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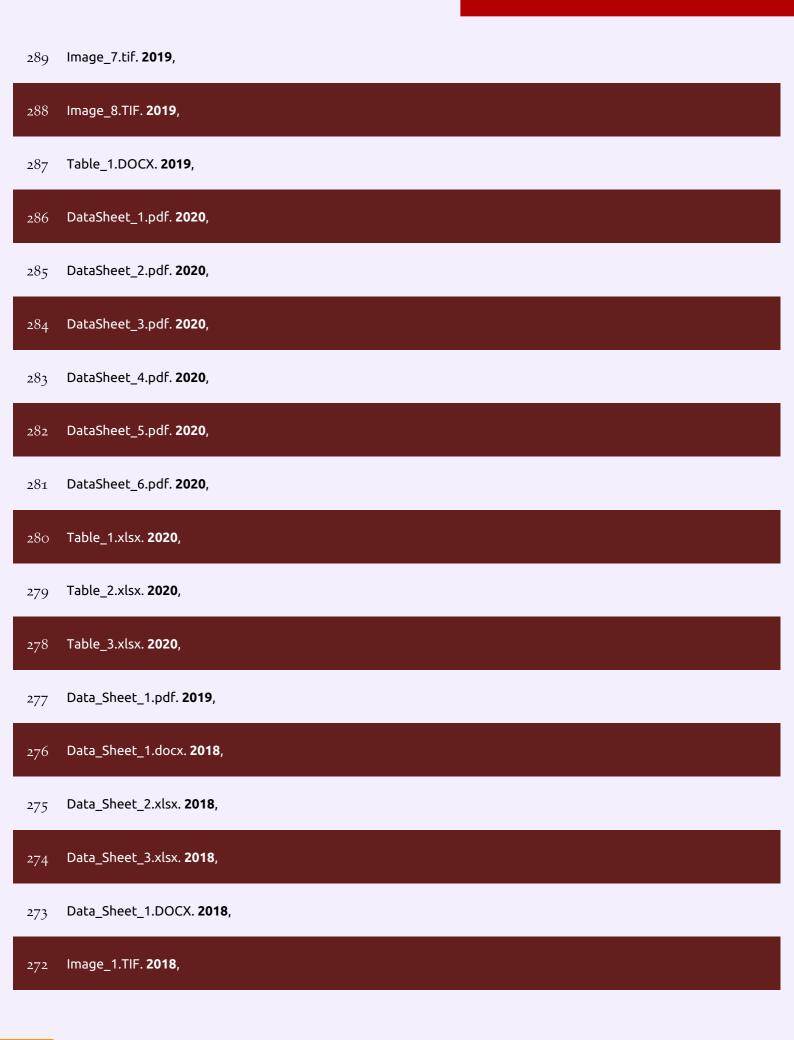
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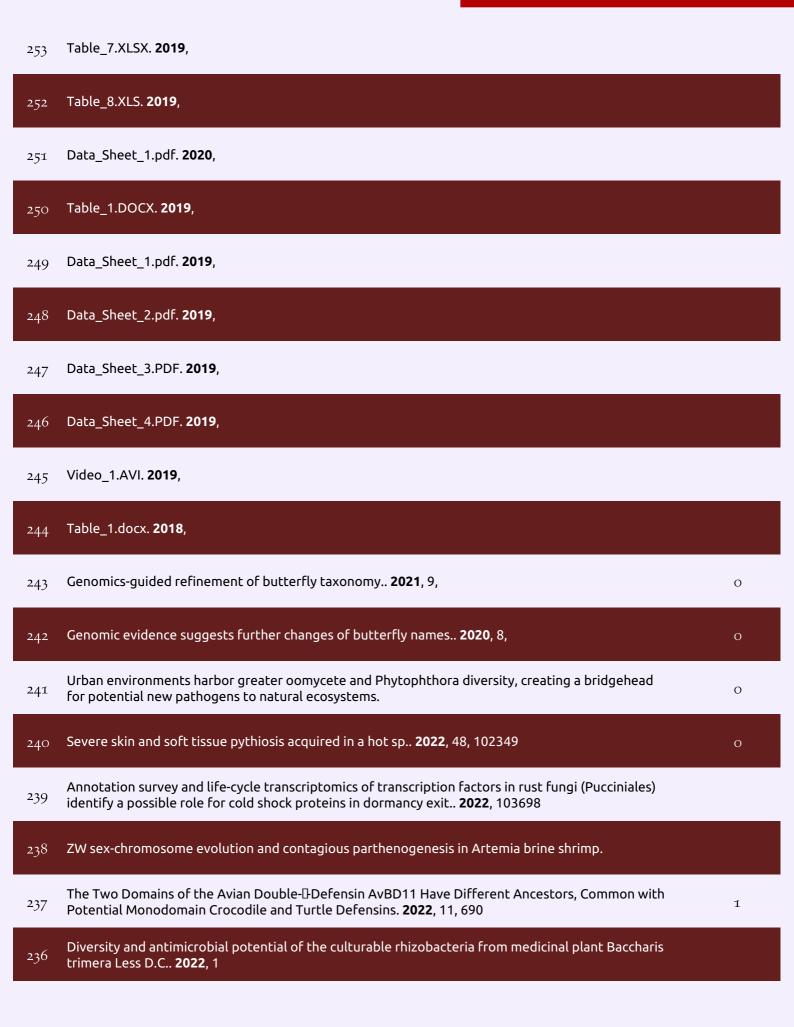
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81	Morphological and molecular evidence reveals a new species of chewing louse Pancola ailurus n. sp. (Phthiraptera: Trichodectidae) from the endangered Chinese red panda Ailurus styani. 2022 ,	О
80	Genotypic and phenotypic differences among phase-variable colony variants conserved acrossGardnerellaspp.	O
79	The Embryonic Origin of Primordial Germ Cells in the TardigradeHypsibius exemplaris.	Ο
78	The complete ITS2 barcoding region for Strongylus vulgaris and Strongylus edentatus.	Ο
77	Genome Characterization of Bacteriophage KPP-1, a Novel Member in the Subfamily Vequintavirinae, and Use of Its Endolysin for the Lysis of Multidrug-Resistant Klebsiella variicola In Vitro. 2023 , 11, 207	0
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74	Myxobolus shuifuensis sp. n. (Myxozoa: Myxobolidae) infecting the exotic mrigal Cirrhinus mrigala feral in China. 2023 , 102732	0

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72	Utilization of microalgae, Chlorella sp. UMT LF2 for bioremediation of Litopenaeus vannamei culture system and harvesting using bio-flocculant, Aspergillus niger. 2023 , 47, 102596	Ο
71	New Negombata species discovered: latrunculin mystery solved.	0
7º	Occurrence of two myxosporean parasites in the gall bladder of white seabream Diplodus sargus (L.) (Teleostei, Sparidae), with the morphological and molecular description of Ceratomyxa sargus n. sp 11, e14599	O
69	Molecular of Anaplasma marginale Theiler (Rickettsiales: Anaplasmataceae) in horseflies (Diptera: Tabanidae) in Uruguay. 2022 , 12,	O
68	Expanded tRNA methyltransferase family member TRMT9B regulates synaptic growth and function.	O
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66	Evolution of TOP1 and TOP1MT Topoisomerases in Chordata.	O
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64	ISSR-assisted spatial genetic structure, population admixture, and biodiversity estimates across locally adopted saffron ecotypes from 18 different provenances of Iran. 2023 , 100467	O
63	A putative xanthine dehydrogenase is critical for Borrelia burgdorferi survival in ticks and mice. 2023 , 169,	O
62	Sugarcane molasses as substrate to soil yeasts: Indole-3-acetic acid production and maize initial growth promotion. 2023 , 47, 102618	1
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60	Biochemical Activation and Regulatory Functions of Trans-Regulatory KLF14 and Its Association with Genetic Polymorphisms. 2023 , 13, 199	O
59	Overview of the Properties of Glutamic Peptidases That Are Present in Plant and Bacterial Pathogens and Play a Role in Celiac Disease and Cancer. 2023 , 62, 672-694	O
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56	Bioremediation of hexavalent chromium by transformation of Escherichia coli DH5 with chromate reductase (ChrR) genes of Pseudomonas putidaisolated from tannery effluent. 2023 , 134,	O

55	Elucidating the life cycle of opossum parasites: DNA sequences reveal the involvement of planorbid snails as intermediate hosts of Rhopalias spp. (Trematoda: Echinostomatidae) in Brazil. 2023 , 18, e0279268	О
54	Acquisition of host-derived carbon in biomass of ectomycorrhizal fungus pisolithus microcarpus is correlated to fungal carbon demand and plant defences.	О
53	Sequencing the genomes of LPP-1, the first isolated cyanophage, and its relative LPP-2 reveal different integration mechanisms in closely related phages. 2023 , 124, 102409	О
52	The embryonic origin of primordial germ cells in the tardigrade Hypsibius exemplaris. 2023, 497, 42-58	O
51	Characterization of chitinases from the GH18 gene family in the myxomycete Physarum polycephalum. 2023 , 1867, 130343	О
50	Population characteristics of Xanthomonas arboricola pv. corylina strains from hazelnut orchards in Turkey. 2023 , 130, 337-349	O
49	Molecular Detection of Candidatus Orientia chuto in Wildlife, Saudi Arabia. 2023 , 29, 402-406	O
48	Novel Species of Brucella Causing Human Brucellosis, French Guiana. 2023 , 29, 333-340	O
47	Mechanistic and evolutionary insights into alkaline phosphatase superfamily through structure-function studies on Sphingomonas alkaline phosphatase. 2023 , 736, 109524	О
46	A novel, non-neuronal acetylcholinesterase of schistosome parasites is essential for definitive host infection. 14,	O
45	Apical Secretory Glycoprotein Complex Contributes to Cell Attachment and Entry by Cryptosporidium parvum. 2023 , 14,	О
44	Diversity and pathogenicity of species of Botrytis , Cladosporium , Neopestalotiopsis and Pestalotiopsis causing flower diseases of macadamia in Australia.	О
43	Identification, classification and characterization of the dermonecrotic toxins in venom glands of Hottentotta saulcyi, Androctonus crassicauda and Hemiscorpius lepturus using transcriptome analysis.	О
42	Ets-1 transcription factor regulates glial cell regeneration and function in planarians.	O
41	Exploring the Relationship between Biosynthetic Gene Clusters and Constitutive Production of Mycosporine-like Amino Acids in Brazilian Cyanobacteria. 2023 , 28, 1420	1
40	Aim18p and Aim46p are chalcone isomerase domaindontaining mitochondrial hemoproteins in Saccharomyces cerevisiae. 2023 , 299, 102981	O
39	Planarian LDB and SSDP proteins scaffold transcriptional complexes for regeneration and patterning.	О
38	Discovery of a coronavirus in the Eurasian badger (Meles meles) belonging to a putative new genus. 2023 , 109, 105406	O

37	Heterotrimeric G proteins regulate planarian regeneration and behavior. 2023, 223,	0
36	Expansion of Kuravirus-like Phage Sequences within the Past Decade, including Escherichia Phage YF01 from Japan, Prompt the Creation of Three New Genera. 2023 , 15, 506	O
35	Biomining Sesuvium portulacastrum for halotolerant PGPR and endophytes for promotion of salt tolerance in Vigna mungo L 14,	O
34	The midbody component Prc1-like is required for microtubule reorganization during cytokinesis and dorsal determinant segregation in the early zebrafish embryo. 2023 , 150,	O
33	Carp edema virus in Ukraine I The evidence for the furthest east presence of CEV genogroup I in Europe. 2023 , 29, 101500	0
32	DnaK and DnaJ proteins from Hsp70/40 family are involved in Rubisco biosynthesis in Synechocystis sp. PCC6803 and sustain the enzyme assembly in a heterologous system. 2023 , 23,	O
31	Fungal diversity notes 1512Il610: taxonomic and phylogenetic contributions on genera and species of fungal taxa. 2022 , 117, 1-272	2
30	A Novel Memory Type inC. elegansExhibits Post-Training Consolidation.	O
29	Eco-sustainable bioremediation of industrial azodye micropollutants by extremophilic plant growth promoting bacterium Halomonas desertis G11. 2022 , 6, 13-19	0
28	Genomic Analysis of Haloarchaea from Diverse Environments, including Permian Halite, Reveals Diversity of Ultraviolet Radiation Survival and DNA Photolyase Gene Variants. 2023 , 11, 607	O
27	Comparative molecular and morphological species delineation of Halammohydra Remane, 1927 (Hydrozoa) with the description of four new species.	O
26	Rickettsia felis DNA recovered from a child who lived in southern Africa 2000 years ago. 2023 , 6,	O
25	Genome-wide identification, characterization and expression analysis of AGO, DCL, and RDR families in Chenopodium quinoa. 2023 , 13,	O
24	Genetic Predisposition to Neuroblastoma Results from a Regulatory Polymorphism that Promotes the Adrenergic Cell State.	O
23	MeC3HDZ1/MeCNA is a strong candidate for cassava storage root productivity improvement.	0
22	Tenacibaculum. 1-26	O
21	The complete mitochondrial genome of a wild-collected Kappaphycus malesianus (Solieriaceae, Rhodophyta). 2023 , 8, 359-363	0
20	Passing the post: Roles of post-translational modifications in the form and function of extracellular matrix.	O

19	Domain Architecture of the Nonreceptor Tyrosine Kinase Ack1. 2023 , 12, 900	0
18	Scleromatobacter humisilvae gen. nov., sp. nov., a novel bacterium isolated from oak forest soil. 2023 , 73,	O
17	Characterization of the l-arabinofuranose-specific GafABCD ABC transporter essential for l-arabinose-dependent growth of the lignocellulose-degrading bacterium Shewanella sp. ANA-3. 2023 , 169,	0
16	Evolution of visual guanylyl cyclases and their activating proteins with respect to clade and species-specific visual system adaptation. 16,	O
15	Novel R2R3 MYB transcription factors regulate anthocyanin synthesis in Aubergine tomato plants. 2023 , 23,	О
14	Recent Progress of Reclassification of the Genus Streptomyces. 2023 , 11, 831	O
13	Improved rock phosphate dissolution from organic acids is driven by nitrate assimilation of bacteria isolated from nitrate and CaCO3-rich soil. 2023 , 18, e0283437	0
12	Rhizosphere and detritusphere habitats modulate expression of soil N-cycling genes during plant development.	0
11	Multi-Knock multi-targeted genome-scale CRISPR toolbox to overcome functional redundancy in plants. 2023 , 9, 572-587	0
10	Human seven-⊡strand (METTL) methyltransferases - conquering the universe of protein lysine methylation. 2023 , 104661	O
9	The Lytic Activity of Bacteriophage ZCSE9 against Salmonella'enterica and Its Synergistic Effects with Kanamycin. 2023 , 15, 912	О
8	Geographical subdivision of Alviniconcha snail populations in the Indian Ocean hydrothermal vent regions. 10,	o
7	Genome-based classification of Pedobacter polysacchareus sp. nov., isolated from Antarctic soil producing exopolysaccharide.	О
6	Origin and Evolution of RAS Membrane Targeting.	O
5	Adaptive evolution characteristics of mitochondrial genomes in genus Aparapotamon (Brachyura, Potamidae) of freshwater crabs. 2023 , 24,	0
4	Genomic characterization of an Omono River virus isolated from Culex tritaeniorhynchus in eastern China. 2023 , 20,	0
3	Presence and activity of nitrogen-fixing bacteria in Scots pine needles in a boreal forest: a nitrogen-addition experiment.	0
2	Complete genome sequence of Aeromonas phage GomatiRiver_11, a novel T4-like bacteriophage that infects Aeromonas hydrophila.	O

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