Residence of Star Tortoise, *Geochelone elegans* (Testudines: Testudinidae) in India. **2023**, 13, 150 ○
- 166 Isolation and Molecular Confirmation of an indole-negative <i>Klebsiella oxytoca</i>; isolated from a patient with cystitis. **2022**, ○
- 165 Genome-scale metabolic reconstruction of 7,302 human microorganisms for personalized medicine. 1
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- 163 Integrative taxonomy for the traditional coccidians (Chromista: Miozoa: Eimeriidae) from island canaries (Aves: Passeriformes: Fringillidae): Worldwide distribution, morphological and molecular characterization, revaluations and establishment of junior synonyms. ○
- 162 Morphological characterization and genetic diversity of a new microsporidium, *Neoflabelliforma dubium* n. sp. from the adipose tissue of *Diaphanosoma dubium* (Crustacea: Sididae). 13, ○
- 161 Genomic analysis of an ultrasmall freshwater green alga, *Medakamo hakoo*. **2023**, 6, ○
- 160 Genome-Wide Identification and Functional Characterization of Stress Related Glyoxalase Genes in *Brassica napus* L.. **2023**, 24, 2130 ○
- 159 *Vibrio sinus* sp. nov., a marine bacterium isolated from coastal seawater. **2023**, 73, ○
- 158 *Vibrio amylolyticus* sp. nov. and *Vibrio gelatinilyticus* sp. nov., two marine bacteria isolated from surface seawater of Qingdao. **2023**, 73, ○
- 157 Diversity and Antimicrobial Activities of Actinobacteria Isolated from Mining Soils in Midelt Region, Morocco. **2023**, 2023, 1-19 ○
- 156 A Wider and Deeper Peptide-Binding Groove for the Class I Molecules from B15 Compared with B19 Chickens Correlates with Relative Resistance to Marek's Disease. ○
- 155 Calnexin, More Than Just a Molecular Chaperone. **2023**, 12, 403 ○
- 154 A new isolate of *Mesorhabditis monhystera* (B schli, 1873) Dougherty, 1955 (Rhabditida: Rhabditidae): re-evaluated with molecular data and scanning electron microscopic observations. **2023**, 97, ○
- 153 *Actinomadura terrae* sp. nov. and *Actinomadura litoris* OS3-89, isolated from rhizosphere soil of cactus. **2023**, 73, ○
- 152 DNA Barcoding of Morphologically Characterized Mosquitoes Belonging to the Genus *Mansonia* from the Atlantic Forest and Brazilian Savanna. **2023**, 14, 109 ○
- 151 *Bacillus litorisediminis* sp. nov., a Thermophilic Bacterium Isolated from Mangrove Sediment. **2023**, 80, ○
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- 149 Complete genome sequence of a novel mycovirus from *Pleurotus citrinopileatus*. **2023**, 168, ○
- 148 Two new species of Hymenochaetaceae from tropical Asia and America. 12, ○
- 147 Characterization of the REVELLE family in Rosaceae and role of PblHY in flowering time regulation. **2023**, 24, ○
- 146 Complete genome sequence of a novel fusarivirus from *Rhizoctonia solani* AG-3 PT strain 3P-2-2. **2023**, 168, ○
- 145 Phylogeography and introgression between *Pinus kesiya* and *P. yunnanensis* in Southeast Asia. ○
- 144 Morphological and molecular characterization of *Hysterothylacium* spp. parasitizing *Pomatomus saltatrix* and *Pagrus pagrus* of the State of S̃ Paulo, Brazil. **2023**, 95, ○
- 143 Field isolates of *Monilinia fructicola* Change resistance pattern to greater sensitivity to thiophanate-methyl in recent populations. ○
- 142 Investigation of B-atp6-orfH79 distributing in Chinese populations of *Oryza rufipogon* and analysis of its chimeric structure. **2023**, 23, ○
- 141 Recent status and distribution pattern of cotton leaf curl disease in Northwest India: an alarming situation in future cotton cultivation. **2017**, 87, 2
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- 136 *Frateuria soli* sp. nov. and *Frateuria edaphi* sp. nov., isolated from greenhouse soil. **2023**, 73, ○
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- 132 Phosphate-solubilizing fungi in riparian sediment of Xiangxi Bay (China) and implications of phosphorus pollution control. ○

- 131 Isolation and Screening of Polyhydroxybutyrate (PHB) Producing Bacteria from Soils. ○
- 130 *Collinsella urealyticum* sp. nov., a urease-positive bacterial strain isolated from swine faeces. **2023**, 205, ○
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- 123 *Ectobacillus ponti* sp. nov., a novel bacterium isolated from Pearl River Estuary. **2023**, 73, ○
- 122 Expression and functional analyses for estrogen receptor and estrogen related receptor of Yesso scallop, *Patinopecten yessoensis*. **2023**, 231, 106302 ○
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- 120 The RALF Signaling Pathway Regulates Cell Wall Integrity during Pollen Tube Growth in Maize. ○
- 119 *Parabacteroides faecalis* sp. nov. Isolated from Swine Faeces. **2023**, 80, ○
- 118 Genetic diversity of *Contraecaecum rudolphii* sp. A (Nematoda: Anisakidae) parasitizing the European Shag *Phalacrocorax aristotelis desmarestii* from the Spanish Mediterranean coast. 10, ○
- 117 *Arthrobacter hankyongi* sp. nov., Isolated From Wet Land. **2023**, 80, ○
- 116 Genomic identification of cotton SAC genes branded ovule and stress-related key genes in *Gossypium hirsutum*. 14, ○
- 115 Two new corticioid species of Phanerochaetaceae (Polyporales, Basidiomycota) from Southwest China. 13, 1
- 114 A New Species of *Neoscytalidium hylocereum* sp. nov. Causing Canker on Red-Fleshed Dragon Fruit (*Hylocereus polyrhizus*) in Southern Thailand. **2023**, 9, 197 ○

- 113 Impacts of Sourdough Technology on the Availability of Celiac Peptides from Wheat Hard End Gliadins: In Silico Approach. **2023**, 3, 39-57 ○
- 112 Diversity of Mycoviruses Present in Strains of Binucleate Rhizoctonia and Multinucleate Rhizoctonia, Causal Agents for Potato Stem Canker or Black Scurf. **2023**, 9, 214 ○
- 111 Identification and Analysis of Phosphatidylethanolamine-Binding Protein Family Genes in the Hangzhou White Chrysanthemum (*Chrysanthemum morifolium* Ramat). **2023**, 13, 374 ○
- 110 Fine Mapping of a Pleiotropic Locus (BnUD1) Responsible for the Up-Curling Leaves and Downward-Pointing Siliques in *Brassica napus*. **2023**, 24, 3069 ○
- 109 *Muricauda spongiicola* sp. nov., isolated from the sponge *Callyspongia elongata*. **2023**, 73, ○
- 108 A single nucleotide substitution in the MATE transporter gene regulates plastochron and many noded dwarf phenotype in barley (*Hordeum vulgare* L.). **2023**, ○
- 107 The Bcl-2-associated athanogene gene family in tobacco (*Nicotiana tabacum*) and the function of NtBAG5 in leaf senescence. 14, ○
- 106 Genetic diversity study of *Chrysoperla carnea* (Neuroptera: Chrysopidae) populations via molecular markers. **2022**, 54, 1295-1312 ○
- 105 Production and physicochemical properties of fungal chitosans with efficacy to inhibit mycelial growth activity of pathogenic fungi. **2023**, 525, 108762 ○
- 104 Overexpression of CcFALDH from spider plant (*Chlorophytum comosum*) enhances the formaldehyde removing capacity of transgenic gloxinia (*Sinningia speciosa*)1. **2023**, 223, 115466 ○
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- 102 Planthopper salivary sheath protein LsSP1 contributes to manipulation of rice plant defenses. **2023**, 14, ○
- 101 Mitochondriomics of Clarias Fishes (Siluriformes: Clariidae) with a New Assembly of *Clarias camerunensis*: Insights into the Genetic Characterization and Diversification. **2023**, 13, 482 ○
- 100 Biodiversity of Fungi Inhabiting European Blueberry in North-Western Russia and in Finland. **2022**, 507, 441-455 ○
- 99 *Lactococcus intestinalis* sp. nov., a new lactic acid bacterium isolated from intestinal contents in Alzheimer's disease mice. ○
- 98 Prevalence and genetic diversity in bovine parechovirus infecting Japanese cattle. **2023**, 168, ○
- 97 *Bacteroides rhinocerotis* sp. nov., isolated from the fresh feces of rhinoceros in Beijing Zoo. ○
- 96 *Barnettozyma menglunensis* f.a., sp. nov., a novel yeast species isolated from rotting wood. **2023**, 73, ○

- 95 Molecular characterization, developmental biology, and life table of ladybird beetle, *Micraspis discolor* and its predatory potential against *Aphis gossypii*. 1-11 ○
- 94 Evaluation of efficacy and mechanism of *Bacillus velezensis* CB13 for controlling peanut stem rot caused by *Sclerotium rolfsii*. 14, ○
- 93 Gut bacteria of woodboring *Streltzoviella insularis* and their involvement in pinoresinol degradation of ash tree (*Fraxinus* spp.) host. ○
- 92 A New Species of the Genus *Achalinus* (Squamata: Xenodermidae) from the Dabie Mountains, Anhui, China. **2023**, 13, 708 ○
- 91 Giardiasis in children and dogs, and the first report of assemblage E in dogs from northeastern Brazil. **2023**, 32, ○
- 90 Genome-Wide Analysis of SIMILAR TO RCD ONE (SRO) Family Revealed Their Roles in Abiotic Stress in Poplar. **2023**, 24, 4146 ○
- 89 The complete mitochondrial genome of *Triplax ainonia* Lewis, 1877 (Coleoptera: Erotylidae). **2023**, 8, 297-300 ○
- 88 Synthesis of 12-aminododecenoic acid by coupling transaminase to oxylipin pathway enzymes. **2023**, 107, 2209-2221 ○
- 87 *Azospirillum Aestuarii* sp. nov., a Novel Nitrogen-Fixing and Aerobic Denitrifying Bacteria Isolated from an Estuary of a Freshwater River. **2023**, 80, ○
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- 85 Incidence and Epidemiology of Citrus Viroids in Greece: Role of Host and Cultivar in Epidemiological Characteristics. **2023**, 15, 605 1
- 84 *Arthrobacter mangrovi* sp. nov., an actinobacterium isolated from the rhizosphere of a mangrove. **2023**, 73, ○
- 83 Protein sequence analysis. **2023**, 217-223 ○
- 82 GENETIC DIVERSITY OF THE EAST EUROPEAN VOLE (*MICROTUS ROSSIAEMERIDIONALIS*) IN BELARUS. **2022**, 33, 119-128 ○
- 81 Improvement of Laccase Production by *Thielavia terrestris* Co3Bag1. Enhancing the Bio-Catalytic Performance of the Native Thermophilic TtLacA via Immobilization in Copper Alginate Gel Beads. **2023**, 9, 308 ○
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- 78 Genome-wide survey of HMA gene family and its characterization in wheat (*Triticum aestivum*). 11, e14920 ○

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- 76 Genome-Wide Identification and Expression Analysis of the HSF Gene Family in Poplar. **2023**, 14, 510 ○
- 75 A contribution to Porogramme (Polyporaceae, Agaricomycetes) and related genera. **2023**, 14, ○
- 74 Genetic diversity of Toll-like receptor 9 in swamp eels (*Monopterus albus*). ○
- 73 Two Novel Species of *Mesophoma* gen. nov. from China. **2023**, 80, ○
- 72 Genome-wide analysis of bromodomain gene family in *Arabidopsis* and rice. 14, ○
- 71 Characterization of a novel mycotombus-like virus from the plant-pathogenic fungus *Phoma matteucciicola*. **2023**, 168, ○
- 70 Intraspecific genetic diversity of the fish-infecting microsporidian parasite *Pseudokabatana alburnus* (Microsporidia). 14, ○
- 69 Genome-wide identification of bHLH transcription factors and their response to salt stress in *Cyclocarya paliurus*. 14, ○
- 68 Barcoding of white-spotted stink bug, *Eysarcoris ventralis* (Westwood) (Hemiptera: Pentatomidae) attacking Egyptian barley and its population density. ○
- 67 Suppression of Fusarium Wilt in Watermelon by *Bacillus amyloliquefaciens* DHA55 through Extracellular Production of Antifungal Lipopeptides. **2023**, 9, 336 ○
- 66 A new species of *Gracixalus* (Anura, Rhacophoridae) from northwestern Vietnam. 1153, 15-35 ○
- 65 The Overwintering Capability of *Ulva prolifera* Spores and Gametes in the Yellow Sea, China. **2023**, 22, 509-516 ○
- 64 Thioredoxin-2 Regulates SqrR-Mediated Polysulfide-Responsive Transcription via Reduction of a Polysulfide Link in SqrR. **2023**, 12, 699 ○
- 63 Detection of hop stunt viroid variants from naturally infected kumquat and limequat trees in Mazandaran Province, Iran. ○
- 62 The chalara-like anamorphs of *Leotiomyces*. **2023**, 119, 213-490 ○
- 61 Molecular phylogeny and morphology reveal four new species of *Thelephora* (Thelephorales, Basidiomycota) from subtropical China, closely related to *T. ganbajun*. 14, ○
- 60 Plastome-based backbone phylogeny of East Asian *Phedimus* (Subgenus *Aizoon*: Crassulaceae), with special emphasis on Korean endemics. 14, ○

- 59 Historical Landscape Evolution Shaped the Phylogeography and Population History of the Cyprinid Fishes of *Acrossocheilus* (Cypriniformes: Cyprinidae) According to Mitochondrial DNA in Zhejiang Province, China. **2023**, 15, 425 ○
- 58 The transcription factor GhWRKY70 from *Gossypium hirsutum* enhances resistance to *Verticillium* wilt via the jasmonic acid pathway. **2023**, 23, ○
- 57 Diverse branching forms regulated by a core auxin transport mechanism in plants. **2023**, 150, ○
- 56 *Microbacterium tenebrionis* sp. nov. and *Microbacterium allomyrinae* sp. nov., isolated from larvae of *Tenebrio molitor* L. and *Allomyrina dichotoma*, respectively. **2023**, 73, ○
- 55 *Paenibacillus mangrovi* sp. nov., a novel endophytic bacterium isolated from bark of *Kandelia candel*. **2023**, 73, ○
- 54 Description of *Alkalimarinus coralli* sp. nov., isolated from tissue of the coral *Pocillopora damicornis*. **2023**, 73, ○
- 53 *Aquiflexum gelatinilyticum* sp. nov., isolated from river water. **2023**, 73, ○
- 52 *Salinibacterium sedimenticola* sp. nov., Isolated from Tidal Flat Sediment. **2023**, 80, ○
- 51 Baseline Analysis of Endophytic Fungal Associates of *Solenopsis invicta* Buren from Mounds across Five Counties of Guangdong Province, China. **2023**, 9, 377 ○
- 50 SCARECROW maintains the stem cell niche in *Arabidopsis* roots by ensuring telomere integrity. ○
- 49 Genetic Diversity of Dengue Vector *Aedes albopictus* Collected from South Korea, Japan, and Laos. **2023**, 14, 297 ○
- 48 Identification, molecular characterization and risk factors of *Theileria* infection among sheep: a first comprehensive report from North India. 1-13 ○
- 47 Genetic Characterization and Insular Habitat Enveloping of Endangered Leaf-Nosed Bat, *Hipposideros nicobarulae* (Mammalia: Chiroptera) in India: Phylogenetic Inference and Conservation Implication. **2023**, 14, 765 ○
- 46 Genome-Wide Identification and Functional Analysis of the Roles of SAUR Gene Family Members in the Promotion of Cucumber Root Expansion. **2023**, 24, 5940 ○
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- 43 Comparative Analysis of Chloroplast Genome of *Desmodium stryacifolium* with Closely Related Legume Genome from the Phaseoloid Clade. **2023**, 24, 6072 ○
- 42 Complete Genome Sequence of *Streptomyces* sp. HP-A2021, a Promising Bacterium for Natural Product Discovery. ○

- 41 Possible species discrimination of a blotched nerite *Nerita albicilla* with their distribution pattern and demographic history in the Indo-Pacific. **2023**, 13, ○
- 40 *Lasiodiplodias* species associated with mango (*Mangifera indica* L.) decline in Burkina Faso and influence of climatic factors on the disease distribution. ○
- 39 Genome-wide identification of the TIFY family reveals JAZ subfamily function in response to hormone treatment in *Betula platyphylla*. **2023**, 23, ○
- 38 Genetic Diversity of Asian *Vigna* Species (Subgenus *Ceratotropis*; Genus *Vigna*) in India Based on ITS2 Sequences Data. ○
- 37 Morpho-molecular characterisation of angular leaf spot pathogen and identification of new sources of resistance in North-western Himalayan common beans. ○
- 36 Phenotypic variations among *Laudakia stellio* populations in Anatolia: Colour, colour pattern, and phylogeny (Reptilia: Agamidae). 1-19 ○
- 35 Bacteriolytic activity in saliva of the hematophagous *Triatoma infestans* (Reduviidae) and novel characterization and expression site of a third lysozyme. ○
- 34 *Euphorbia helioscopia* a Putative Plant Reservoir of Pathogenic *Curtobacterium flaccumfaciens*. **2023**, 80, ○
- 33 *Sulfurospirillum tamanensis* sp. nov., a Facultatively Anaerobic Alkaliphilic Bacterium from a Terrestrial Mud Volcano. **2023**, 92, 21-29 ○
- 32 *Meretrix taiwanica* (Bivalvia: Veneridae), a previously misidentified new species in Taiwan. 1-10 ○
- 31 Whole-genome identification and salt- and ABA-induced expression trends of the *Nicotiana tabacum* CKX gene family. ○
- 30 *Bacteroides rhinocerotis* sp. nov., isolated from the fresh feces of rhinoceros in Beijing Zoo. **2023**, 205, ○
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- 27 *Limimanicola litoreus* sp. nov., isolated from intertidal sand of the Yellow Sea. **2023**, 73, ○
- 26 MORPHOLOGICAL AND MOLECULAR CHARACTERIZATION OF *Isostichopus* sp. IN THE COLOMBIAN CARIBBEAN SEA. **2018**, 29, 33-48 ○
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- 23 Plant recognition by *Trichoderma harzianum* elicits upregulation of a novel secondary metabolite cluster required for colonization. ○
- 22 *Fusarium* diversity from the Golden Gate Highlands National Park. 14, ○
- 21 *Paraflavisolibacter caeni* gen. nov., sp. nov., a novel taxon within the family Chitinophagaceae isolated from sludge. **2023**, 73, ○
- 20 Cassava MeRS40 is required for the regulation of plant salt tolerance. **2023**, ○
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- 16 Phylogenetic analyses based on molecular and morphological data reveal a new species of *Strigea* *Abildgaard*, 1790 (Digenea: Strigeidae) and taxonomic changes in strigeids infecting Neotropical birds of prey. **2023**, 97, ○
- 15 Characterization of the complete mitochondrial genome of *Culex vishnui* (Diptera: Culicidae), one of the major vectors of Japanese encephalitis virus. ○
- 14 Unbiased analyses of ITS folding motifs in a taxonomically confusing lineage: *Anagnostidinema visiae* sp. nov. (CYANOBACTERIA). ○
- 13 *Aridibaculum aurantiacum* gen. nov., sp. nov., isolated from the Kumtag Desert soil. **2023**, 73, ○
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- 4 An antagonist of *Caenorhabditis elegans* controls root-knot disease and promotes tomato plant growth. 1-13 ○
- 3 Identification of *Spodoptera frugiperda* (Lepidoptera: Noctuidae) and its two host strains in China by PCR-RFLP. ○
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