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1503	Genome analysis of a major urban malaria vector mosquito, Anopheles stephensi. <b>2014</b> , 15, 459		80
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1501	Decay of interspecific avian flock networks along a disturbance gradient in Amazonia. <b>2014</b> , 281, 20132	599	61
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1497	Hierarchy of climate and hydrological uncertainties in transient low-flow projections. <b>2016</b> , 20, 3651-36	72	34
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1494	Artificial Seawater Media Facilitate Cultivating Members of the Microbial Majority from the Gulf of Mexico. <b>2016</b> , 1,		33
1493	Failure of Colorectal Surgical Site Infection Predictive Models Applied to an Independent Dataset: Do They Add Value or Just Confusion?. <b>2016</b> , 222, 431-8		26
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1491	Co-acclimation of bacterial communities under stresses of hydrocarbons with different structures. <b>2016</b> , 6, 34588		38
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1481	Male apoE*3-Leiden.CETP mice on high-fat high-cholesterol diet exhibit a biphasic dyslipidemic response, mimicking the changes in plasma lipids observed through life in men. <b>2017</b> , 5, e13376	12
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1105	: A Manual Collection of Circulating microRNAs of Prognostic and Diagnostic Value. 2021, 13,	1
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1100	Nuclear-cytoplasmic balance: whole genome duplications induce elevated organellar genome copy number. <b>2021</b> , 108, 219-230	1
1099	vlda: An R package for statistical visualization of multidimensional longitudinal data. <b>2021</b> , 28, 369-391	
1098	Genetic variability and nitrogen response indices in common bean (Phaseolus vulgaris) cultivars under contrasting nitrogen environments. <b>2021</b> , 140, 907	Ο
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865	Computational Assessment of Botrytis cinerea Lipase for Biofuel Production. <b>2021</b> , 11, 1319	Ο
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842	Protein-Protein Interactions Shape Genomic Autoimmunity in the Adaptively-Evolving Rhino-Deadlock-Cutoff (RDC) Complex.	
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840	Le lynx Lynx pardinus spelaeus Boule, 1910 du Pl^ Istoc^ Ēe moyen de la grotte de l'Escale (Bouches-du-Rh^ Ēe, France) : donn^ ès pal^ bntologiques et taphonomiques. <b>2020</b> , 108-137	2
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813	Single organoid RNA-sequencing reveals high organoid-to-organoid variability.	O
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811	Regulation of PP2A, PP4, and PP6 holoenzyme assembly by carboxyl-terminal methylation. <b>2021</b> , 11, 23031	O
810	Diel Movement by White Suckers (Catostomus commersonii) in Connected Northern Lakes. <b>2021</b> , 28,	
809	Deciphering Multidrug-Resistant Acinetobacter baumannii from a Pediatric Cancer Hospital in Egypt. <b>2021</b> , e0072521	2
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804	Thermal-metabolic phenotypes of the lizard Podarcis muralis differ across elevation, but converge in high elevation hypoxia. <b>2021</b> ,	1
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## (2021-2021)

802	Seasonal Dynamics of Bathyarchaeota-Dominated Benthic Archaeal Communities Associated with Seagrass (Zostera japonica) Meadows. <b>2021</b> , 9, 1304	1
801	Spatial and Temporal Constraints on the Composition of Microbial Communities in Subsurface Boreholes of the Edgar Experimental Mine. <b>2021</b> , e0063121	O
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799	Peptidomic analysis characterising proteolysis in thaw-aging of beef short plate <b>2021</b> , 3, 100051	O
798	PRDX-1 supports the survival and antitumor activity of primary and CAR-modified NK cells under oxidative stress. <b>2021</b> ,	3
797	Low parental socioeconomic position results in longer post-Norwood length of stay 2021,	6
796	Revisiting proboscidean phylogeny and evolution through total evidence and palaeogenetic analyses including ancient DNA <b>2022</b> , 25, 103559	3
795	Smoothing of Time-Varying Graph With the Generalized LASSO. <b>2021</b> , 9, 162348-162358	
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793	Disrupting Personal (In)Security? The Role of Ride-Hailing Service Features, Commute Strategies, and Gender in Mexico City.	
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784	Complex hydrological knowledge to support digital soil mapping. <b>2022</b> , 409, 115638	O
783	Protected areas in South Asia: Status and prospects <b>2021</b> , 811, 152316	O
782	An^ llsis del flujo comercial en Am^ fica Latina y el Caribe. <b>2021</b> , 1, 36-45	
781	Single-cell-resolved dynamics of chromatin architecture delineate cell and regulatory states in zebrafish embryos. <b>2022</b> , 2, 100083	1
780	Microbiomes of air dust collected during the ground-based closed bioregenerative life support experiment "Lunar Palace 365" <b>2022</b> , 17, 4	O
779	Fuzzy Cognitive Maps as a Tool for Identifying Cognitive Conflicts That Hinder the Adoption of Healthy Habits <b>2022</b> , 19,	O
778	Hotspots for social and ecological impacts from freshwater stress and storage loss 2022, 13, 439	6
777	Unlocking the Potential of Deep Learning for Migratory Waterbirds Monitoring Using Surveillance Video. <b>2022</b> , 14, 514	
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774	Integrative time-scale and multi-omic analysis of host-responses to Hop stunt viroid infection.	2
773	Analysis of SNP and SSR markers for the estimates of genetic diversity by using two oilseed Brassica napus populations carrying genome contents of Brassica oleracea.	
77²	The landscape of eating disorders research: A 40-year bibliometric analysis 2022,	O
771	Changes in chemical composition, structural and functional microbiome during alfalfa () ensilage with PS-8 <b>2022</b> , 9, 100-109	1
77°	Breeding Schemes: What Are They, How to Formalize Them, and How to Improve Them?. <b>2021</b> , 12, 791859	O
769	C:N:P stoichiometry in six distinct habitats of a glacier terminus in the source area of the Yangtze River. <b>2022</b> , 158, 181	1
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767	Effects of environmental conditions and jellyfish blooms on small pelagic fish and fisheries from the Western Mediterranean Sea. <b>2022</b> , 264, 107699	O

766	Environmental Factors Driving Seed Hydration Status of Soil Seed Banks and the Implications for Post-fire Recruitment <b>2021</b> , 12, 795003	0
765	Does revegetation cause soil microbiota recovery? Evidence from revisiting a revegetation chronosequence six years after initial sampling.	O
764	Applying biostimulants boosts forage productivity without affecting soil biotic and abiotic parameters on a Central Coast California rangeland.	
763	Should I use fixed effects or random effects when I have fewer than five levels of a grouping factor in a mixed-effects model?. <b>2022</b> , 10, e12794	4
762	Virus Infection and Host Plant Suitability Affect Feeding Behaviors of Cannabis Aphid (Hemiptera: Aphididae), a Newly Described Vector of Potato Virus Y <b>2022</b> ,	O
761	Establishment of a transparent soil system to study Bacillus subtilis chemical ecology.	O
760	Peanut Rotation and Flooding Induce Rhizobacteriome Variation With Opposing Influences on the Growth and Medicinal Yield of <b>2021</b> , 12, 779302	
759	A Bacteriophage Cocktail Significantly Reduces without Deleterious Impact on the Commensal Gut Microbiota under Simulated Gastrointestinal Conditions <b>2022</b> , 14,	1
758	Maintenance of Intraspecific Diversity in Response to Species Competition and Nutrient Fluctuations <b>2022</b> , 10,	
757	Mapping the Green-Lipped Mussel (Perna canaliculus) Microbiome: A Multi-Tissue Analysis of Bacterial and Fungal Diversity <b>2022</b> , 79, 76	O
756	Organic matter and metal loadings influence the spatial gradient of the benthic bacterial community in a temperate estuary. <b>2022</b> ,	O
755	Validation of a deep learning-based image analysis system to diagnose subclinical endometritis in dairy cows <b>2022</b> , 17, e0263409	O
754	Shunt resistance is associated with clinically important outcomes after the Norwood operation. <b>2022</b> ,	O
753	Rapid epidemic expansion of the SARS-CoV-2 Omicron variant in southern Africa 2022,	205
752	Are There Bad ICU Rooms? Temporal Relationship between Patient and ICU Room Microbiome, and Influence on Vancomycin-Resistant Colonization <b>2022</b> , e0100721	O
751	Monocarboxylate Transporters are Involved in Extracellular Matrix Remodelling in Pancreatic Ductal Adenocarcinoma.	1
750	GWAS identifies genetic loci underlying nitrogen responsiveness in the climate resilient C4 model Setaria italica (L.). <b>2022</b> ,	O
749	Arguments for cycling as a mechanism for sustainable modal shifts in Bogot Îl <b>2022</b> , 99, 103291	O

748	Length estimation of Atlantic bluefin tuna (Thunnus thynnus) using vertebrae.	O
747	Allelic and haplotypic data of MHC class II Alu insertions in Ngazidja (Comoros archipelago) and insight on its historical biology <b>2022</b> ,	
746	Microbial Nitrogen Transformation Potential in Sediments of Two Contrasting Lakes Is Spatially Structured but Seasonally Stable <b>2022</b> , e0101321	1
745	Humoral immune response to COVID-19 mRNA vaccination in relation to selenium status <b>2022</b> , 50, 102242	1
744	Insight into the mechanism of indium toxicity in rice <b>2022</b> , 429, 128265	0
743	Resistance to bacterial wilt caused by Ralstonia solanacearum depends on the nutrient condition in soil and applied fertilizers: A meta-analysis. <b>2022</b> , 329, 107874	O
742	Time Series Ovarian Transcriptome Analyses of the Porcine Estrous Cycle Reveals Gene Expression Changes during Steroid Metabolism and Corpus Luteum Development <b>2022</b> , 12,	0
741	Transmission potential of human schistosomes can be driven by resource competition among snail intermediate hosts <b>2022</b> , 119,	2
740	Associations and recovery dynamics of the nasopharyngeal microbiota during influenza-like illness in the aging population <b>2022</b> , 12, 1915	0
739	Climate Stability Index maps, a global high resolution cartography of climate stability from Pliocene to 2100 <b>2022</b> , 9, 48	O
738	Endogenous CRISPR-Cas Systems in Group I and Do Not Directly Target the Botulinum Neurotoxin Gene Cluster <b>2021</b> , 12, 787726	2
737	Polymerase II-Associated Factor 1 Complex-Regulated -Clade Genes Repress Flowering in Response to Chilling <b>2022</b> , 13, 817356	1
736	RNA-seq profiling of white and brown adipocyte differentiation treated with epigallocatechin gallate <b>2022</b> , 9, 41	
735	Size doesnEmatter shape does: A morphological study of pitcher plant in distinct forest canopy structures. <b>2022</b> , 976, 012065	
734	Population Structure and Genetic Diversity Analysis in Sugarcane (Saccharum spp. hybrids) and Six Related Saccharum Species. <b>2022</b> , 12, 412	
733	Microeukaryotic gut parasites in wastewater treatment plants: diversity, activity, and removal <b>2022</b> , 10, 27	O
732	MatchMiner: An open source platform for cancer precision medicine.	
731	DNABERT-based explainable lncRNA identification in plant genome assemblies.	

## (2021-2022)

730	Extracellular electron transfer increases fermentation in lactic acid bacteria via a hybrid metabolism <b>2022</b> , 11,	3
729	Long-term land use in Amazon influence the dynamic of microbial communities in soil and rhizosphere. <b>2022</b> , 21, 100482	O
728	The cannabinoid profile and growth of hemp (Cannabis sativa L.) is influenced by tropical daylengths and temperatures, genotype and nitrogen nutrition. <b>2022</b> , 178, 114605	3
727	Prospective crop yield and income return based on a retrospective analysis of a long-term rainfed agriculture experiment in Nebraska. <b>2022</b> , 198, 103384	O
726	Non-syntrophic methanogenic hydrocarbon degradation by an archaeal species 2021,	10
725	Temperature-responsive optogenetic probes of cell signaling 2021,	4
724	Rapid epidemic expansion of the SARS-CoV-2 Omicron variant in southern Africa.	12
723	Comparative Genomics of Clinical and Environmental Isolates of spp. of Colombia: Implications of Traits Associated with Virulence and Resistance <b>2021</b> , 10,	1
722	Transposons Hidden in Genome Assembly Gaps and Mobilization of Non-Autonomous LTR Retrotransposons Unravelled by Nanotei Pipeline <b>2021</b> , 10,	0
721	Chemical constituents from Linn. leaves as potential cytotoxic, EGFR and aromatase (CYP19A) inhibitors; a study supported by molecular docking <b>2022</b> , 12, 9154-9162	1
720	Archival Mitogenomes Uncover the Synergistic Roles of Environment and Infection in an Amphibian Extinction.	
719	Genetic Variation of Strawberry Fusarium Wilt Pathogen Population in Korea <b>2022</b> , 50, 82-88	1
718	Healthy aging and muscle function are positively associated with NAD+ abundance in humans. <b>2022</b> , 2, 254-263	4
717	Establishing a Prognostic Model Based on Ulceration and Immune Related Genes in Melanoma Patients and Identification of EIF3B as a Therapeutic Target <b>2022</b> , 13, 824946	1
716	Reactive Oxygen Species Differentially Modulate the Metabolic and Transcriptomic Response of Endothelial Cells <b>2022</b> , 11,	2
715	Functional Genomic Analysis of Amphetamine Sensitivity in <b>2022</b> , 13, 831597	
714	Captivity Shifts Gut Microbiota Communities in White-Lipped Deer () <b>2022</b> , 12,	O
713	Transcriptome Co-expression Network and Metabolome Analysis Identifies Key Genes and Regulators of Proanthocyanidins Biosynthesis in Brown Cotton <b>2021</b> , 12, 822198	O

712	Clade-wide variation in bite-force performance is determined primarily by size, not ecology <b>2022</b> , 289, 20212493	1
711	Investigation and characterization of human gut phageome in advanced liver cirrhosis of defined etiologies <b>2022</b> , 14, 9	
710	Unveiling the genomic potential of type strains for discovering new natural products 2022, 8,	O
709	Classification and Regression Models for Genomic Selection of Skewed Phenotypes: A Case for Disease Resistance in Winter Wheat (L.) <b>2022</b> , 13, 835781	O
708	Seasonality and Biting Behavior of Mansonia (Diptera, Culicidae) in Rural Settlements Near Porto Velho, State of Rond <sup>^</sup> Bia, Brazil <b>2022</b> ,	1
707	Comparative Analysis of Seed Transcriptome and Coexpression Analysis Reveal Candidate Genes for Enhancing Seed Size/Weight in <b>2022</b> , 13, 814486	1
706	A deeper look at carrier proteome effects for single-cell proteomics <b>2022</b> , 5, 150	2
705	Towards Heat Tolerant Runner Bean (Phaseolus coccineus L.) by Utilizing Plant Genetic Resources. <b>2022</b> , 12, 612	1
704	Identification of immune-infiltrated hub genes as potential biomarkers of Moyamoya disease by bioinformatics analysis <b>2022</b> , 17, 80	О
703	Electronic Cigarette Use Promotes a Unique Periodontal Microbiome <b>2022</b> , e0007522	1
702	Changes in the Metabolism of Sphingomyelin and Ceramide in the Brain Structures and Spinal Cord of Transgenic Mice (FUS(1-359)) Modeling Amyotrophic Lateral Sclerosis. <b>2022</b> , 48, 178-189	1
701	Lifetime and current depression in the German National Cohort (NAKO). <b>2021</b> , 1-31	3
700	Biosolids for safe land application: does wastewater treatment plant size matters when considering antibiotics, pollutants, microbiome, mobile genetic elements and associated resistance genes?. <b>2022</b> ,	О
699	Genome-based reclassification of the genus Meiothermus along with the proposal of a new genus Allomeiothermus gen. nov <b>2022</b> , 115, 645	2
698	Transcriptional Regulation of Pine Male and Female Cone Initiation and Development: Key Players Identified Through Comparative Transcriptomics <b>2022</b> , 13, 815093	
697	Serum miRNAs as biomarkers for the rare types of muscular dystrophy 2022,	1
696	A community and functional comparison of coral and reef fish assemblages between four decades of coastal urbanisation and thermal stress <b>2022</b> , 12, e8736	О
695	Microbiome Variation Across Populations of Desert Halophyte <b>2022</b> , 13, 841217	

694	When Humans Work Like Beavers: Riparian Restoration Enhances Invertebrate Gamma Diversity and Habitat Heterogeneity.	0
693	Thermal processing reduces PFAS concentrations in blue food - A systematic review and meta-analysis <b>2022</b> , 119081	O
692	Plants with arbuscular mycorrhizal fungi efficiently acquire Nitrogen from substrate additions by shaping the decomposer community composition and their net plant carbon demand. 1	0
691	Response of meiobenthos to migrating ripples in sandy lowland streams. <b>2022</b> , 849, 1905-1921	O
690	Model Integration in Computational Biology: The Role of Reproducibility, Credibility and Utility. <b>2022</b> , 2,	1
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687	IL-22 Plays a Dual Role in the Amniotic Cavity: Tissue Injury and Host Defense against Microbes in Preterm Labor <b>2022</b> , 208, 1595-1615	1
686	Identifying potential drivers of distribution patterns of invasive relative to native freshwater mussels (Unionidae) across spatial scales <b>2022</b> , 12, e8737	0
685	Guitar2: An improved version of the Guitar for sketching the transcriptomic view of RNA-related biological features <b>2022</b> ,	O
684	Genomic signatures of adaptive divergence in lacustrine copepods.	0
683	Alterations to cavefish red blood cells provide evidence of adaptation to reduced subterranean oxygen <b>2022</b> , 12, 3735	1
682	Comparative Genome Analysis Across 128 Isolates Reveal Species-Specific Microsatellite Distribution and Localized Evolution of Compartmentalized Genomes <b>2022</b> , 13, 806398	1
681	Screening the Significant Hub Genes by Comparing Tumor Cells, Normoxic and Hypoxic Glioblastoma Stem-like Cell Lines Using Co-Expression Analysis in Glioblastoma <b>2022</b> , 13,	O
680	Range expansion of Kellicottia bostoniensis (Rousselet, 1908) (Rotifera) throughout a biogeographic boundary between the Brazilian savanna and the Amazon. <b>2022</b> , 84, 1	0
679	Fantastic Beasts: Unfolding Mixoplankton Temporal Variability in the Belgian Coastal Zone Through DNA-Metabarcoding. <b>2022</b> , 9,	O
678	Co-expression network analyses of anthocyanin biosynthesis genes in Ruellia (Wild Petunias; Acanthaceae) <b>2022</b> , 22, 27	1
677	Weaning Time Affects the Archaeal Community Structure and Functional Potential in Pigs <b>2022</b> , 13, 845621	0

Discordant genotype calls across technology platforms elucidate variants with systematic errors in next-generation sequencing. 676

675	Molecular hallmarks of heterochronic parabiosis at single-cell resolution 2022,	5
674	The early life microbiota mediates maternal effects on offspring growth in a nonhuman primate <b>2022</b> , 25, 103948	О
673	Mitochondrial DNA and other lines of evidence clarify species diversity in the Peromyscus truei species group (Cricetidae: Neotominae). <b>2022</b> ,	
672	Monocarboxylate Transporters Are Involved in Extracellular Matrix Remodelling in Pancreatic Ductal Adenocarcinoma <b>2022</b> , 14,	0
671	Metatranscriptomic Analyses Unravel Dynamic Changes in the Microbial and Metabolic Transcriptional Profiles in Artisanal Austrian Hard-Cheeses During Ripening <b>2022</b> , 13, 813480	0
670	Transcriptome Sequencing-Based Mining of Genes Associated With Pubertal Initiation in Dolang Sheep <b>2022</b> , 13, 818810	
669	Range shifts in butternut, a rare, endangered tree, in response to past climate and modern conditions.	0
668	Urbanization comprehensively impairs biological rhythms in coral holobionts 2022,	1
667	Effect of Long-Term Supplementation With Silkworm Pupae Oil on the Methane Yield, Ruminal Protozoa, and Archaea Community in Sheep <b>2022</b> , 13, 780073	0
666	Reproducible Propagation of Species-Rich Soil Bacterial Communities Suggests Robust Underlying Deterministic Principles of Community Formation <b>2022</b> , e0016022	1
665	Genomic architecture of 5S rDNA cluster and its variations within and between species <b>2022</b> , 23, 238	
664	Genetic Diversity and Association Analysis for Carotenoid Content among Sprouts of Cowpea ( L. Walp) <b>2022</b> , 23,	0
663	RGMQL: scalable and interoperable computing of heterogeneous omics big data and metadata in R/Bioconductor <b>2022</b> , 23, 123	O
662	Energy flux across multitrophic levels drives ecosystem multifunctionality: Evidence from nematode food webs. <b>2022</b> , 108656	2
661	Optimal Dose and Type of Exercise to Improve Cognitive Function in Older Adults: A Systematic Review and Bayesian Model-Based Network Meta-Analysis of RCTs <b>2022</b> , 101591	3
660	Discovery of Resistance Genes in Rye by Targeted Long-Read Sequencing and Association Genetics <b>2022</b> , 11,	O
659	Short-term experimental support for bird diversity retention measures during thinning in European boreal forests. <b>2022</b> , 509, 120084	O

658	The genetic architecture underlying body-size traits plasticity over different temperatures and developmental stages in Caenorhabditis elegans <b>2022</b> ,	О
657	Long-Term Stability of Bacterial Associations in a Microcosm of (Chlorophyta, Mamiellophyceae) <b>2022</b> , 13, 814386	
656	High throughput pH bioassay demonstrates pH adaptation of Rhizobium strains isolated from the nodules of Trifolium subterraneum and T. repens <b>2022</b> , 195, 106455	2
655	Responses of the Soil Bacterial Community, Resistome, and Mobilome to a Decade of Annual Exposure to Macrolide Antibiotics <b>2022</b> , e0031622	2
654	A new approach for the classification of grassland utilization in Inner Mongolia based on ecological sites and state-and-transition models. <b>2022</b> , 137, 108733	1
653	Genomics and informatics, conjoined tools vital for understanding and protecting plant health. <b>2021</b> ,	
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651	Screening for suitable mixed microbial consortia from anaerobic sludge and animal dungs for biodegradation of brewery spent grain. <b>2022</b> , 159, 106396	O
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649	Analysis of carbohydrate-active enzymes and sugar transporters in Penicillium echinulatum: A genome-wide comparative study of the fungal lignocellulolytic system <b>2022</b> , 822, 146345	1
648	Depth as a driver of evolution and diversification of ancient squat lobsters (Decapoda, Galatheoidea, Phylladiorhynchus) <b>2022</b> , 107467	1
647	Secondary succession and parent material drive soil bacterial community composition in terraced abandoned olive groves from a Mediterranean hyper-humid mountainous area. <b>2022</b> , 332, 107932	O
646	Temporal variability of Cs concentrations in coastal sediments off Fukushima 2022, 154670	O
645	Bisphenols impact hormone levels in animals: A meta-analysis <b>2022</b> , 154533	O
644	Sugarcane polyphenol and fiber to affect production of short-chain fatty acids and microbiota composition using in vitro digestion and pig faecal fermentation model <b>2022</b> , 385, 132665	2
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638	The epidemic volatility index, a novel early warning tool for identifying new waves in an epidemic. <b>2021</b> , 11, 23775	Ο
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636	Awareness and preparedness of healthcare workers against the first wave of the COVID-19 pandemic: A cross-sectional survey across 57 countries <b>2021</b> , 16, e0258348	1
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632	Differential Transcriptomic Response of Rainbow Trout to Infection with Two Strains of IPNV <b>2021</b> , 14,	О
631	Discovery of a Novel Leaf Rust () Resistance Gene in Rye ( L.) Using Association Genomics <b>2021</b> , 11,	O
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621	Genomic analysis reveals geography rather than culture as the predominant factor shaping genetic variation in northern Kenyan human populations.	
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618	Allele-specific aberration of imprinted domain chromosome architecture associates with large offspring syndrome <b>2022</b> , 25, 104269	О
617	Impact of Climate and Slope Aspects on the Composition of Soil Bacterial Communities Involved in Pedogenetic Processes along the Chilean Coastal Cordillera. <b>2022</b> , 10, 847	1
616	Identification of diffusion routes of O/EA-3 topotype of foot-and-mouth disease virus in Africa and Western Asia between 1974 and 2019 - a phylogeographic analysis <b>2022</b> ,	1
615	Drinking water chlorination has minor effects on the intestinal flora and resistomes of Bangladeshi children <b>2022</b> ,	2
614	Genomic and metabolic adaptations of biofilms to ecological windows of opportunity in glacier-fed streams <b>2022</b> , 13, 2168	1
613	Potential Pleiotropic Genes and Shared Biological Pathways in Epilepsy and Depression Based on GWAS Summary Statistics <b>2022</b> , 2022, 6799285	
612	Identification of novel PHD-finger genes in pepper by genomic re-annotation and comparative analyses <b>2022</b> , 22, 206	0
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610	Therapeutic impact of BET inhibitor BI 894999 treatment: backtranslation from the clinic 2022,	2
609	Morpho-physiology and cannabinoid concentrations of hemp (Cannabis sativa L.) are affected by potassium fertilisers and microbes under tropical conditions. <b>2022</b> , 182, 114907	O
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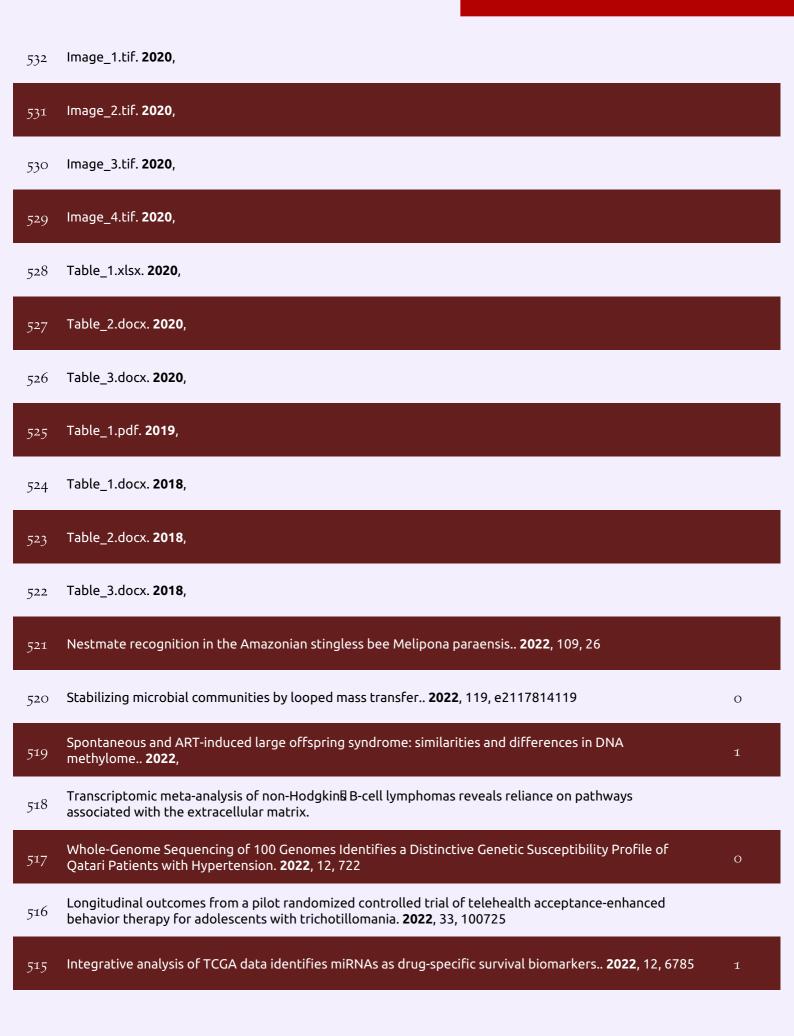
## (2020-2019)

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## (2020-2020)





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513	Extreme mitochondrial DNA divergence underlies genetic conflict over sex determination 2022,	1
512	Little clams with big potential: nutrient release by invasive Corbicula fluminea can exceed co-occurring freshwater mussel (Unionidae) assemblages.	1
511	Plant-plant interactions determine natural restoration of plant biodiversity over time, in a degraded mined land <b>2022</b> , 12, e8878	1
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509	Effects of Different Formulations of Glyphosate on Rumen Microbial Metabolism and Bacterial Community Composition in the Rumen Simulation Technique System <b>2022</b> , 13, 873101	
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507	The evolving SARS-CoV-2 epidemic in Africa: Insights from rapidly expanding genomic surveillance.	O
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505	Sequence Analysis of the Complete Mitochondrial Genome of a Medicinal Plant, Vitex rotundifolia Linnaeus f. (Lamiales: Lamiaceae). <b>2022</b> , 13, 839	o
504	Metabolic Profile and Metabolite Analyses in Extreme Weight Responders to Gastric Bypass Surgery. <b>2022</b> , 12, 417	O
503	Comparing genome-based estimates of relatedness for use in pedigree-based conservation management <b>2022</b> ,	О
502	The role of the primary motor cortex in motor imagery: A theta burst stimulation study 2022, e14077	
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500	Physiological and molecular aspects of seed longevity: exploring intra-species variation in eight Pisum sativum L. accessions <b>2022</b> , e13698	1
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497	Preferential associations of soil fungal taxa under mixed compositions of eastern American tree species <b>2022</b> ,	

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495	The Illicit Ecosystem of Hacking: A Longitudinal Network Analysis of Website Defacement Groups. 08944393	2210978
494	High-resolution 3D spatiotemporal transcriptomic maps of developing Drosophila embryos and larvae <b>2022</b> ,	1
493	Continued Emergence and Evolution of Omicron in South Africa: New BA.4 and BA.5 lineages.	7
492	is a human-adapted lineage in the 'complex <b>2022</b> , 8,	1
491	Alpha and beta phylogenetic diversities jointly reveal ant community assembly mechanisms along a tropical elevational gradient <b>2022</b> , 12, 7728	O
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481	Effect of Prenatal Opioid Exposure on the Human Placental Methylome. <b>2022</b> , 10, 1150	O
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478	Assessing climate change impacts on Pacific salmon and trout using bioenergetics and spatiotemporal explicit river temperature predictions under varying riparian conditions <b>2022</b> , 17, e0266871	1
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436	A parasite outbreak in notothenioid fish in an Antarctic fjord. 2022, 104588	
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430	Can Sphagnum removal reverse the undesired succession of rich fens under different alkalinity and fertility levels?.	O
429	Influence of short and long term processes on SAR11 communities in open ocean and coastal systems.	O
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426	Temporal Analysis of Gene Expression and Isoform Switching in Brown Bears (Ursus arctos).	O
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421	High Serum Vitamin D Concentrations, Induced via Diet, Trigger Immune and Intestinal Microbiota Alterations Leading to Type 1 Diabetes Protection in NOD Mice. 13,	o
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407	Detection of Stable Elite Haplotypes and Potential Candidate Genes of Boll Weight Across Multiple Environments via GWAS in Upland Cotton. 13,	O

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401	Emergence of SARS-CoV-2 Omicron lineages BA.4 and BA.5 in South Africa.	43
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399	Understanding drivers of phylogenetic clustering and terminal branch lengths distribution in epidemics of Mycobacterium tuberculosis. 11,	O
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385	Double, Triple, and Quadruple Jeopardy: Entering Pregnancy With Two or More Multimorbid Diagnoses and Increased Risk of Severe Maternal Morbidity and Postpartum Readmission. <b>2022</b> ,	O
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272	Sugars and Jasmonic Acid Concentration in Root Exudates Affect Maize Rhizosphere Bacterial Communities. <b>2022</b> , 88,	1
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267	IIritonEA Disease-resistant Cantaloupe Hybrid. <b>2022</b> , 57, 1106-1109	O
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257	The Vinca minor genome highlights conserved evolutionary traits in monoterpene indole alkaloid synthesis.	0
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252	Uncovering the Genetic of Cadmium Accumulation in the Rice 3K Panel. 2022, 11, 2813	О
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248	Presence of algal symbionts affects denitrifying bacterial communities in the sea anemone Aiptasia coral model. <b>2022</b> , 2,	O
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246	Mismatch negativity and clinical trajectories in psychotic disorders: Five-year stability and predictive utility. 1-11	О
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243	Multi-omics reveals diet-induced metabolic disorders and liver inflammation via microbiota-gut-liver axis. <b>2022</b> , 109183	O
242	Species distribution modeling of Aedes aegypti in Maricopa County, Arizona from 2014 to 2020. 10,	О
241	Do currents shape global patterns of hybrid richness in coral reef fishes?.	O
240	The South Asian monsoon maintains the disjunction of Rumex hastatus between the western Himalayas and the Hengduan Mountains, southwest China.	О
239	Indoor Air Quality Intervention in Schools: Effectiveness of a Portable HEPA Filter Deployment in Five Schools Impacted by Roadway and Aircraft Pollution Sources. <b>2022</b> , 13, 1623	O
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236	Environmental DNA (eDNA) detects temporal and habitat effects on community composition and endangered species in ephemeral ecosystems: A case study in vernal pools.	О
235	Bilinguals produce language-specific voice onset time in two true-voicing languages.	O
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231	Moderators and processes of change in a pilot randomized controlled trial of acceptance-enhanced behavior therapy for trichotillomania in adolescents. <b>2022</b> , 100757	O
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228	Benthic exometabolites and their ecological significance on threatened Caribbean coral reefs. <b>2022</b> , 2,	O
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226	The ability of Phaeobacter inhibens to produce tropodithietic acid influences the community dynamics of a microalgal microbiome. <b>2022</b> , 2,	0
225	Microbial life strategy with high rRNA operon copy number facilitates the energy and nutrient flux in anaerobic digestion. <b>2022</b> , 226, 119307	o
224	Tailoring and optimizing fatty acid production by oleaginous yeasts through the systematic exploration of their physiological fitness. <b>2022</b> , 21,	О
223	Soybean disease identification using original field images and transfer learning with convolutional neural networks. <b>2022</b> , 203, 107449	О
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221	Development of epigenetic biomarkers with diagnostic and prognostic value to assess the lasting effects of early temperature changes in farmed fish. <b>2023</b> , 563, 738918	O
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219	Drought then wildfire reveals a compound disturbance in a resprouting forest.	O
218	Integrative cross-species analysis of GABAergic neuron cell types and their functions in Alzheimer disease. <b>2022</b> , 12,	О
217	Methylome and transcriptome data integration reveals potential roles of DNA methylation and candidate biomarkers of cow Streptococcus uberis subclinical mastitis. <b>2022</b> , 13,	O
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215	Virulence of Beauveria sp. and Metarhizium sp. fungi towards Fall armyworm (Spodoptera frugiperda).	O
214	Grasping follows Weber's law: How to use response variability as a proxy for JND. <b>2022</b> , 22, 13	О
213	Identification of hub genes regulating the cell activity and function of adipose-derived stem cells under oxygen-glucose deprivation. 9,	O
212	Cross-sectional study of antimicrobial resistance and ecology in gastrointestinal and oral microbial communities of urban Pakistani adults.	O
211	Abnormally Enhanced Midfrontal Theta-Band Activity During Response Monitoring in Youth with Obsessive-Compulsive Disorder. <b>2022</b> ,	o
210	Alternative approaches to identify core bacteria in Fucus distichus microbiome and assess their distribution and host-specificity. <b>2022</b> , 17,	О
209	A chromosomal-scale reference genome of the New World Screwworm, Cochliomyia hominivorax.	O

208	Development of sequencing-based methodologies to distinguish viable from non-viable cells in a bovine milk matrix: A pilot study. 13,	0
207	EGFR-induced suppression of HPV E6/E7 is mediated by microRNA-9-5p silencing of BRD4 protein in HPV-positive head and neck squamous cell carcinoma. <b>2022</b> , 13,	Ο
206	Mimicry promotes morphological diversification but not speciation in mimetic swallowtail butterflies.	0
205	Multivariate analyses reveal key traits in the identification of nitrogen use efficient cultivars of common beans.	O
204	De Novo Hybrid Assembly of the Salvia miltiorrhiza Mitochondrial Genome Provides the First Evidence of the Multi-Chromosomal Mitochondrial DNA Structure of Salvia Species. <b>2022</b> , 23, 14267	1
203	Aquaculture rearing systems induce no legacy effects in Atlantic cod larvae or their rearing water bacterial communities. <b>2022</b> , 12,	O
202	Influence of short and long term processes on SAR11 communities in open ocean and coastal systems. <b>2022</b> , 2,	0
201	Contributions of water-borne corticosterone as one non-invasive biomarker in assessing nitrate pollution stress in tadpoles of Rana temporaria. <b>2023</b> , 331, 114164	O
200	Long-term grazing exclusion affects the population genetics and functional traits of Artemisia frigida in Mongolia. <b>2023</b> , 209, 104900	0
199	Connections between and within extended psychosis and autistic phenotypes and social relationships in the general population. <b>2022</b> ,	O
198	Polymeric structure of the Cannabis sativa L. mitochondrial genome identified with an assembly graph model. <b>2023</b> , 853, 147081	0
197	Finite element and multivariate random forests modelling for stress shield attenuation in customized hip implants. <b>2023</b> , 10, 100151	O
196	Water-use characteristics of two dominant plant species in different community types in the Mu Us Desert. <b>2023</b> , 221, 106803	0
195	A regional-scale study of the contribution of local, management and climate factors to the infestation of processing tomato fields with Amaranthus species. <b>2023</b> , 143, 126722	0
194	Al-DeMat: A web-based expert system platform for computationally expensive models in materials design. <b>2023</b> , 176, 103398	1
193	Meiotic and mitotic aneuploidies drive arrest of in vitro fertilized human preimplantation embryos.	O
192	mTORC1 links pathology in experimental models of Still disease and macrophage activation syndrome. <b>2022</b> , 13,	0
191	The mechanical regulation of RNA binding protein hnRNPC in the failing heart. <b>2022</b> , 14,	O

190	Metal- and Organ-Specific Response to Heavy Metal-Induced Stress Mediated by Antioxidant Enzymes (Activities, Polyamines, and Plant Hormones Levels in Populus deltoides. <b>2022</b> , 11, 3246	2
189	Diversity and genomics of giant viruses in the North Pacific Subtropical Gyre. 13,	Ο
188	Global Expansion of SARS-CoV-2 Variants of Concern: Dispersal Patterns and Influence of Air Travel.	1
187	Abundant fungi dominate the complexity of microbial networks in soil of contaminated site: High-precision community analysis by full-length sequencing. <b>2022</b> , 160563	O
186	Identifying magnetosome-associated genes in the extended CtrA regulon in Magnetospirillum magneticum AMB-1 using a combinational approach.	0
185	Influence of Geochemistry in the Tropical Hot Springs on Microbial Community Structure and Function. <b>2023</b> , 80,	Ο
184	Epigenetic Stress and Long-Read cDNA Sequencing of Sunflower (Helianthus annuus L.) Revealed the Origin of the Plant Retrotranscriptome. <b>2022</b> , 11, 3579	О
183	Spermatozoa, acts as an external cue and alters the cargo and production of the extracellular vesicles derived from oviductal epithelial cells in vitro.	O
182	Genomic imprinting-like monoallelic paternal expression determines sex of channel catfish. 2022, 8,	0
181	Integrated Analysis of Bulk RNA-Seq and Single-Cell RNA-Seq Unravels the Influences of SARS-CoV-2 Infections to Cancer Patients. <b>2022</b> , 23, 15698	O
180	Ozone modified hypothalamic signaling enhancing thermogenesis in the TDP-43A315T transgenic model of Amyotrophic Lateral Sclerosis. <b>2022</b> , 12,	1
179	Human microbiota drives hospital-associated antimicrobial resistance dissemination in the urban environment and mirrors patient case rates. <b>2022</b> , 10,	O
178	Soil Carbon Emission Sources Differ Under Litter and Nutrient Addition During Secondary Succession: Evidence from a Mesocosm Study Using a Three-Transfer-Pool Model.	О
177	Suppression of Streptococcosis and Modulation of the Gut Bacteriome in Nile Tilapia (Oreochromis niloticus) by the Marine Sediment Bacteria Bacillus haynesii and Advenella mimigardefordensis. <b>2022</b> , 10,	O
176	Linking soil microbial community with the changes in soil physicochemical properties in response to long-term agricultural land use change of different chronosequences and depth layers. <b>2022</b> , 145, 109727	0
175	Optogenetic clustering and membrane translocation of the BcLOV4 photoreceptor.	O
174	Variable organization of symbiont-containing tissue across planthoppers hosting different heritable endosymbionts.	О
173	Proteomic-Based Approach Reveals the Involvement of Apolipoprotein A-I in Related Phenotypes of Autism Spectrum Disorder in the BTBR Mouse Model. <b>2022</b> , 23, 15290	O

172	Fermentation Characteristics, Microbial Compositions, and Predicted Functional Profiles of Forage Oat Ensiled with Lactiplantibacillus plantarum or Lentilactobacillus buchneri. <b>2022</b> , 8, 707	0
171	De novo birth of functional microproteins in the human lineage. <b>2022</b> , 41, 111808	O
170	Roadmap to recovery revealed through the reintroduction of an IUCN Red List species.	0
169	Transcriptomic, histological and biochemical analyses of Macrobrachium nipponense response to acute heat stress.	O
168	A comparative analysis of mitochondrial ORFs provides new insights on expansion of mitochondrial genome size in Arcidae. <b>2022</b> , 23,	O
167	Anthropogenic nest materials in two distinct populations of migratory bird in Europe	O
166	CloudSEN12, a global dataset for semantic understanding of cloud and cloud shadow in Sentinel-2. <b>2022</b> , 9,	O
165	ONE-seq: epitranscriptome and gene-specific profiling of NAD-capped RNA.	O
164	De novo genome assembly and genome skims reveal LTRs dominate the genome of a limestone endemic Mountainsnail (Oreohelix idahoensis). <b>2022</b> , 23,	O
163	8-HQA Adjusts the Number and Diversity of Bacteria in the Gut Microbiome of Spodoptera littoralis.	O
162	Ectomycorrhizal Fungi Modulate Pedunculate Oak Heat Stress Responses through the Alternation of Polyamines, Phenolics, and Osmotica Content. <b>2022</b> , 11, 3360	1
161	Vulnerability to climate change increases with trophic level in terrestrial organisms. 2022, 161049	O
160	Is perioperative fast-track management the future of proximal aortic repair?.	1
159	Interplay between Cruciferous Vegetables and the Gut Microbiome: A Multi-Omic Approach. <b>2023</b> , 15, 42	O
158	Dietary supplementation of Bacillus subtilis or antibiotics modified intestinal microbiome of weaned pigs under enterotoxigenic Escherichia coli infection. 13,	O
157	Actl7b-deficiency leads to mislocalization of LC8 type dynein light chains and disruption of murine spermatogenesis.	O
156	Functional and sex-specific dynamics of ectoparasite size evolution in marine isopodfish interactions: Harrison rule and increasing variance.	О
155	Dysregulated ceRNA network modulated by copy number variation-driven lncRNAs in breast cancer: A comprehensive analysis.	O

154	Hydropower dam alters the microbial structure of fish gut in different habitats in upstream and downstream rivers.	O
153	Corrosion-influencing microorganisms in petroliferous regions on a global scale: systematic review, analysis, and scientific synthesis of 16S amplicon metagenomic studies. 11, e14642	1
152	CPGView : A package for visualizing detailed chloroplast genome structures.	9
151	Novel integrative elements and genomic plasticity in ocean ecosystems. <b>2023</b> , 186, 47-62.e16	O
150	Genome-wide association and RNA-seq analyses identify loci for pod orientation in rapeseed (Brassica napus). 13,	0
149	Community risks for SARS-CoV-2 infection among fully vaccinated US adults by rurality: A retrospective cohort study from the National COVID Cohort Collaborative. <b>2023</b> , 18, e0279968	O
148	Expanded catalogue of metagenome-assembled genomes reveals resistome characteristics and athletic performance-associated microbes in horse. <b>2023</b> , 11,	0
147	Genetic diversity and population structure of barley landraces from Southern Ethiopia Gumer district: Utilization for breeding and conservation. <b>2023</b> , 18, e0279737	O
146	Historical ecology of forest garden management in Laxyuubm Tsthsyen and beyond. 2023, 19,	0
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144	8-HQA adjusts the number and diversity of bacteria in the gut microbiome of Spodoptera littoralis. 14,	0
143	Regionalized protein localization domains in the hair cell kinocilium.	0
142	Changes in the Ixodes ricinus microbiome associated with artificial tick feeding. 13,	0
141	Somatic mutation but not aneuploidy differentiates lung cancer in never-smokers and smokers.	o
140	Effect on microbial communities in apple orchard soil when exposed short-term to climate change abiotic factors and different orchard management practices.	0
139	Re-evaluating homoploid reticulate evolution in Helianthus sunflowers.	O
138	Spatio-temporal diversity of dietary preferences and stress sensibilities of early and middle Miocene Rhinocerotidae from Eurasia: impact of climate changes. 3,	0
137	Conceptual combination during novel and existing compound word reading in context: A self-paced reading study.	O

136	Fungi and cercozoa regulate methane-associated prokaryotes in wetland methane emissions. 13,	O
135	The role of unfolded protein response-associated miRNAs in immunogenic cell death amplification: A literature review and bioinformatics analysis. <b>2023</b> , 314, 121341	O
134	Anaerobic digestion using cocoa residues as substrate: Systematic review and meta-analysis. <b>2023</b> , 72, 265-277	1
133	Spatiotemporal evolution of global long-term patterns of soil moisture. <b>2023</b> , 867, 161470	O
132	A hierarchy of mindreading strategies in joint action participation. <b>2021</b> , 16, 844-897	Ο
131	What can Verbal Derivation Tell us about Proper Names?. <b>2022</b> ,	0
130	Do brachycephaly and nose size predict the severity of obstructive sleep apnea (OSA)? A sample-based geometric morphometric analysis of craniofacial variation in relation to OSA syndrome and the role of confounding factors.	О
129	Flower transcriptional response to long term hot and cold environments in Antirrhinum majus. 14,	O
128	TCR_Explore: A novel webtool for T cell receptor repertoire analysis. <b>2023</b> , 21, 1272-1282	Ο
127	Effects of forest fragment on avian breeding activity.	O
126	Cross-domain interactions induce community stability to benthic biofilms in proglacial streams.	0
125	Ontogenetic variability of the intertympanic sinus distinguishes lineages within Crocodylia.	O
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123	A Cross-Sectional Study of Potential Antimicrobial Resistance and Ecology in Gastrointestinal and Oral Microbial Communities of Young Normoweight Pakistani Individuals. <b>2023</b> , 11, 279	O
122	Reintroducing genetic diversity in populations from cryopreserved material: The case of Abondance, a French local dairy cattle breed.	0
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116	Addressing the pervasive scarcity of structural annotation in eukaryotic algae. 2023, 13,	1
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113	Dietary niche reconstruction of Pliocene and Pleistocene Equidae from the Linxia Basin of northwestern China based on stable isotope analysis. <b>2023</b> , 111416	O
112	Genomic view of the diversity and functional role of archaea and bacteria in the skeleton of the reef-building corals Porites lutea and Isopora palifera. <b>2022</b> , 12,	1
111	Distinct microbiome profiles in convergent wisdom tooth impactions. 2023, 134,	0
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108	Commercial production of Florida pompano (Trachinotus carolinus) larvae at low salinity induces variable changes in whole-larvae microbial diversity, gene expression, and gill histopathology. 10,	0
107	Ancient Fish Lineages Illuminate Toll-Like Receptor Diversification in Early Vertebrate Evolution.	O
106	A comprehensive metabolic map reveals major quality regulations in red-flesh kiwifruit ( Actinidia chinensis ).	O
105	Transcriptionally active nitrogen fixation and biosynthesis of diverse secondary metabolites by Dolichospermum and Aphanizomenon-like Cyanobacteria in western Lake Erie Microcystis blooms. <b>2023</b> , 124, 102408	O
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103	Keeping in the lane! Investigating drivers[performance handling silent vs. alerted lateral control failures in monotonous partially automated driving. <b>2023</b> , 95, 103429	O
102	Disturbance of primary producer communities disrupts the thermal limits of the associated aquatic fauna. <b>2023</b> , 872, 162135	О
101	Short-term prescribed fire-induced changes in soil microbial communities and nutrients in native rangelands of Florida. <b>2023</b> , 189, 104914	O

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86	Forest structure predicts species richness and functional diversity in Amazonian mixed-species bird flocks. <b>2023</b> , 55, 467-479	O
85	European oak metabolites shape digestion and fitness of the herbivore Tortrix viridana.	O
84	A bacteriocyte symbiont determines whitefly sex ratio by regulating mitochondrial function. <b>2023</b> , 42, 112102	0
83	Ad^ lie penguins north and east of the Ad^ lie gapleontinue to thrive in the face of dramatic declines elsewhere in the Antarctic Peninsula region. <b>2023</b> , 13,	O

82	Aplicaciones interactivas basadas en el paquete SHINY/R para explicar conceptos estad <sup>^</sup> Eticos. <b>2023</b> , 17, 1-15	O
81	Time-dependent recruitment of GAF, ISGF3 and IRF1 complexes shapes IFN and IFN activated transcriptional responses and explains mechanistic and functional overlap.	О
80	Putative chemical cue from Gyrodactylus-infected guppies subtly alters behaviour but prior exposure decreases parasite intensity. <b>2023</b> , 150, 434-445	О
79	Host Determinants of Fungal Species Composition and Symptom Manifestation in the Sorghum Grain Mold Disease Complex. <b>2023</b> , 107, 315-325	О
78	Ultra-small bacteria and archaea exhibit genetic flexibility towards groundwater oxygen content, and adaptations for attached or planktonic lifestyles. <b>2023</b> , 3,	О
77	Whole-Genome Comparison Reveals Structural Variations behind Heading Leaf Trait in Brassica oleracea. <b>2023</b> , 24, 4063	О
76	Effects of dielectric barrier discharge plasma and plasma-activated water on the surface microbial diversity of fresh goji berries during storage. <b>2023</b> , 313, 111920	О
75	Seasonal and genetic effects on lipid profiles of juvenile Atlantic salmon.	O
74	Autonomous sensory meridian response is associated with a larger heartbeat-evoked potential amplitude without differences in interoceptive awareness.	O
73	Assessing the viability of estimating baleen whale abundance from tourist vessels. 10,	O
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70	Aging Atlas Reveals Cell-Type-Specific Regulation of Pro-longevity Strategies.	О
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65	Increase in Linguistic Complexity in Older Adults During COVID-19. 1-19	O

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61	Surface Bacterioplankton Community Structure Crossing the Antarctic Circumpolar Current Fronts. <b>2023</b> , 11, 702	O
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54	Biological soil crust development affects bacterial communities in the Caragana microphylla community in alpine sandy areas. 14,	О
53	Excess labile carbon promotes diazotroph abundance in heat-stressed octocorals. 2023, 10,	1
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51	Reproducible chemostat cultures to eliminate eukaryotic viruses from fecal transplant material.	О
50	The distribution of the Lansing Effect across animal species. <b>2023</b> , 77, 608-615	0
49	Predicting diversity in benthic macro-scale communities associated with mussel matrices in three Pacific ecoregions. <b>2023</b> , 44,	O
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44	Relatedness within colonies of three North American species of carpenter ants (Subgenus: Camponotus) and a comparison with relatedness estimates across Formicinae.	0
43	A reproducible enteric phage community improves blood glucose regulation in an obesity mouse model.	o
42	Blood flow restricted walking alters gait kinematics. 1-10	0
41	Conditional Forest Models Built Using Metagenomic Data Accurately Predicted Salmonella Contamination in Northeastern Streams. <b>2023</b> , 11,	o
40	Variable organization of symbiont-containing tissue across planthoppers hosting different heritable endosymbionts. 14,	0
39	High nucleotide diversity accompanies differential DNA methylation in naturally diverging populations.	o
38	Fine-scale mapping of physicochemical and microbial landscapes of the coral skeleton.	0
37	High landscape-scale forest cover favours cold-adapted plant communities in agriculture <b>f</b> orest mosaics.	o
36	Comparative Study of Transcriptome Profile and Immune-related Genes Network Associated with Intestinal Epithelial Tissue based on Microarray Data in Poultry with Coccidiosis. <b>2022</b> , 13, 154-162	O
35	Highly accurate long reads are crucial for realizing the potential of biodiversity genomics. 2023, 24,	O
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33	Emerging investigator series: differential effects of carbon nanotubes and graphene on the tomato rhizosphere microbiome.	o
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29	Impacts of anthropogenic climate change on tropical montane forests: an appraisal of the evidence.	o

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25	T9GPred: A Comprehensive Computational Tool for the Prediction of Type 9 Secretion System, Gliding Motility and the Associated Secreted Proteins.	O
24	Spatial and seasonal variation in thermal sensitivity within North American bird species.	0
23	The distribution of vertebrate roadkill varies by season, surrounding environment, and animal class. <b>2023</b> , 69,	O
22	Seasonal changes in photoperiod and temperature lead to changes in cuticular hydrocarbon profiles and affect mating success in Drosophila suzukii. <b>2023</b> , 13,	0
21	Differential gene expression underlying the biosynthesis of Dufour's gland signals in Bombus impatiens. <b>2023</b> , 3, 100056	0
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19	Integrative taxonomy using the plant core DNA barcodes in Sumatra's Burseraceae. <b>2023</b> , 13,	O
18	Environmental drivers of spatial and temporal water quality variability in four coastal wetlands of Lake Ontario. <b>2023</b> ,	O
17	Response of Soil Microbial Communities to Elevation Gradient in Central Subtropical Pinus taiwanensis and Pinus massoniana Forests. <b>2023</b> , 14, 772	O
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7	Reintroducing genetic diversity in populations from cryopreserved material: the case of Abondance, a French local dairy cattle breed. <b>2023</b> , 55,	O
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3	Field scale biodegradation of total petroleum hydrocarbons and soil restoration by Ecopiles: microbiological analysis of the process. 14,	O
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