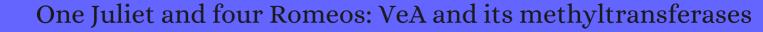
CITATION REPORT List of articles citing



DOI: 10.3389/fmicb.2015.00001 Frontiers in Microbiology, 2015, 6, 1.

Source: https://exaly.com/paper-pdf/62329186/citation-report.pdf

Version: 2024-04-05

This report has been generated based on the citations recorded by exaly.com for the above article. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

#	Paper	IF	Citations
1021	Distinct RNAi Pathways in the Regulation of Physiology and Development in the Fungus Mucor circinelloides. 2015 , 91, 55-102		16
1020	Novel Signal-Enhancing Approaches for Optical Detection of Nucleic Acids © oing beyond Target Amplification. 2015 , 3, 224-240		9
1019	Synthetic biology of fungal natural products. <i>Frontiers in Microbiology</i> , 2015 , 6, 775	5.7	30
1018	Levan Enhances Associated Growth of Bacteroides, Escherichia, Streptococcus and Faecalibacterium in Fecal Microbiota. 2015 , 10, e0144042		38
1017	The RNAi machinery controls distinct responses to environmental signals in the basal fungus Mucor circinelloides. 2015 , 16, 237		35
1016	Isolation and Characterization of a Novel Virulent Phage of Lactobacillus casei ATCC 393. 2015 , 7, 333-4	∤1	11
1015	Epigenetics as an emerging tool for improvement of fungal strains used in biotechnology. 2015 , 99, 610	57-81	30
1014	The effect of mineral and organic nutrient input on yields and nitrogen balances in western Kenya. 2015 , 214, 10-20		15
1013	Effects of 3-nitrooxypropanol on methane production using the rumen simulation technique (Rusitec). 2015 , 209, 98-109		14
1012	Regulation of secondary metabolite production in the fungal tomato pathogen Cladosporium fulvum. 2015 , 84, 52-61		16
1011	Use of UHPLC high-resolution Orbitrap mass spectrometry to investigate the genes involved in the production of secondary metabolites in Aspergillus flavus. 2015 , 32, 1656-73		13
1010	Analysis of the xynB5 gene encoding a multifunctional GH3-BglX Eglucosidase-Ekylosidase-Barabinosidase member in Caulobacter crescentus. 2015 , 108, 993-1007		9
1009	Nickel has biochemical, physiological, and structural effects on the green microalga Ankistrodesmus falcatus: An integrative study. 2015 , 169, 27-36		36
1008	The overproduction of Synechocystis sp. PCC 6803 heat-shock protein (Sll0170) protects Escherichia coli against high-temperature stress. 2015 , 29, 1201-1207		1
1007	The role of the genetic elements bla oxa and IS Aba 1 in the Acinetobacter calcoaceticus-Acinetobacter baumannii complex in carbapenem resistance in the hospital setting. 2016 , 49, 433-40		16
1006	Isolation and selection of plant growth-promoting bacteria associated with sugarcane. 2016 , 46, 149-15		33
1005	Survival and Evolution of a Large Multidrug Resistance Plasmid in New Clinical Bacterial Hosts. 2016 , 33, 2860-2873		130

(2016-2016)

10	004	Effects of Propidium Monoazide (PMA) Treatment on Mycobiome and Bacteriome Analysis of Cystic Fibrosis Airways during Exacerbation. 2016 , 11, e0168860	17
10	003	Proteomics and Metabolomics Analyses to Elucidate the Desulfurization Pathway of Chelatococcus sp. 2016 , 11, e0153547	22
10	002	Antifungal Imidazole-Decorated Cationic Amphiphiles with Markedly Low Hemolytic Activity. 2016 , 22, 11148-51	18
10	001	A phosphorylation code of the Aspergillus nidulans global regulator VelvetA (VeA) determines specific functions. 2016 , 99, 909-24	22
10	000	Functional metagenomic selection of ribulose 1, 5-bisphosphate carboxylase/oxygenase from uncultivated bacteria. 2016 , 18, 1187-99	21
9	99	Hepatitis C in Cameroon: What is the progress from 2001 to 2016?. 2016 , 4, 162-169	10
9:	98	Dose-response effect of nitrate on hydrogen distribution between rumen fermentation end products: an in vitro approach. 2016 , 56, 224	12
9	97	Bacteria-type-specific biparental immune priming in the pipefish. 2016 , 6, 6735-6757	9
9	96	Metagenomic approach to characterize soil microbial diversity of Phumdi at Loktak Lake. 2016 , 74, 2075-2086	16
9	95	Methicillin resistant Staphylococcus aureus in Ethiopia: a meta-analysis. 2016 , 16, 689	23
9	94	Marinopyrones A D , ⊕yrones from marine-derived actinomycetes of the family Nocardiopsaceae. 2016 , 57, 1997-2000	12
9	93	Effects of macrophyte-associated nitrogen cycling bacteria on ANAMMOX and denitrification in river sediments in the Taihu Lake region of China. 2016 , 93, 82-90	21
9:	92	Chinese herb medicine against Sortase A catalyzed transformations, a key role in gram-positive bacterial infection progress. 2016 , 31, 184-196	10
9	91	Awakening of Fungal Secondary Metabolite Gene Clusters. 2016 , 253-273	18
9	90	Dibenzothiophene desulfurization capability and evolutionary divergence of newly isolated bacteria. 2016 , 198, 509-19	20
9	89	Chitosan nanoparticles in drug therapy of infectious and inflammatory diseases. 2016 , 13, 1177-94	65
9	88	Biology of the Whiskey Fungus. 2016 , 413-428	Ο
9	87	Development of a fluorescent transgenic zebrafish biosensor for sensing aquatic heavy metal pollution. 2016 , 25, 617-27	13

986	Influence of periparturient and postpartum diets on rumen methanogen communities in three breeds of primiparous dairy cows. 2016 , 16, 78	18
985	A review of the ecology, ecophysiology and biodiversity of microalgae in Arctic soil crusts. 2016 , 39, 2227-22	4039
984	Effect of effluent recirculation rate on the performance of anaerobic bio-filter treating coal gasification wastewater under co-digestion conditions. 2016 , 6, 87926-87934	10
983	Generation of acid mine drainage around the Karaerik copper mine (Espiye, Giresun, NE Turkey): implications from the bacterial population in the Acšu effluent. 2016 , 20, 673-85	7
982	Unraveling the dose-response puzzle of : A mechanistic approach. 2016 , 1, 101-114	5
981	Complete Genome Sequence of the MRSA Isolate HC1335 from ST239 Lineage Displaying a Truncated AgrC Histidine Kinase Receptor. 2016 , 8, 3187-3192	8
980	Toward understanding long-distance extracellular electron transport in an electroautotrophic microbial community. 2016 , 9, 3544-3558	50
979	Chemical and physical strategies in onychomycosis topical treatment: A review. 2017 , 55, 461-475	11
978	Protein phosphorylation and its role in archaeal signal transduction. 2016 , 40, 625-47	51
977	Methane emissions, ruminal characteristics and nitrogen utilisation changes after refaunation of protozoa-free sheep. 2016 , 144, 48-55	9
976	Biogenic methane in shale gas and coal bed methane: A review of current knowledge and gaps. 2016 , 165, 106-120	67
975	Tetraodon nigroviridis: A model of Vibrio parahaemolyticus infection. 2016 , 56, 388-396	10
974	Aspergillus fumigatus spore proteomics and genetics reveal that VeA represses DefA-mediated DNA damage response. 2016 , 148, 26-35	2
973	Effects of 3-nitrooxypropanol and monensin on methane production using a forage-based diet in Rusitec fermenters. 2016 , 220, 67-72	9
972	Phialomyces macrosporus decreases anthracnose severity on coffee seedlings by competition for nutrients and induced resistance. 2016 , 103, 119-128	3
971	Assessment of Genetic Diversity in Indigenous Plants from Northeast India Using Molecular Marker Technology. 2016 , 181-192	
970	Altered antibiotic tolerance in anaerobic digesters acclimated to triclosan or triclocarban. 2016 , 163, 22-26	17
969	Intrinsic factors of Peltigera lichens influence the structure of the associated soil bacterial microbiota. 2016 , 92,	11

(2016-2016)

968	origins of endolithic algae. 2016 , 6, 31508	65
967	Hierarchical effects of pro-inflammatory cytokines on the post-influenza susceptibility to pneumococcal coinfection. 2016 , 6, 37045	35
966	Fungal endophytes for sustainable crop production. 2016 , 92,	166
965	RNA regulators responding to ribosomal protein S15 are frequent in sequence space. 2016 , 44, 9331-9341	3
964	Rapid colorimetric sensing platform for the detection of Listeria monocytogenes foodborne pathogen. 2016 , 86, 1061-1066	77
963	Biogas Production: Microbiology and Technology. 2016 , 156, 195-234	32
962	How do toxic metals affect harmful cyanobacteria? An integrative study with a toxigenic strain of Microcystis aeruginosa exposed to nickel stress. 2016 , 133, 36-46	14
961	Perspectives on polymeric nanostructures for the therapeutic application of antimicrobial peptides. 2016 , 11, 1729-44	32
960	Regulation of Secondary Metabolism in the Gray Mold Fungus Botrytis cinerea. 2016, 201-216	6
959	Genome-Wide Chromatin Immunoprecipitation Sequencing Analysis of the Penicillium chrysogenum Velvet Protein PcVelA Identifies Methyltransferase PcLlmA as a Novel Downstream Regulator of Fungal Development. 2016 , 1,	7
958	Novel Genomic and Evolutionary Insight of WRKY Transcription Factors in Plant Lineage. 2016 , 6, 37309	38
957	Characterization of the resistome in manure, soil and wastewater from dairy and beef production systems. 2016 , 6, 24645	63
956	Overexpression of the Global Regulator LaeA in Chaetomium globosum Leads to the Biosynthesis of Chaetoglobosin Z. 2016 , 79, 2487-2494	26
955	Phylogenetical coherence of Pseudomonas in unexplored soils of Himalayan region. 2016 , 6, 170	3
954	Competition and coexistence between a syntrophic consortium and a metabolic generalist, and its effect on productivity. 2016 , 404, 348-360	7
953	Sulfated levan from Halomonas smyrnensis as a bioactive, heparin-mimetic glycan for cardiac tissue engineering applications. 2016 , 149, 289-96	47
952	Insights into the single cell draft genome of "Candidatus Achromatium palustre". 2016 , 11, 28	6
951	Stachybotrys mycotoxins: from culture extracts to dust samples. 2016 , 408, 5513-26	13

950	Cellulases and beyond: the first 70 years of the enzyme producer Trichoderma reesei. 2016 , 15, 106	280
949	MBsbauer spectroscopy and X-ray fluorescence studies on sediments from the methanic zone of the Helgoland mud area, North Sea. 2016 , 237, 1	
948	Copper pollution decreases the resistance of soil microbial community to subsequent dry-rewetting disturbance. 2016 , 39, 155-164	17
947	Inactivation of Bacillus amyloliquefaciens spores by continuous high-pressure-assisted thermal sterilization in an oil-in-water (o/w) emulsion with 10 % soybean oil. 2016 , 242, 935-942	9
946	RNAi pathways in Mucor: A tale of proteins, small RNAs and functional diversity. 2016 , 90, 44-52	24
945	Inactivation of the global regulator LaeA in Monascus ruber results in a species-dependent response in sporulation and secondary metabolism. 2016 , 120, 297-305	32
944	Next-Generation Sequencing in the Mycology Lab. 2016 , 10, 37-42	32
943	Effect of Evolutionary Adaption on Xylosidase Activity in Thermotolerant Yeast Isolates Kluyveromyces marxianus NIRE-K1 and NIRE-K3. 2016 , 179, 1143-54	11
942	Enhancement in xylose utilization using Kluyveromyces marxianus NIRE-K1 through evolutionary adaptation approach. 2016 , 39, 835-43	28
941	Vector competence of Aedes vexans (Meigen), Culex poicilipes (Theobald) and Cx. quinquefasciatus Say from Senegal for West and East African lineages of Rift Valley fever virus. 2016 , 9, 94	30
940	Climate change: Production performance, health issues, greenhouse gas emissions and mitigation strategies in sheep and goat farming. 2016 , 135, 50-59	54
939	The phage P.E1 isolated from hospital sewage reduces the growth of Escherichia coli. 2016 , 26, 181-188	14
938	Effect of rice straw application on microbial community and activity in paddy soil under different water status. 2016 , 23, 5941-8	24
937	Linear antimicrobial peptides with activity against herpes simplex virus 1 and Aichi virus. 2017 , 108, e22871	21
936	The evolution of no-cost resistance at sub-MIC concentrations of streptomycin in Streptomyces coelicolor. 2017 , 11, 1168-1178	34
935	NMR-based metabolomics of transgenic and non-transgenic sweet orange reveals different responses in primary metabolism during citrus canker development. 2017 , 13, 1	8
934	Composition and copper binding properties of aquatic fulvic acids in eutrophic Taihu Lake, China. 2017 , 172, 496-504	32
933	Retrospective study of clinical and lesion characteristics of patients undergoing surgical treatment for Pulmonary Tuberculosis in Georgia. 2017 , 56, 200-207	5

932	Human pathogens in plant biofilms: Formation, physiology, and detection. 2017 , 114, 1403-1418	13
931	Bacterial and Fungal Proteolytic Enzymes: Production, Catalysis and Potential Applications. 2017 , 183, 1-19	74
930	Enhanced Biotransformation of Triclocarban by Ochrobactrum sp. TCC-1 Under Anoxic Nitrate Respiration Conditions. 2017 , 74, 491-498	13
929	Symbiotic performance, nitrogen flux and growth of lima bean (Phaseolus lunatus L.) varieties inoculated with different indigenous strains of rhizobia. 2017 , 73, 117-124	7
928	Mechanistic Investigation on ROS Resistance of Phosphorothioated DNA. 2017, 7, 42823	14
927	Development of fungal cell factories for the production of secondary metabolites: Linking genomics and metabolism. 2017 , 2, 5-12	66
926	Advances in industrial microbiome based on microbial consortium for biorefinery. 2017, 4, 11	43
925	Efficiency of Ohmic assisted hydrodistillation for the extraction of essential oil from oregano (Origanum vulgare subsp. viride) spices. 2017 , 41, 172-178	64
924	Development and evaluation of a trickle bed bioreactor for enhanced mass transfer and methanol production from biogas. 2017 , 122, 103-114	24
923	Biodeterioration. 2017 , 1-30	3
923 922	Biodeterioration. 2017, 1-30 A review on microbiological decontamination of fresh produce with nonthermal plasma. 2017, 122, 1438-145	
922	A review on microbiological decontamination of fresh produce with nonthermal plasma. 2017 , 122, 1438-145 Influence of agricultural activities in the structure and metabolic functionality of paramo soil	5 ₇₇
922	A review on microbiological decontamination of fresh produce with nonthermal plasma. 2017, 122, 1438-145 Influence of agricultural activities in the structure and metabolic functionality of paramo soil samples in Colombia studied using a metagenomics analysis in dynamic state. 2017, 351, 63-76	5 ₇₇
922 921 920	A review on microbiological decontamination of fresh produce with nonthermal plasma. 2017, 122, 1438-145 Influence of agricultural activities in the structure and metabolic functionality of paramo soil samples in Colombia studied using a metagenomics analysis in dynamic state. 2017, 351, 63-76 Alleviation of environmental stress in plants: The role of beneficial Pseudomonas spp 2017, 47, 372-407 Involvement of a velvet protein ClVelB in the regulation of vegetative differentiation, oxidative	5 ₇₇ 6 24
922 921 920 919	A review on microbiological decontamination of fresh produce with nonthermal plasma. 2017, 122, 1438-145 Influence of agricultural activities in the structure and metabolic functionality of paramo soil samples in Colombia studied using a metagenomics analysis in dynamic state. 2017, 351, 63-76 Alleviation of environmental stress in plants: The role of beneficial Pseudomonas spp 2017, 47, 372-407 Involvement of a velvet protein CIVelB in the regulation of vegetative differentiation, oxidative stress response, secondary metabolism, and virulence in Curvularia lunata. 2017, 7, 46054 From Nano to Micro: using nanotechnology to combat microorganisms and their multidrug	5 ₇₇ 6 24
922921920919918	A review on microbiological decontamination of fresh produce with nonthermal plasma. 2017, 122, 1438-145 Influence of agricultural activities in the structure and metabolic functionality of paramo soil samples in Colombia studied using a metagenomics analysis in dynamic state. 2017, 351, 63-76 Alleviation of environmental stress in plants: The role of beneficial Pseudomonas spp 2017, 47, 372-407 Involvement of a velvet protein ClVelB in the regulation of vegetative differentiation, oxidative stress response, secondary metabolism, and virulence in Curvularia lunata. 2017, 7, 46054 From Nano to Micro: using nanotechnology to combat microorganisms and their multidrug resistance. 2017, 41, 302-322 Bacilli as Biological Nano-factories Intended for Synthesis of Silver Nanoparticles and Its	5 ₇₇ 6 24 10 116

914	Characterization of non-dialyzable constituents from cranberry juice that inhibit adhesion, co-aggregation and biofilm formation by oral bacteria. 2017 , 8, 1955-1965	18
913	BusyBee Web: metagenomic data analysis by bootstrapped supervised binning and annotation. 2017 , 45, W171-W179	43
912	The Trichodesmium consortium: conserved heterotrophic co-occurrence and genomic signatures of potential interactions. 2017 , 11, 1813-1824	38
911	Citrate-Coated Silver Nanoparticles Growth-Independently Inhibit Aflatoxin Synthesis in Aspergillus parasiticus. 2017 , 51, 8085-8093	30
910	Ecological Aspects on Rumen Microbiome. 2017 , 367-389	O
909	A niche for cyanobacteria producing chlorophyll f within a microbial mat. 2017 , 11, 2368-2378	40
908	Bacterioplankton assemblages in coastal ponds reflect the influence of hydrology and geomorphological setting. 2017 , 93,	4
907	Fermented Vegetables. 2017 , 537-584	5
906	Synergy between azoles and 1,4-dihydropyridine derivative as an option to control fungal infections. 2017 , 110, 1219-1226	2
905	Optimization and characterization of a murine lung infection model for the evaluation of novel therapeutics against Burkholderia cenocepacia. 2017 , 139, 181-188	2
904	Dissolved copper (dCu) biogeochemical cycling in the subarctic Northeast Pacific and a call for improving methodologies. 2017 , 196, 47-61	24
903	Genetic engineering of Trichoderma reesei cellulases and their production. 2017 , 10, 1485-1499	122
902	Screening of enzymatic activities within different enological non- yeasts. 2017, 54, 1555-1564	39
901	Transcriptome analysis of the two unrelated fungal flactam producers Acremonium chrysogenum and Penicillium chrysogenum: Velvet-regulated genes are major targets during conventional strain improvement programs. 2017 , 18, 272	13
900	Presence of Oenococcus oeni and other lactic acid bacteria in grapes and wines from Priorat (Catalonia, Spain). 2017 , 81, 326-334	21
899	Isolation, Identification And Screening Antibacterial Activity from Marine Sponge-Associated Fungi Against Multidrug-Resistant (MDR)Escherichia coli. 2017 , 55, 012028	3
898	Toxicological features of maleilated polyflavonoids from Pinus radiata (D. Don.) as potential functional additives for biomaterials design. 2017 , 109, 1069-1078	2
897	Prokaryotic Abundance and Activity in Permafrost of the Northern Victoria Land and Upper Victoria Valley (Antarctica). 2017 , 74, 402-415	13

(2017-2017)

896	Inactivation of Staphylococcus saprophyticus in chicken meat and purge using thermal processing, high pressure processing, gamma radiation, and ultraviolet light (254 nm). 2017 , 75, 78-82	12
895	The Effect of Membrane Type on the Performance of Microbial Electrosynthesis Cells for Methane Production. 2017 , 164, H3015-H3023	24
894	Engineering photosynthetic production of L-lysine. 2017 , 44, 273-283	25
893	Effect of an active label based on benzyl isothiocyanate on the morphology and ochratoxins production of Aspergillus ochraceus. 2017 , 101, 61-72	7
892	Model microbial communities for ecosystems biology. 2017 , 6, 51-57	21
891	Dehydration and rehydration - induced temporal changes in cytosolic and membrane proteome of the nitrogen fixing cyanobacterium Anabaena sp. PCC 7120. 2017 , 27, 244-258	10
890	Processing of baby food using pressure-assisted thermal sterilization (PATS) and comparison with thermal treatment. 2017 , 37, 579-593	4
889	Antimicrobial edible coatings and films from micro-emulsions and their food applications. 2017 , 263, 9-16	34
888	Microbial Functions of the Rhizosphere. 2017 , 177-210	10
887	Mixed sulfate-reducing bacteria-enriched microbial fuel cells for the treatment of wastewater containing copper. 2017 , 189, 134-142	58
886	The development of seaweed-derived bioactive compounds for use as prebiotics and nutraceuticals using enzyme technologies. 2017 , 70, 20-33	65
885	Co-inoculum of Beauveria brongniartii and B. bassiana shows in vitro different metabolic behaviour in comparison to single inoculums. 2017 , 7, 13102	10
884	Anti-candidal Activity and Chemical Composition of Essential Oil of Clove (Syzygium aromaticum). 2017 , 20, 951-958	11
883	Small-scale variation of ammonia oxidisers within intertidal sediments dominated by ammonia-oxidising bacteria Nitrosomonas sp. amoA genes and transcripts. 2017 , 7, 13200	12
882	Phytoplankton functional diversity increases ecosystem productivity and stability. 2017, 361, 184-196	43
881	Prospective of Microbial Exopolysaccharide for Heavy Metal Exclusion. 2017 , 183, 582-600	50
880	Interventions for promoting participation in shared decision-making for children and adolescents with cystic fibrosis. 2017 ,	1
879	Seasonality can induce coexistence of multiple bet-hedging strategies in Dictyostelium discoideum via storage effect. 2017 , 426, 104-116	17

878	Potential biotechnological capabilities of cultivable mycobiota from carwash effluents. 2017, 6, e00498	7
877	Links between Natural Variation in the Microbiome and Host Fitness in Wild Mammals. 2017 , 57, 756-769	40
876	Micro-electrolysis/retinervus luffae-based simultaneous autotrophic and heterotrophic denitrification for low C/N wastewater treatment. 2017 , 24, 16651-16658	11
875	Food Hazards and Quality Control in Table Olive Processing with a Special Reference to Functional Compounds. 2017 , 343-352	
874	Taxonomic and functional patterns across soil microbial communities of global biomes. 2017 , 609, 1064-1074	24
873	Factors involved in Bacillus spore's resistance to cold atmospheric pressure plasma. 2017 , 43, 173-181	23
872	Nonsteroidal Anti-inflammatory Drugs (NSAIDS) Inhibit the Growth and Reproduction of Chaetomium globosum and Other Fungi Associated with Water-Damaged Buildings. 2017 , 182, 1025-1036	3
871	Single-cell sequencing unveils the lifestyle and CRISPR-based population history of Hydrotalea sp. in acid mine drainage. 2017 , 26, 5541-5551	6
870	A glimpse into the future - new therapeutic targets could transform the way we treat staphylococcal infections. 2017 , 8, 1508-1510	2
869	Genomic Characterisation of Starter Cultures and Probiotic Bacteria. 2017 , 37-65	
868	Autoinhibitory sterol sulfates mediate programmed cell death in a bloom-forming marine diatom. 2017 , 8, 1292	44
867	Vaccine candidate discovery for the next generation of malaria vaccines. 2017 , 152, 195-206	26
866	Polyphenols and tri-terpenoids from Olea europaea L. in alleviation of enteric pathogen infections through limiting bacterial virulence and attenuating inflammation. 2017 , 36, 132-143	23
865	Hanseniaspora uvarum enhanced with trehalose induced defense-related enzyme activities and relative genes expression levels against Aspergillus tubingensis in table grapes. 2017 , 132, 162-170	31
864	Single Cell Transcriptomics, Mega-Phylogeny, and the Genetic Basis of Morphological Innovations in Rhizaria. 2017 , 34, 1557-1573	29
863	6-methylmercaptopurine riboside, a thiopurine nucleoside with antiviral activity against canine distemper virus in vitro. 2017 , 14, 124	3
862	The deep, hot biosphere: Twenty-five years of retrospection. 2017 , 114, 6895-6903	49
861	Formation, physiology, ecology, evolution and clinical importance of bacterial persisters. 2017 , 41, 219-251	179

(2017-2017)

860	Thermostabilization of the uronate dehydrogenase from Agrobacterium tumefaciens by semi-rational design. 2017 , 7, 103	10
859	The Aspergillus nidulans Velvet-interacting protein, VipA, is involved in light-stimulated heme biosynthesis. 2017 , 105, 825-838	7
858	Free nitrous acid and pH determine the predominant ammonia-oxidizing bacteria and amount of NO in a partial nitrifying reactor. 2017 , 101, 1673-1683	29
857	Diversity and hydrocarbon-degrading potential of epiphytic microbial communities on Platanus x acerifolia leaves in an urban area. 2017 , 220, 650-658	27
856	Seasonality of freshwater bacterioplankton diversity in two tropical shallow lakes from the Brazilian Atlantic Forest. 2017 , 93,	22
855	Are There Any Other Compounds Isolated From Dermacoccus spp at All?. 2017 , 74, 132-144	8
854	Phage-Enabled Nanomedicine: From Probes to Therapeutics in Precision Medicine. 2017 , 56, 1964-1992	98
853	Key role of LaeA and velvet complex proteins on expression of Elactam and PR-toxin genes in Penicillium chrysogenum: cross-talk regulation of secondary metabolite pathways. 2017 , 44, 525-535	25
852	Combination of Fenton processes and biotreatment for wastewater treatment and soil remediation. 2017 , 574, 1599-1610	236
851	Nanomedizin auf Phagenbasis: von Sonden zu Therapeutika f∃eine Pr⊠isionsmedizin. 2017 , 129, 1992-2022	7
850	Isolation and characterization of roridin E. 2017 , 55, 337-340	2
8 ₅ 0	Isolation and characterization of roridin E. 2017 , 55, 337-340 Biodesulfurization: How to Make it Work?. 2017 , 42, 1-9	38
849	Biodesulfurization: How to Make it Work?. 2017 , 42, 1-9 Paper-based microfluidic sampling and separation of analytes for potentiometric ion sensing. 2017 ,	38
849 848	Biodesulfurization: How to Make it Work?. 2017 , 42, 1-9 Paper-based microfluidic sampling and separation of analytes for potentiometric ion sensing. 2017 , 243, 346-352	38
849 848 847	Biodesulfurization: How to Make it Work?. 2017, 42, 1-9 Paper-based microfluidic sampling and separation of analytes for potentiometric ion sensing. 2017, 243, 346-352 Aquatic viral metagenomics: Lights and shadows. 2017, 239, 87-96 Ebola virus: A gap in drug design and discovery - experimental and computational perspective. 2017	38 29 11
849 848 847 846	Biodesulfurization: How to Make it Work?. 2017, 42, 1-9 Paper-based microfluidic sampling and separation of analytes for potentiometric ion sensing. 2017, 243, 346-352 Aquatic viral metagenomics: Lights and shadows. 2017, 239, 87-96 Ebola virus: A gap in drug design and discovery - experimental and computational perspective. 2017, 89, 297-308 A metagenome-derived thermostable Eglucanase with an unusual module architecture which	38 29 11
849 848 847 846	Biodesulfurization: How to Make it Work?. 2017, 42, 1-9 Paper-based microfluidic sampling and separation of analytes for potentiometric ion sensing. 2017, 243, 346-352 Aquatic viral metagenomics: Lights and shadows. 2017, 239, 87-96 Ebola virus: A gap in drug design and discovery - experimental and computational perspective. 2017, 89, 297-308 A metagenome-derived thermostable Eglucanase with an unusual module architecture which defines the new glycoside hydrolase family GH148. 2017, 7, 17306 The sensitivity of yeasts and yeasts-like fungi to copper and sulfur could explain lower yeast	38 29 11 12

842	Searching for Antibiotic Resistance Genes in a Pristine Arctic Wetland. 2017 , 160, 42-59	14
841	Sulfated Alginates as Heparin Analogues: A Review of Chemical and Functional Properties. 2017 , 22,	28
840	The Application of Whole Cell-Based Biosensors for Use in Environmental Analysis and in Medical Diagnostics. 2017 , 17,	156
839	Structural insights into RapZ-mediated regulation of bacterial amino-sugar metabolism. 2017 , 45, 10845-1086	024
838	Pherotype Polymorphism in Streptococcus pneumoniae Has No Obvious Effects on Population Structure and Recombination. 2017 , 9, 2546-2559	8
837	Characterization of bacterial community associated with phytoplankton bloom in a eutrophic lake in South Norway using 16S rRNA gene amplicon sequence analysis. 2017 , 12, e0173408	46
836	Characterization of spatial distribution of the bacterial community in the South Sea of Korea. 2017 , 12, e0174159	18
835	Denitrification potential of the eastern oyster microbiome using a 16S rRNA gene based metabolic inference approach. 2017 , 12, e0185071	45
834	Dynamic cellular complexity of anoxygenic phototrophic sulfur bacteria in the chemocline of meromictic Lake Cadagno. 2017 , 12, e0189510	13
833	Curcuminoid Analogs via Microbial Biotransformation With Improved Therapeutic Properties. 2017 , 251-275	1
832	CbtA toxin of Escherichia coli inhibits cell division and cell elongation via direct and independent interactions with FtsZ and MreB. 2017 , 13, e1007007	23
831	Conversion of an inactive xylose isomerase into a functional enzyme by co-expression of GroEL-GroES chaperonins in Saccharomyces cerevisiae. 2017 , 17, 71	15
830	Antibiotic resistance genes in an urban river as impacted by bacterial community and physicochemical parameters. 2017 , 24, 23753-23762	83
829	Diversity of the Potential 2-Methylisoborneol-Producing Genotypes in Thai Strains of Planktothricoides (Cyanobacteria). 2017 , 60,	3
828	Functional Genomics of Pathogenesis. 2017 , 123-142	
827	Trypanocidal Effects of Essential Oils from Selected Medicinal Plants. Synergy among the Main Components. 2017 , 12, 1934578X1701200	5
826	Biochemical process of ureolysis-based microbial CaCO precipitation and its application in self-healing concrete. 2018 , 102, 3121-3132	36
825	References. 2018 , 313-359	_

824	Engineering chemical interactions in microbial communities. 2018 , 47, 1705-1729	19
823	Efficacy of a bacterial preparation of Aneurinibacillus migulanus against downy mildew of cucumber (Pseudoperonospora cubensis). 2018 , 151, 439-450	4
822	Does fungal endophyte inoculation affect the responses of aspen seedlings to carbon dioxide enrichment?. 2018 , 33, 24-31	5
821	BTK drives ibrutinib resistance via ERK1/2 and protects BTK MYD88-mutated cells by a paracrine mechanism. 2018 , 131, 2047-2059	48
820	Anti-elastase and anti-collagenase potential of Lactobacilli exopolysaccharides on human fibroblast. 2018 , 46, 1051-1061	15
819	Shrub range expansion alters diversity and distribution of soil fungal communities across an alpine elevation gradient. 2018 , 27, 2461-2476	25
818	A novel anode modified by 1,5-dihydroxyanthraquinone/multiwalled carbon nanotubes composite in marine sediment microbial fuel cell and its electrochemical performance. 2018 , 42, 2574-2582	5
817	Recent Advances in Intrinsic Self-Healing Cementitious Materials. 2018, 30, e1705679	108
816	In vitro photodynamic inactivation effects of benzylidene cyclopentanone photosensitizers on clinical fluconazole-resistant Candida albicans. 2018 , 22, 178-186	6
815	Composition and functional diversity of microbial community across a mangrove-inhabited mudflat as revealed by 16S rDNA gene sequences. 2018 , 633, 518-528	62
814	Fungal Morphogenesis, from the Polarized Growth of Hyphae to Complex Reproduction and Infection Structures. 2018 , 82,	141
813	Ready-to-eat meat products as a source of Listeria monocytogenes. 2018 , 62, 49-55	42
812	Dual-species biofilm of Listeria monocytogenes and Escherichia coli on stainless steel surface. 2018 , 34, 61	11
811	Development and characterization of the gummy supplements, enriched with probiotics and prebiotics. 2018 , 16, 580-587	10
810	Recent advances in reconstructing microbial secondary metabolites biosynthesis in Aspergillus spp. 2018 , 36, 739-783	46
809	Methane oxidation in industrial biogas plants-Insights in a novel methanotrophic environment evidenced by pmoA gene analyses and stable isotope labelling studies. 2018 , 270, 77-84	2
808	Smart Fertilizers as a Strategy for Sustainable Agriculture. 2018 , 147, 119-157	87
807	Fruiting-Body Development in Ascomycetes. 2018 , 1-56	17

806	The nasal microbiome mirrors and potentially shapes olfactory function. 2018 , 8, 1296	44
805	Impact of microbiota on the use and effects of isoflavones in the relief of climacteric symptoms in menopausal women 🖪 review. 2018 , 41, 100-111	12
804	Biogeographical patterns of bacterial and archaeal communities from distant hypersaline environments. 2018 , 41, 139-150	24
803	Simulated climate change affects how biocrusts modulate water gains and desiccation dynamics after rainfall events. 2018 , 11, e1935	25
802	Systems Biology Modeling to Study Pathogen-Host Interactions. 2018 , 1734, 97-112	4
801	N-Acyl-homoserine lactones and autoinducer-2-mediated quorum sensing during wastewater treatment. 2018 , 102, 1119-1130	25
800	Unraveling the lipid and pigment biosynthesis in Coelastrella sp. M-60: Genomics-enabled transcript profiling. 2018 , 29, 277-289	13
799	Biodesulfurization of Thiophenic Compounds by a 2-Hydroxybiphenyl-Resistant Gordonia sp. HS126-4N Carrying dszABC Genes. 2018 , 75, 597-603	16
798	Application of Rotating Magnetic Fields Increase the Activity of Antimicrobials Against Wound Biofilm Pathogens. 2018 , 8, 167	10
797	Analysis of the spectrum and antibiotic resistance of uropathogens in outpatients at a tertiary hospital. 2018 , 30, 145-149	19
796	Nutrient dynamics and stream order influence microbial community patterns along a 2914 kilometer transect of the Mississippi River. 2018 , 63, 1837-1855	24
795	The occurrence of antibiotic resistance genes in tap water 🗈 review. 2018 , 30, 01012	1
794	Enzymatic decolourization of water using loofa sponge as cellular carrier: Immobilization and dye degradation performance. 2018 , 96, 2321-2333	5
793	Temporal Variability of Faecal Contamination from On-Site Sanitation Systems in the Groundwater of Northern Thailand. 2018 , 61, 939-953	15
792	Pressure- and heat-induced protein unfolding in bacterial cells: crowding vs. sticking. 2018, 592, 1357-1365	12
791	Incubation with macroalgae induces large shifts in water column microbiota, but minor changes to the epibiota of co-occurring macroalgae. 2018 , 27, 1966-1979	18
790	Rosmarinic acid ameliorates the negative effects of salinity in in vitro-regenerated potato explants (Solanum tuberosum L.). 2018 , 40, 1	4
789	Thermal inactivation kinetics of Paenibacillus sanguinis 2301083PRC and Clostridium sporogenes JCM1416MGA in full and low fat BequeijB cremoso[12018, 84, 395-402	25

(2018-2018)

788	Micronutrient metal speciation is controlled by competitive organic chelation in grassland soils. 2018 , 120, 283-291	19
787	Stress Responses of Bacterial Cells as Mechanism of Development of Antibiotic Tolerance (Review). 2018 , 54, 108-127	20
786	Bio-active nanocomposite films based on nanocrystalline cellulose reinforced styrylquinoxalin-grafted-chitosan: Antibacterial and mechanical properties. 2018 , 114, 733-740	28
7 ⁸ 5	Dynamics of the human gut phageome during antibiotic treatment. 2018 , 74, 420-427	10
784	In Vitro Antimicrobial Assessment of Selected Plant Extracts from Pakistan. 2018 , 42, 267-272	3
783	Antioxidant and antibacterial activity of different extracts from herbs obtained by maceration or supercritical technology. 2018 , 133, 58-64	46
782	Potential for gulls to transport bacteria from human waste sites to beaches. 2018 , 615, 123-130	31
781	Occurrence, populations, diversity, and growth potential of spore-forming bacteria in E equeij ö cremoso[] 2018 , 89, 24-31	11
780	Saltatory Rolling Circle Amplification (SRCA): a Novel Nucleic Acid Isothermal Amplification Technique Applied for Rapid Detection of Shigella Spp. in Vegetable Salad. 2018 , 11, 504-513	13
779	MybA, a new player driving survival of the conidium of the human pathogen Aspergillus fumigatus. 2018 , 64, 141-146	7
778	The superior effect of nature based solutions in land management for enhancing ecosystem services. 2018 , 610-611, 997-1009	389
777	Biocontrol of Fusarium wilt of Capsicum annuum by rhizospheric bacteria isolated from turmeric endowed with plant growth promotion and disease suppression potential. 2018 , 150, 831-846	15
776	The second International Symposium on Fungal Stress: ISFUS. 2018 , 122, 386-399	8
775	Optimization of rhamnolipid production from PBS towards application for microbial enhanced oil recovery. 2018 , 8, 20	34
774	Scanning electron microscopic observations of early stages of interaction of Trichoderma harzianum, Gliocladium virens and Bacillus subtilis with Acaulospora colombiana. 2018 , 50, 227-229	3
773	Physicochemical and nutraceutical properties of moringa (Moringa oleifera) leaves and their effects in an in vivo AOM/DSS-induced colorectal carcinogenesis model. 2018 , 105, 159-168	47
772	Fluorescent pseudomonads pursue media-dependent strategies to inhibit growth of pathogenic Verticillium fungi. 2018 , 102, 817-831	5

770	Listeria monocytogenes Adaptation and Growth at Low Temperatures. 2018, 227-248	4
7 69	Isolation of folate-producing Weissella spp. from Thai fermented fish (Plaa Som Fug). 2018 , 89, 388-391	6
768	Prescription Opioid Use is Associated with Virologic Failure in People Living with HIV. 2018 , 22, 1323-1328	13
767	Determination of the treatment period of banana seedlings with rhizobacteria in the control of Meloidogyne javanica. 2018 , 40,	
766	MONITORING OF BIOSURFACTANT PRODUCTION BY Bacillus subtilis USING BEET PEEL AS CULTURE MEDIUM VIA THE DEVELOPMENT OF A NEURAL SOFT-SENSOR IN AN ELECTRONIC SPREADSHEET. 2018 , 35, 1355-1367	2
765	Quantifying the Spatiotemporal Dynamics of Plant Root Colonization by Beneficial Bacteria in a Microfluidic Habitat. 2018 , 2, 1800048	11
764	SIMULTANEOUS PRODUCTION OF BIOSURFACTANTS AND LIPASES FROM Aspergillus niger AND OPTIMIZATION BY RESPONSE SURFACE METHODOLOGY AND DESIRABILITY FUNCTIONS. 2018 , 35, 857-868	6
763	Probiotics for the treatment of irritable bowel syndrome in children. 2018,	78
762	Is the undergraduate microbiology curriculum preparing students for careers in their field?: an assessment of biology majors' conceptions of growth and control of microorganisms. 2018 , 5, 42	3
761	Promoting fruit seedling growth by encapsulated microorganisms. 2018, 40,	1
760	Mycorrhizal fungi mediate the direction and strength of plant-soil feedbacks differently between arbuscular mycorrhizal and ectomycorrhizal communities. 2018 , 1, 196	34
759	Delay in the dispersal of flocks moving in unbounded space using long-range interactions. 2018 , 8, 15872	7
758	The transcription factor PRO44 and the histone chaperone ASF1 regulate distinct aspects of multicellular development in the filamentous fungus Sordaria macrospora. 2018 , 19, 112	8
757	Review on biological process of soil improvement in the mitigation of liquefaction in sandy soil. 2018 , 250, 01017	5
756	Structural Insights into the FtsQ/FtsB/FtsL Complex, a Key Component of the Divisome. 2018 , 8, 18061	12
755	Sugarcane apoplast fluid modulates the global transcriptional profile of the diazotrophic bacteria Paraburkholderia tropica strain Ppe8. 2018 , 13, e0207863	8
754	Contemporary IncI1 plasmids involved in the transmission and spread of antimicrobial resistance in Enterobacteriaceae. 2021 , 118, 102392	28
753	Diversity and seasonal distribution of ammonia-oxidizing archaea in the water column of a tropical estuary along the southeast Arabian Sea. 2018 , 34, 188	4

(2018-2018)

752	1-Aminocyclopropane-1-carboxylic acid deaminase producing beneficial rhizobacteria ameliorate the biomass characters of Panicum maximum Jacq. by mitigating drought and salt stress. 2018 , 8, 17513	49
75 ¹	Influence of water quality on diversity and composition of fungal communities in a tropical river. 2018 , 8, 14799	8
75°	Application of NGS for studying the antibiotic resistant microbiome in Wroc∃w tap water. 2018 , 44, 00161	2
749	Genomes from uncultivated prokaryotes: a comparison of metagenome-assembled and single-amplified genomes. 2018 , 6, 173	43
748	Production, purification and characterization of an acid/alkali and thermo tolerant cellulase from Schizophyllum commune NAIMCC-F-03379 and its application in hydrolysis of lignocellulosic wastes. 2018 , 8, 173	50
747	Host genetics and the rumen microbiome jointly associate with methane emissions in dairy cows. 2018 , 14, e1007580	89
746	Engineering of the Filamentous Fungus as Cell Factory for Natural Products. <i>Frontiers in Microbiology</i> , 2018 , 9, 2768	35
745	Fungal persister cells: The basis for recalcitrant infections?. 2018 , 14, e1007301	52
744	Association between functional lactase variants and a high abundance of Bifidobacterium in the gut of healthy Japanese people. 2018 , 13, e0206189	21
743	Major 3'-5' Exoribonucleases in the Metabolism of Coding and Non-coding RNA. 2018 , 159, 101-155	12
742	Natural Food Antimicrobials: Recent Trends in Their Use, Limitations, and Opportunities for Their Applications in Food Preservation. 2018 , 25-43	1
741	Off-label use of ceftiofur in one-day chicks triggers a short-term increase of ESBL-producing E. coli in the gut. 2018 , 13, e0203158	12
740	Importance of the agar-media in the evaluation of bacteriocin activity against the same test-microorganisms. 2018 , 54,	4
739	Sporulation environment influences spore properties in Bacillus: evidence and insights on underlying molecular and physiological mechanisms. 2018 , 42, 614-626	46
738	Potential Treatment Options in a Post-antibiotic Era. 2018 , 1052, 51-61	28
737	Effectiveness of polymeric coated films containing bacteriocin-producer living bacteria for Listeria monocytogenes control under simulated cold chain break. 2018 , 76, 173-179	12
736	Investigations on VELVET regulatory mutants confirm the role of host tissue acidification and secretion of proteins in the pathogenesis of Botrytis cinerea. 2018 , 219, 1062-1074	33
735	Implication of orphan histidine kinase (OhkAsp) in biosynthesis of doxorubicin and daunorubicin in Streptomyces peucetius ATCC 27952. 2018 , 214, 37-46	4

734	Drought-Tolerant Pseudomonas spp. Improve the Growth Performance of Finger Millet (Eleusine coracana (L.) Gaertn.) Under Non-Stressed and Drought-Stressed Conditions. 2018 , 28, 227-240	54
733	Impact of Pseudomonas graminis strain CPA-7 on respiration and ethylene production in fresh-cut Golden delicious apple according to the maturity stage and the preservation strategy. 2018 , 144, 36-45	1
732	Zinc is an inhibitor of the LdtR transcriptional activator. 2018 , 13, e0195746	1
731	Antifungal photodynamic inactivation against dermatophyte Trichophyton rubrum using nanoparticle-based hybrid photosensitizers. 2018 , 23, 202-208	9
730	Nisin and other antimicrobial peptides: Production, mechanisms of action, and application in active food packaging. 2018 , 48, 179-194	97
729	Impact of Unit Operations From Farm to Fork on Microbial Safety and Quality of Foods. 2018 , 85, 131-175	2
728	Microbial Aspects of Acid Mine DrainageMining Environmental Pollution and Control. 2018, 395-432	
727	Rhizosphere mediated nutrient management in Allium hookeri Thwaites by using phosphate solubilizing rhizobacteria and tricalcium phosphate amended soil. 2018 , 13, 256-269	5
726	Microbial anaerobic Fe(II) oxidation - Ecology, mechanisms and environmental implications. 2018 , 20, 3462-3483	79
725	Presence of fluoroquinolone resistance with persistent occurrence of gyrA gene mutations in a municipal wastewater treatment plant in India. 2018 , 211, 817-825	14
724	Studies on the potential application of various blends of essential oils as antioxidant and antimicrobial preservatives in emulsion based chicken sausages. 2018 , 120, 1398-1411	8
723	Whole genome sequences to assess the link between antibiotic and metal resistance in three coastal marine bacteria isolated from the mummichog gastrointestinal tract. 2018 , 135, 514-520	7
722	Velvet domain protein VosA represses the zinc cluster transcription factor SclB regulatory network for Aspergillus nidulans asexual development, oxidative stress response and secondary metabolism. 2018 , 14, e1007511	13
721	Development and characterisation of self-assembled graphene hydrogel-based anodes for bioelectrochemical systems 2018 , 8, 26755-26763	5
720	Diversity and functionality of archaeal, bacterial and fungal communities in deep Archaean bedrock groundwater. 2018 , 94,	18
719	Feeding sows resistant starch during gestation and lactation impacts their faecal microbiota and milk composition but shows limited effects on their progeny. 2018 , 13, e0199568	15
718	The application of growth-no growth models to directly assess the stability of wholemeal multigrain bread towards Penicillium paneum LMQA-002 and Paecilomyces variotii LMQA-001. 2018 , 97, 231-237	7
717	Description of New and Amended Clades of the Genus Photobacterium. 2018 , 6,	6

(2018-2018)

716	Evolutionary insights into Trm112-methyltransferase holoenzymes involved in translation between archaea and eukaryotes. 2018 , 46, 8483-8499	25
715	Quorum sensing intervened bacterial signaling: Pursuit of its cognizance and repression. 2018 , 16, 239-252	27
714	Harnessing fungi to mitigate CH in natural and engineered systems. 2018, 102, 7365-7375	4
713	Biofilm formation and electron transfer in bioelectrochemical systems. 2018 , 7, 220-234	18
712	Production, characterization and environmental applications of biosurfactants from Bacillus amyloliquefaciens and Bacillus subtilis. 2018 , 16, 132-139	28
711	Effect of antiprotozoal molecules on hypnospores of Perkinsus spp. parasite. 2018 , 192, 25-35	2
710	Characterisation of an ABC transporter of a resistant Candida glabrata clinical isolate. 2018 , 113, e170484	4
709	Ubiquitous giants: a plethora of giant viruses found in Brazil and Antarctica. 2018 , 15, 22	23
708	Antimicrobial activity of extracts and phthalides occurring in Apiaceae plants. 2018 , 32, 1459-1487	9
707	Role of acetyl-CoA Synthetase and LovE Regulator Protein of Polyketide Biosynthesis in Lovastatin Production by Wild-Type and Overproducing Aspergillus terreus Strains. 2018 , 54, 188-197	7
706	Soil Microorganisms. 2018 , 457-482	2
705	Characterization of extended-spectrum and CMY-2 Elactamases, and associated virulence genes in Escherichia coli from food of animal origin in Maico. 2018 , 120, 1457-1473	1
704	ConnectedAlign: a PPI network alignment method for identifying conserved protein complexes across multiple species. 2018 , 19, 286	3
703	The Type III Secretion System of Pathogenic Escherichia coli. 2018 , 416, 51-72	20
702	Evaluating the impact of sewage discharges on the marine environment with a lightweight AUV. 2018 , 135, 714-722	8
701	Antimicrobial potentiality of actinobacteria isolated from two microbiologically unexplored forest ecosystems of Northeast India. 2018 , 18, 71	22
700	Semi-continuous treatment of naphthenic acids using aerobic granular sludge. 2018 , 3, 191-199	8
699	Self-sustaining closed-loop multienzyme-mediated conversion of amines into alcohols in continuous reactions. 2018 , 1, 452-459	76

698	The effect of calcium on biofilm formation in dairy wastewater. 2018, 13, 400-409	7
697	Biofilm monitoring as a tool to assess the efficiency of artificial reefs as substrates: Toward 3D printed reefs. 2018 , 120, 230-237	8
696	Photochemistry beyond the red limit in chlorophyll f-containing photosystems. 2018, 360, 1210-1213	129
695	Comprehensive analysis of Verticillium nonalfalfae in silico secretome uncovers putative effector proteins expressed during hop invasion. 2018 , 13, e0198971	12
694	Fungal genotype determines survival of Drosophila melanogaster when competing with Aspergillus nidulans. 2018 , 13, e0190543	3
693	Ex vivo expansion of alveolar macrophages with Mycobacterium tuberculosis from the resected lungs of patients with pulmonary tuberculosis. 2018 , 13, e0191918	8
692	Genomic insights into the broad antifungal activity, plant-probiotic properties, and their regulation, in Pseudomonas donghuensis strain SVBP6. 2018 , 13, e0194088	18
691	Host: Microbiome co-metabolic processing of dietary polyphenols - An acute, single blinded, cross-over study with different doses of apple polyphenols in healthy subjects. 2018 , 112, 108-128	48
690	Metabolic and proteomic mechanism of bisphenol A degradation by Bacillus thuringiensis. 2018 , 640-641, 714-725	13
689	Validation of Self-Healing Properties of Construction Materials through Nondestructive and Minimal Invasive Testing. 2018 , 5, 1800179	14
688	Optimization and purification of glucansucrase produced by Leuconostoc mesenteroides DRP2-19 isolated from Chinese Sauerkraut. 2018 , 48, 465-473	5
687	Actinobacteria. 2018 , 191-205	4
686	Potential of Brevibacillus sp. AVN 13 isolated from crude oil contaminated soil for biosurfactant production and its optimization studies. 2018 , 6, 4347-4356	19
685	How leaking and overproducing resources affect the evolutionary robustness of cooperative cross-feeding. 2018 , 454, 278-291	3
684	Low-dose addition of silver nanoparticles stresses marine plankton communities. 2018 , 5, 1965-1980	10
683	Essential oils from Lippia turbinata and Tagetes minuta persistently reduce in vitro ruminal methane production in a continuous-culture system. 2019 , 59, 709	9
682	Glucose electro-fermentation as main driver for efficient H2-producing bacteria selection in mixed cultures. 2019 , 44, 2230-2238	15
681	Optimization of biosurfactant production by N3-1P using the brewery waste as the carbon source. 2019 , 40, 3371-3380	28

680	Biofertilizers and Their Role in Sustainable Agriculture. 2019 , 285-300	5
679	Plant-Microbiome Interactions in Agroecosystem: An Application. 2019 , 251-291	3
678	Rhizobial Inoculants for Sustainable Agriculture: Prospects and Applications. 2019, 245-283	5
677	A first insight into the structure and function of rhizosphere microbiota in Antarctic plants using shotgun metagenomic. 2019 , 42, 1825-1835	8
676	Evaluation of a new strategy in the elaboration of culture media to produce surfactin from hemicellulosic corncob liquor. 2019 , 24, e00364	10
675	Enhancement of Etarotene production by regulating the autophagy-carotenoid biosynthesis seesaw in Chlamydomonas reinhardtii. 2019 , 292, 121937	15
674	Mycorrhizosphere: Microbial Interactions for Sustainable Agricultural Production. 2019, 321-338	7
673	Mammalian cell entry operons; novel and major subset candidates for diagnostics with special reference to subspecies infection. 2019 , 39, 65-75	8
672	Release behavior of 1-methylcylopropene coated paper-based shellac solution in response to stepwise humidity changes to develop novel functional packaging for fruit. 2019 , 32, 523-533	11
671	Detection of pathogenic bacteria in hot tap water using the qPCR method: preliminary research. 2019 , 1, 1	8
670	Anthropogenic impacts on sulfonamide residues and sulfonamide resistant bacteria and genes in Larut and Sangga Besar River, Perak. 2019 , 688, 1335-1347	13
669	High-value chemicals from Botryococcus braunii and their current applications - A review. 2019 , 291, 121911	12
668	1,3-Diaminopropane and Spermidine Upregulate Lovastatin Production and Expression of Lovastatin Biosynthetic Genes in Aspergillus terreus via LaeA Regulation. 2019 , 55, 243-254	10
667	COP9 Signalosome Interaction with UspA/Usp15 Deubiquitinase Controls VeA-Mediated Fungal Multicellular Development. 2019 , 9,	7
666	Integration of Sensor Cells into Hardware Platforms. 2019 , 1-23	
665	Algae colonisation of brick pavement at the University of Venda: A potential slippery hazard. 2019 , 11, 689	1
664	High-throughput sequencing revealed differences of microbial community structure and diversity between healthy and diseased Caulerpa lentillifera. 2019 , 19, 225	7
663	Antimicrobial and antioxidant properties of tomato processing byproducts and their correlation with the biochemical composition. 2019 , 116, 108558	36

662	An expansin-like protein expands forage cell walls and synergistically increases hydrolysis, digestibility and fermentation of livestock feeds by fibrolytic enzymes. 2019 , 14, e0224381	4
661	Latitudinal gradient of cyanobacterial diversity in tidal flats. 2019 , 14, e0224444	1
660	An Overview of Biotechnological Processes in the Food Industry. 2019 , 1-19	2
659	Metagenomic study of endophytic bacterial community of sweet potato (Ipomoea batatas) cultivated in different soil and climatic conditions. 2019 , 35, 176	8
658	Lactobacillus rhamnosus Lcr35 as an effective treatment for preventing Candida albicans infection in the invertebrate model Caenorhabditis elegans: First mechanistic insights. 2019 , 14, e0216184	14
657	Biomonitoring, physico-chemical, and biomarker evaluations of abattoir effluent discharges into the Ogun River from Kara Market, Ogun State, Nigeria, using Clarias gariepinus. 2018 , 191, 44	10
656	Light in the Fungal World: From Photoreception to Gene Transcription and Beyond. 2019 , 53, 149-170	44
655	Genomic signatures and co-occurrence patterns of the ultra-small Saccharimonadia (phylum CPR/Patescibacteria) suggest a symbiotic lifestyle. 2019 , 28, 4259-4271	33
654	Evaluation of surfactin synthesis in a genome reduced Bacillus subtilis strain. 2019, 9, 84	15
653	Community structure and functional genes in radionuclide contaminated soils in Chernobyl and Fukushima. 2019 , 366,	11
652	Intraspecific variation on epiphytic bacterial community from Laguncularia racemosa phylloplane. 2019 , 50, 1041-1050	O
651	Microbial diversity in grape musts from Austrian and South African grape varieties and regions. 2019 , 12, 02028	1
650	Protocol adjustment improves the extraction of high-quality total RNA from common bean stems infected by Sclerotinia sclerotiorum. 2019 , 43,	2
649	Capability for arsenic mobilization in groundwater is distributed across broad phylogenetic lineages. 2019 , 14, e0221694	6
648	Screening chemical modulators of benzoic acid derivatives to improve lipid accumulation in SR21 with metabolomics analysis. 2019 , 12, 209	13
647	The bacterial community and fermentation quality of mulberry (Morus alba) leaf silage with or without Lactobacillus casei and sucrose. 2019 , 293, 122059	41
646	Alginates along the filament of the brown alga Ectocarpus help cells cope with stress. 2019 , 9, 12956	14
645	A newly constructed Agrobacterium-mediated transformation system revealed the influence of nitrogen sources on the function of the LaeA regulator in Penicillium chrysogenum. 2019 , 123, 830-842	3

644	The Effect of Blue Light on the Production of Citrinin in M9 by Regulating the Gene through lncRNA. 2019 , 11,	8
643	Identification and Characterization of Two New S-Adenosylmethionine-Dependent Methyltransferase Encoding Genes Suggested Their Involvement in Stipe Elongation of. 2019 , 47, 441-448	2
642	An electrochemical investigation of interfacial electron uptake by the sulfur oxidizing bacterium Thioclava electrotropha ElOx9. 2019 , 324, 134838	12
641	Nonreductive biomineralization of uranium by Bacillus subtilis ATCC-6633 under aerobic conditions. 2019 , 208-209, 106027	13
640	Dissection of patulin biosynthesis, spatial control and regulation mechanism in Penicillium expansum. 2019 , 21, 1124-1139	40
639	Nanobiosensors: Point-of-care approaches for cancer diagnostics. 2019 , 130, 147-165	63
638	Sustainable Nanocomposites in Food Packaging. 2019 , 413-436	O
637	Suppression of arbuscular mycorrhizal fungal activity in a diverse collection of non-cultivated soils. 2019 , 95,	13
636	Identification of Escherichia coli from broiler chickens in Jordan, their antimicrobial resistance, gene characterization and the associated risk factors. 2019 , 15, 159	45
635	New glycopolymers containing both D- and L-rhamnopyranoses from Rathayibacter iranicus VKM Ac-1602 cell wall. 2019 , 482, 107728	6
634	A neural network model predicts community-level signaling states in a diverse microbial community. 2019 , 15, e1007166	2
633	Secondary Metabolites: Metabolomics for Secondary Metabolites. 2019 , 333-344	2
632	Advances in whole cell-based biosensors in environmental monitoring. 2019 , 263-284	7
631	Anaerobic hydrocarbon degradation in candidate phylum 'Atribacteria' (JS1) inferred from genomics. 2019 , 13, 2377-2390	29
630	Impact of the self-healing agent composition on material characteristics of bio-based self-healing concrete. 2019 , 11, e00250	12
629	Genome-facilitated discovery of RND efflux pump-mediated resistance to cephalosporins in Vibrio spp. isolated from the mummichog fish gut. 2019 , 19, 294-300	4
628	Cadmium stress effects indicating marine pollution in different species of sea urchin employed as environmental bioindicators. 2019 , 24, 675-687	18
627	Soil quality indicators under management systems in a Quilombola community in the Brazilian Cerrado. 2019 , 76, 518-526	2

626	Mediterranean grassland soil C-N compound turnover is dependent on rainfall and depth, and is mediated by genomically divergent microorganisms. 2019 , 4, 1356-1367	70
625	Substrate and spatial variables are major determinants of fungal community in karst caves in Southwest China. 2019 , 46, 1504	7
624	Control of Development, Secondary Metabolism and Light-Dependent Carotenoid Biosynthesis by the Velvet Complex of. 2019 , 212, 691-710	9
623	Genetic Engineering Applications to Improve Cellulase Production and Efficiency: Part II. 2019 , 227-260	1
622	Cultivable hydrocarbon degrading bacteria have low phylogenetic diversity but highly versatile functional potential. 2019 , 142, 43-51	7
621	Environmental factors driving the dominance of the harmful bloom-forming cyanobacteria Microcystis and Aphanocapsa in a tropical water supply reservoir. 2019 , 91, 1466-1478	11
620	Novel Stenotrophomonas maltophilia temperate phage DLP4 is capable of lysogenic conversion. 2019 , 20, 300	16
619	Medical prescribing and antibiotic resistance: A game-theoretic analysis of a potentially catastrophic social dilemma. 2019 , 14, e0215480	14
618	The spleen morphophysiology of fruit bats. 2019 , 48, 315-324	5
617	Decoding the Fatty Acid Profile of Bacillus licheniformis I89 and Its Adaptation to Different Growth Conditions to Investigate Possible Biotechnological Applications. 2019 , 54, 245-253	2
616	Effects of elemental magnesium and magnesium oxide on hydrogen, methane and volatile fatty acids production in in vitro rumen batch cultures. 2019 , 252, 74-82	3
615	A review on the biosynthesis of metal and metal salt nanoparticles by microbes 2019 , 9, 12944-12967	276
614	Effect of linoleic acid and methyl jasmonate on astaxanthin content of Scenedesmus acutus and Chlorella sorokiniana under heterotrophic cultivation and salt shock conditions. 2019 , 31, 2811-2822	10
613	Assembly of a heptameric STRIPAK complex is required for coordination of light-dependent multicellular fungal development with secondary metabolism in Aspergillus nidulans. 2019 , 15, e1008053	16
612	Effect of different pretreatment on the microbial diversity of fermented potato revealed by high-throughput sequencing. 2019 , 290, 125-134	16
611	Anti-E. coli cellulose-based materials. 2019 , 107, 325-330	3
610	The genus Aeromonas: A general approach. 2019 , 130, 81-94	54
609	Diversity and functional analysis of salivary microflora of Indian Antarctic expeditionaries. 2019 , 11, 1581513	7

608	Gene regulation of Sclerotinia sclerotiorum during infection of Glycine max: on the road to pathogenesis. 2019 , 20, 157	19
607	Relative stability of the Pocillopora acuta microbiome throughout a thermal stress event. 2019 , 38, 373-386	20
606	Carbon Cycle Implications of Soil Microbial Interactions. 2019 , 1-29	
605	Molecular detection of Shiga toxin-producing and antibiotic-resistant Escherichia coli isolates from buffaloes in southwest of Iran. 2019 , 51, 1725-1736	3
604	The Effect of Land-Use Change on Soil CH4 and N2O Fluxes: A Global Meta-Analysis. 2019, 22, 1424-1443	25
603	Evolution of hierarchy in bacterial metabolic networks. 2019 , 180, 71-78	2
602	Retail liver juices enhance the survivability of Campylobacter jejuni and Campylobacter coli at low temperatures. 2019 , 9, 2733	14
601	Biodeterioration of mortar coating in historical buildings: Microclimatic characterization, material, and fungal community. 2019 , 155, 195-209	17
600	Polyextremophilic Bacteria from High Altitude Andean Lakes: Arsenic Resistance Profiles and Biofilm Production. 2019 , 2019, 1231975	10
599	Maize synthesized benzoxazinoids affect the host associated microbiome. 2019 , 7, 59	87
598	Epicoccum layuense a potential biological control agent of esca-associated fungi in grapevine. 2019 , 14, e0213273	22
597	Evaluation of MALDI-TOF MS and an expanded custom reference spectra database for the identification and differentiation of Taylorella equigenitalis and Taylorella asinigenitalis. 2019 , 94, 326-330	1
596	Mapping the Fungal Battlefield: Using Chemistry and Deletion Mutants to Monitor Interspecific Chemical Interactions Between Fungi. <i>Frontiers in Microbiology</i> , 2019 , 10, 285	22
595	Toll-like receptor 3 regulates Zika virus infection and associated host inflammatory response in primary human astrocytes. 2019 , 14, e0208543	30
594	High hydrostatic pressure processing of sliced fermented sausages: A quantitative exposure assessment for Listeria monocytogenes. 2019 , 52, 406-419	15
593	Linking BmicsIto function unlocks the biotech potential of non-model fungi. 2019 , 14, 9-17	11
592	Chemical Composition, Antifungal, and Cytotoxicity Activities of (Sw.) Willd Leaves. 2019 , 2019, 9423658	5
591	Moderate plant water stress improves larval development, and impacts immunity and gut microbiota of a specialist herbivore. 2019 , 14, e0204292	8

590	Physiological and proteomic analysis of grown under the iron-limited condition. 2019 , 14, 1417-1428	1
589	Acute lung injury and repair induced by single exposure of in immunocompetent mice. 2019 , 14, 1511-1525	3
588	Vitamin E for Prevention of Biofilm-caused Healthcare-associated Infections. 2018, 15, 14-21	11
587	The Spatial Distribution of the Microbial Community in a Contaminated Aquitard below an Industrial Zone. 2019 , 11, 2128	9
586	Exploring a glycosylation methodology for the synthesis of hydroxamate-modified alginate building blocks. 2019 , 17, 9321-9335	6
585	Advances in Fe(III) bioreduction and its application prospect for groundwater remediation: A review. 2019 , 13, 1	14
584	Chlorpromazine-impregnated catheters as a potential strategy to control biofilm-associated urinary tract infections. 2019 , 14, 1023-1034	6
583	Extended spectrum flactamase producing uropathogenic Escherichia coli and the correlation of biofilm with antibiotics resistance in Nepal. 2019 , 18, 42	15
582	Changes of the denitrifying communities in a multi-stage free water surface constructed wetland. 2019 , 650, 1419-1425	24
581	Genomic and Functional Characterization of the Endophytic Bacillus subtilis 7PJ-16 Strain, a Potential Biocontrol Agent of Mulberry Fruit Sclerotiniose. 2019 , 77, 651-663	18
580	Molecular Characterization of Fusarium spp. Associated with Sugarcane Pokkah Boeng from the Philippines Using Partial Translation Elongation Factor-1∰TEF-1∄Gene Sequences. 2019 , 21, 619-630	2
579	Conversion of renewable substrates for biosurfactant production by Rhizopus arrhizus UCP 1607 and enhancing the removal of diesel oil from marine soil. 2019 , 38, 40-48	24
578	Effect of tree foliage supplementation of tropical grass diet on in vitro digestibility and fermentation, microbial biomass synthesis and enteric methane production in ruminants. 2019 , 51, 893-904	11
577	Evaluation of the microbiological contamination of food processing environments through implementing surface sensors in an iberian pork processing plant: An approach towards the control of Listeria monocytogenes. 2019 , 99, 40-47	21
576	Biosurfactants and Bioemulsifiers from Marine Algae. 2019 , 169-188	2
575	Physicochemical and microbiological monitoring of anaerobic digestion at laboratory and low cost tubular digester. 2019 , 274, 321-328	O
574	Secondary metabolite profiles of small-spored Alternaria support the new phylogenetic organization of the genus. 2019 , 291, 135-143	18
573	Molecularly imprinted nanoparticles based plasmonic sensors for real-time Enterococcus faecalis detection. 2019 , 126, 608-614	56

(2020-2019)

572	Mass Spectrometry-Based Microbial Metabolomics: Techniques, Analysis, and Applications. 2019 , 1859, 11-69	6
571	Setting-up a fast and reliable cytokinin biosensor based on a plant histidine kinase receptor expressed in Saccharomyces cerevisiae. 2019 , 289, 103-111	6
570	Self-assembly of supramolecular structure based on copper-lipopeptides isolated from e-waste bioleaching liquor. 2019 , 368, 63-71	4
569	Newly isolated microorganisms with potential application in biotechnology. 2019 , 37, 319-339	36
568	Microbial Role in the Ecology of Antarctic Plants. 2019 , 257-275	0
567	The microbiomic and environmental analysis of sediments in the Indo-Pacific humpback dolphin (Sousa chinensis) habitat in the Northern Beibu Gulf, China. 2019 , 26, 6957-6970	5
566	Sugar and iron: Toward understanding the antibacterial effect of ciclopirox in Escherichia coli. 2019 , 14, e0210547	5
565	Exploring the Molecular Mechanisms of Extracellular Electron Transfer for Harnessing Reducing Power in METs: Methodologies and Approaches. 2019 , 261-293	O
564	Oxygen loss from seagrass roots coincides with colonisation of sulphide-oxidising cable bacteria and reduces sulphide stress. 2019 , 13, 707-719	48
563	Polyketide and Its Derivatives. 2019 , 219-227	
563 562	Polyketide and Its Derivatives. 2019, 219-227 Molecular modelling and competitive inhibition of a Mycobacterium tuberculosis multidrug-resistance efflux pump. 2019, 87, 98-108	7
	Molecular modelling and competitive inhibition of a Mycobacterium tuberculosis	7 57
562	Molecular modelling and competitive inhibition of a Mycobacterium tuberculosis multidrug-resistance efflux pump. 2019 , 87, 98-108 Discovery of several novel, widespread, and ecologically distinct marine Thaumarchaeota viruses	
562	Molecular modelling and competitive inhibition of a Mycobacterium tuberculosis multidrug-resistance efflux pump. 2019, 87, 98-108 Discovery of several novel, widespread, and ecologically distinct marine Thaumarchaeota viruses that encode amoC nitrification genes. 2019, 13, 618-631	57
562 561 560	Molecular modelling and competitive inhibition of a Mycobacterium tuberculosis multidrug-resistance efflux pump. 2019, 87, 98-108 Discovery of several novel, widespread, and ecologically distinct marine Thaumarchaeota viruses that encode amoC nitrification genes. 2019, 13, 618-631 Opening the Black Box of Thermophilic Autotrophic Bacterial Diversity. 2019, 333-343	57 1
562 561 560	Molecular modelling and competitive inhibition of a Mycobacterium tuberculosis multidrug-resistance efflux pump. 2019, 87, 98-108 Discovery of several novel, widespread, and ecologically distinct marine Thaumarchaeota viruses that encode amoC nitrification genes. 2019, 13, 618-631 Opening the Black Box of Thermophilic Autotrophic Bacterial Diversity. 2019, 333-343 Structural and Functional Diversity of Microbial Metallothionein Genes. 2019, 387-407 Exploring Microbial Diversity and Function in Petroleum Hydrocarbon Associated Environments	57 1 8
562 561 560 559 558	Molecular modelling and competitive inhibition of a Mycobacterium tuberculosis multidrug-resistance efflux pump. 2019, 87, 98-108 Discovery of several novel, widespread, and ecologically distinct marine Thaumarchaeota viruses that encode amoC nitrification genes. 2019, 13, 618-631 Opening the Black Box of Thermophilic Autotrophic Bacterial Diversity. 2019, 333-343 Structural and Functional Diversity of Microbial Metallothionein Genes. 2019, 387-407 Exploring Microbial Diversity and Function in Petroleum Hydrocarbon Associated Environments Through Omics Approaches. 2019, 171-194	57 1 8

554	Defining trait-based microbial strategies with consequences for soil carbon cycling under climate change. 2020 , 14, 1-9	169
553	Chemical screening identifies an extract from marine Pseudomonas spPTR-08 as an anti-aging agent that promotes fission yeast longevity by modulating the Pap1-ctt1 pathway and the cell cycle. 2020 , 47, 33-43	8
552	Combination of bioelectrochemical systems and electrochemical capacitors: Principles, analysis and opportunities. 2020 , 39, 107456	34
551	Energy-converting hydrogenases: the link between H metabolism and energy conservation. 2020 , 77, 1461-1481	17
550	Quorum sensing for population-level control of bacteria and potential therapeutic applications. 2020 , 77, 1319-1343	39
549	The effect of an additive containing three Lactobacillus species on the fermentation pattern and microbiological status of silage. 2020 , 100, 1174-1184	12
548	Artisan fresh cheese from raw cow's milk as a possible route of transmission in a toxoplasmosis outbreak, in Brazil. 2020 , 67, 122-129	6
547	Structural equation models to disentangle the biological relationship between microbiota and complex traits: Methane production in dairy cattle as a case of study. 2020 , 137, 36-48	9
546	Land use strongly influences soil organic carbon and bacterial community export in runoff in tropical uplands 2020 , 31, 118-132	10
545	Is there potential in elevational assisted migration for the endangered Magnolia vovidesii?. 2020 , 53, 125782	4
544	Energy conservation involving 2 respiratory circuits. 2020 , 117, 1167-1173	9
543	Probiotic potential and amylolytic properties of lactic acid bacteria isolated from Chinese fermented cereal foods. 2020 , 111, 107057	25
542	Effect of high pressure processing combined with temperature on the inactivation and germination of Alicyclobacillus acidoterrestris spores: Influence of heat-shock on the counting of survivors. 2020 , 118, 108781	7
541	Shifts in metabolic patterns of soil bacterial communities on exposure to metal engineered nanomaterials. 2020 , 189, 110012	14
540	Apc.LaeA and Apc.VeA of the velvet complex govern secondary metabolism and morphological development in the echinocandin-producing fungus Aspergillus pachycristatus. 2020 , 47, 155-168	4
539	Antimicrobial activity of gaseous Citrus limon var pompia leaf essential oil against Listeria monocytogenes on ricotta salata cheese. 2020 , 87, 103386	29
538	Next-Generation Sequencing: An Eye-Opener for the Surveillance of Antiviral Resistance in Influenza. 2020 , 38, 360-367	19
537	Inclusion of a blend of copaiba, cashew nut shell and castor oil in the protein-energy supplement for grazing beef cattle improves rumen fermentation, nutrient intake and fibre digestibility. 2020 , 60, 1039	4

(2020-2020)

536	Effects of environmental factors on denitrifying bacteria and functional genes in sediments of Bohai Sea, China. 2020 , 160, 111621	4
535	Penicillium expansum: biology, omics, and management tools for a global postharvest pathogen causing blue mould of pome fruit. 2020 , 21, 1391-1404	27
534	Effects of aflatoxin contaminated maize on Folsomia candida (Collembola). 2020, 101, 103233	
533	Antibacterial green tea catechins from a molecular perspective: mechanisms of action and structure-activity relationships. 2020 , 11, 9370-9396	18
532	Seawater Pb concentration and isotopic composition response to daily time scale dust storms in the Gulf of Aqaba, Red Sea. 2020 , 227, 103895	3
531	Penicillium oxalicum putative methyltransferase Mtr23B has similarities and differences with LaeA in regulating conidium development and glycoside hydrolase gene expression. 2020 , 143, 103445	2
530	Biopsychism: life between computation and cognition. 2020 , 45, 315-330	1
529	Biofilm-based fermentation: a novel immobilisation strategy for Saccharomyces cerevisiae cell cycle progression during ethanol production. 2020 , 104, 7495-7505	5
528	Releasing the microbes from old bones: the effect of different DNA extraction protocols on microbial community profiling. 2020 , 6, 1-15	1
527	Predicting non-carcinogenic hazard quotients of heavy metals in pepper (Capsicum annum L.) utilizing electromagnetic waves. 2020 , 14, 1	2
526	Neutrophil extracellular traps from healthy donors and HIV-1-infected individuals restrict HIV-1 production in macrophages. 2020 , 10, 19603	3
525	Rhizobium indicum sp. nov., isolated from root nodules of pea (Pisum sativum) cultivated in the Indian trans-Himalayas. 2020 , 43, 126127	12
524	Survey of rumen microbiota of domestic grazing yak during different growth stages revealed novel maturation patterns of four key microbial groups and their dynamic interactions. 2020 , 2, 23	7
523	Proteomic characterization of kefir milk by two-dimensional electrophoresis followed by mass spectrometry. 2020 , 55, e4635	6
522	Two representatives of lamiaceae essential oils and their main components cause changes in glutathione related enzymatic activities. 2020 , 1-7	
521	Molecular diversity of the microbial community in coloured snow from the Fildes Peninsula (King George Island, Maritime Antarctica). 2020 , 43, 1391-1405	7
520	Antiparasitic Activity of Bromotyrosine Alkaloids and New Analogues Isolated from the Fijian Marine Sponge Aplysinella rhax. 2020 , 17, e2000335	5
519	Bacterial and fungal diversity in the lorandite (TlAsS2) mine 'Allchar' in the Republic of North Macedonia. 2020 , 96,	5

518	The earthworm microbiome is resilient to exposure to biocidal metal nanoparticles. 2020, 267, 115633	7
517	Whole-genome sequence of multi-drug resistant Pseudomonas aeruginosa strains UY1PSABAL and UY1PSABAL2 isolated from human broncho-alveolar lavage, Yaound Cameroon. 2020 , 15, e0238390	8
516	Alpha-amylase conjugated biogenic silver nanoparticles as innovative strategy against biofilm-forming multidrug resistant bacteria. 2020 , 29, 101784	10
515	The gut bacterial community affects immunity but not metabolism in a specialist herbivorous butterfly. 2020 , 10, 8755-8769	4
514	Identification and enumeration of microbial organisms in grey water in the Khayelitsha informal settlement, Western Cape, South Africa. 2020 , 20, 1114-1126	1
513	Stable Carbon Isotope Studies of CH4 Dynamics Via Water and Plant Pathways in a Tropical Thai Paddy: Insights Into Diel CH4 Transportation. 2020 , 125, e2019JG005112	3
512	The Lipoxygenase Lox1 Is Involved in Light- and Injury-Response, Conidiation, and Volatile Organic Compound Biosynthesis in the Mycoparasitic Fungus. <i>Frontiers in Microbiology</i> , 2020 , 11, 2004 5.7	8
511	Effects of spatial heterogeneity on bacterial genetic circuits. 2020 , 16, e1008159	1
510	Genotyping and plant-derived glycan utilization analysis of Bifidobacterium strains from mother-infant pairs. 2020 , 20, 277	O
509	Quantitative analysis of the effects of morphological changes on extracellular electron transfer rates in cyanobacteria. 2020 , 13, 150	3
508	Characterization of salt-tolerant plant growth-promoting rhizobacteria and the effect on growth and yield of saline-affected rice. 2020 , 15, e0238537	31
507	Methylene blue active substances in plaque of Bacillus subtilis subsp. subtilis and enrichment by supplemental calcium in culture media. 2020 , 71, 550-556	
506	Comprehensive genome based analysis of Vibrio parahaemolyticus for identifying novel drug and vaccine molecules: Subtractive proteomics and vaccinomics approach. 2020 , 15, e0237181	7
505	Water disinfection by ozonation has advantages over UV irradiation in a brackish water recirculation aquaculture system for Pacific white shrimp (Litopenaeus vannamei). 2020 , 43, 1259-1285	1
504	High-Yielding Lovastatin Producer Aspergillus terreus Shows Increased Resistance to Inhibitors of Polyamine Biosynthesis. 2020 , 10, 8290	1
503	Current trends in polymerase chain reaction based detection of three major human pathogenic vibrios. 2020 , 1-19	3
502	Dimethylformamide Inhibits Fungal Growth and Aflatoxin B Biosynthesis in by Down-Regulating Glucose Metabolism and Amino Acid Biosynthesis. 2020 , 12,	1
501	Cyanobacterial polyhydroxybutyrate for sustainable bioplastic production: Critical review and perspectives. 2020 , 8, 104007	23

(2020-2020)

500	Differentiating microbial taxonomic and functional responses to physical disturbance in bulk and rhizosphere soils. 2020 , 31, 2858-2871	3
499	Differential Immune Response of Lactobacillus plantarum 286 Against Salmonella Typhimurium Infection in Conventional and Germ-Free Mice. 2021 , 1323, 1-17	2
498	Cold Atmospheric Plasma Inactivation of Microbial Spores Compared on Reference Surfaces and Powder Particles. 2020 , 13, 827-837	11
497	Novel recombinant Mce-truncated protein based ELISA for the diagnosis of Mycobacterium avium subsp. paratuberculosis infection in domestic livestock. 2020 , 15, e0233695	2
496	Veterinary antibiotic oxytetracycline effect on the soil microbial community. 2020, 44,	7
495	Response of sediment microbial communities to crude oil contamination in marine sediment microbial fuel cells under ferric iron stimulation. 2020 , 263, 114658	16
494	Bacterial synthesized metal and metal salt nanoparticles in biomedical applications: An up and coming approach. 2020 , 34, e5810	9
493	Antileishmanial activity of amphiphilic chlorin derivatives mediated by photodynamic therapy. 2020 , 31, 101769	5
492	Sustainable Production of Biosurfactants and Their Applications. 2020, 159-183	17
491	Experimental investigation of particle size effect on the self-healing performance of microcapsule for cemented coral sand. 2020 , 256, 119343	15
490	Antibacterial activity of Cymbopogon citratus against clinically important bacteria. 2020, 34, 26-30	3
489	Conjugation and Transport Processes. 2020 , 263-329	
488	Toxicity of three rare earth elements, and their combinations to algae, microcrustaceans, and fungi. 2020 , 201, 110795	14
487	Nuclear innate sensors for nucleic acids in immunity and inflammation. 2020 , 297, 162-173	12
486	Bacterioplankton abundance and community structure during post-monsoon in mangrove dominated estuaries of the Indian Sundarbans; An insight to biogeochemical processes. 2020 , 243, 106895	2
485	The distribution of functional N-cycle related genes and ammonia and nitrate nitrogen in soil profiles fertilized with mineral and organic N fertilizer. 2020 , 15, e0228364	4
484	Biarsenical fluorescent probes for multifunctional site-specific modification of proteins applicable in life sciences: an overview and future outlook. 2020 , 12, 1179-1207	2
483	Estimation of the Genome-Wide Mutation Rate and Spectrum in the Archaeal Species. 2020 , 215, 1107-1116	4

Synthesis of 3,15-Disuccinate-12-Coumarin Substituted Andrographolide Derivatives and Their Antiplatelet Aggregation Activities In Vitro. **2020**, 15, 1934578X2091086

481	In Vitro Antibacterial and Anti-Inflammatory Effects of Novel Insect Fungus Extract and Its Constituents against. 2020 , 9,	4
480	Modulation of bacterial multicellularity via spatio-specific polysaccharide secretion. 2020, 18, e3000728	18
479	Physical, Physiological, and Biological Measurements. 2020 , 35-57	
478	Determining incremental coulombic efficiency and physiological parameters of early stage Geobacter spp. enrichment biofilms. 2020 , 15, e0234077	4
477	Direct observation of independently moving replisomes in Escherichia coli. 2020 , 11, 3109	14
476	Identification and characterization of differentially expressed genes in Caenorhabditis elegans in response to pathogenic and nonpathogenic Stenotrophomonas maltophilia. 2020 , 20, 170	2
475	Epidemiology and antimicrobial resistance of spp. in animals in Sub-Saharan Africa: A systematic review. 2020 , 6, e03537	15
474	bloodstream infection in an allogeneic hematopoietic stem cell transplantation patient: Case report and review of literature. 2020 , 20, e00724	1
473	Gut microbiome shifts with urbanization and potentially facilitates a zoonotic pathogen in a wading bird. 2020 , 15, e0220926	28
472	Combined analysis of microbial metagenomic and metatranscriptomic sequencing data to assess in situ physiological conditions in the premature infant gut. 2020 , 15, e0229537	3
471	Thaw Transitions and Redox Conditions Drive Methane Oxidation in a Permafrost Peatland. 2020 , 125, e2019JG005526	9
470	The microbiome of modern microbialites in Bacalar Lagoon, Mexico. 2020 , 15, e0230071	11
469	Advancements in the gold standard: Measuring steroid sex hormones by mass spectrometry. 2020 , 82, 21-32	14
468	Vibrio splendidus O-antigen structure: a trade-off between virulence to oysters and resistance to grazers. 2020 , 22, 4264-4278	4
467	Impact of plants on the diversity and activity of methylotrophs in soil. 2020 , 8, 31	14
466	Aflatoxin Biosynthesis and Genetic Regulation: A Review. 2020 , 12,	65
465	Antibacterial chitosan electrostatic/covalent coating onto biodegradable poly (-lactic acid). 2020 , 105, 105835	10

(2020-2020)

464	Availability of Hydrogen Shapes the Microbial Abundance in Biofilm Anodes based on Geobacter Enrichment. 2020 , 7, 3720-3724	9
463	Seed biopriming with P- and K-solubilizing Enterobacter hormaechei sp. improves the early vegetative growth and the P and K uptake of okra (Abelmoschus esculentus) seedling. 2020 , 15, e0232860	12
462	Policymaking: from policy to the public with a stop at the lab. 2020 , 271-300	
461	The potential of drug repurposing to face bacterial and fungal biofilm infections. 2020 , 307-328	2
460	From hazard analysis to risk control using rapid methods in microbiology: A practical approach for the food industry. 2020 , 19, 1877-1907	10
459	Packaging for Various Applications. 2020 , 239-316	
458	Food, Pharmaceutical, and Medical Packaging. 2020 , 317-365	
457	Catalytic activity regulation through post-translational modification: the expanding universe of protein diversity. 2020 , 122, 97-125	1
456	Improvement of ruminal feed fermentation by addition of eucalyptus based mix essential oil. 2020 , 425, 012086	
455	Double-stranded DNA virioplankton dynamics and reproductive strategies in the oligotrophic open ocean water column. 2020 , 14, 1304-1315	31
454	In situ groundwater remediation with bioelectrochemical systems: A critical review and future perspectives. 2020 , 137, 105550	47
453	Terpinen-4-ol attenuates quorum sensing regulated virulence factors and biofilm formation in. 2020 , 15, 127-142	14
452	Mitigating the carbon footprint and improving productivity of ruminant livestock agriculture using a red seaweed. 2020 , 259, 120836	68
451	A potential antifungal and growth-promoting bacterium Bacillus sp. KTMA4 from tomato rhizosphere. 2020 , 142, 104049	13
450	Mobilizable antibiotic resistance genes are present in dust microbial communities. 2020 , 16, e1008211	16
449	Cyanobacterial sigma factors: Current and future applications for biotechnological advances. 2020 , 40, 107517	13
448	Body site-specific microbiota reflect sex and age-class among wild spotted hyenas. 2020 , 96,	10
447	Bacillus cereus spores and toxins - The potential role of biofilms. 2020 , 90, 103493	31

446	Synchronous degradation of phenol and aniline by sp.strain PB-1entrapped in sodium alginate-bamboo charcoal-chitosan beads. 2021 , 42, 4405-4414	1
445	In silico and functional analyses of immunomodulatory peptides encrypted in the human gut metaproteome. 2020 , 70, 103969	2
444	Draft Genome Sequence of an Escherichia coli Strain Harboring , , Aminoglycoside, Tetracycline, and Sulfonamide Resistance Genes, Isolated from a Costa Rican Wastewater Treatment Plant. 2020 , 9,	О
443	A dynamic microbial sulfur cycle in a serpentinizing continental ophiolite. 2020 , 22, 2329-2345	8
442	Reverse transcriptase enzyme and priming strategy affect quantification and diversity of environmental transcripts. 2020 , 22, 2383-2402	2
441	Ruminal fermentation, microbial population and lipid metabolism in gastrointestinal nematode-infected lambs fed a diet supplemented with herbal mixtures. 2020 , 15, e0231516	5
440	Engineering bacterial ACC deaminase for improving plant productivity under stressful conditions. 2020 , 259-277	9
439	Systematic evaluation of machine learning methods for identifying human-pathogen protein-protein interactions. 2021 , 22,	8
438	Modeling Methods and Numerical Tools for Process Validation/Control During High Pressure Processing. 2021 , 141-159	
437	Soil available nitrogen and phosphorus affected by functional bacterial community composition and diversity as ecological restoration progressed. 2021 , 32, 183-198	4
436	Production and potential biotechnological applications of microbial surfactants: An overview. 2021 , 28, 669-679	24
435	Advancements in protein nanoparticle vaccine platforms to combat infectious disease. 2021 , 13, e1681	13
434	Microalgal biosorption of heavy metals: A comprehensive bibliometric review. 2021 , 402, 123431	78
433	High Pressure Homogenization in Fruit and Vegetable Juice and Puree Processing: Effects on Quality, Stability and Phytochemical Profile. 2021 , 337-358	
432	Guideline for the management of feline sporotrichosis caused by Sporothrix brasiliensis and literature revision. 2021 , 52, 107-124	21
431	Recent advances in synthetic biology-enabled and natural whole-cell optical biosensing of heavy metals. 2021 , 413, 73-82	6
430	Impact of a reduced water salinity on the composition of Vibrio spp. in recirculating aquaculture systems for Pacific white shrimp (Litopenaeus vannamei) and its possible risks for shrimp health and food safety. 2021 , 44, 89-105	5
429	Effects of sulfonamide antibiotics on aquatic microbial community composition and functions. 2021 , 146, 106198	9

428	Inactivation of Listeria innocua on blueberries by novel ultrasound washing processes and their impact on quality during storage. 2021 , 121, 107580	9
427	A review of the biotransformations of priority pharmaceuticals in biological wastewater treatment processes. 2021 , 188, 116446	48
426	Phosphonates and SAR-trigger elicitors reducing the intensity of white mold in common bean. 2021 , 128, 273-278	
425	Ferric iron stimulation in marine SMFCs: Impact on the microbial structure evolution in contaminated sediments with low and high molecular weight PAHs. 2021 , 280, 111636	2
424	Gene gain and loss and recombination shape evolution of Listeria bacteriophages of the genus Pecentumvirus. 2021 , 113, 411-419	1
423	Complete genome sequence of highly pathogenic Vibrio parahaemolyticus isolated from mariculture Penaeus vannamei reveals virulence factor genes. 2021 , 52, 1401-1413	1
422	Development and application of galactomannan and essential oil-based edible coatings applied to \square coalho \square theese. 2021 , 45,	3
421	Map-based cloning identifies velvet A as a critical component of virulence in Fusarium pseudograminearum during infection of wheat heads. 2021 , 125, 191-200	1
420	Novel buffalo rumen metagenome derived acidic cellulase Cel-3.1 cloning, characterization, and its application in saccharifying rice straw and corncob biomass. 2021 , 170, 239-250	1
419	Sediment-based biochar facilitates highly efficient nitrate removal: Physicochemical properties, biological responses and potential mechanism. 2021 , 405, 126645	14
418	Growth inhibition of foodborne pathogens by co-microencapsulation of lactobacilli cell free and propolis extracts. 2021 , 41,	4
		Ĺ
417	Characteristics of plant growth-promoting rhizobacteria SCPG-7 and its effect on the growth of Capsicum annuum L. 2021 , 28, 11323-11332	1
417		
	Capsicum annuum L. 2021 , 28, 11323-11332 Microbial fuel cell-upflow biofilter coupling system for deep denitrification and power recovery:	1
416	Capsicum annuum L. 2021, 28, 11323-11332 Microbial fuel cell-upflow biofilter coupling system for deep denitrification and power recovery: Efficiencies, bacterial succession and interactions. 2021, 196, 110331 Genetic diversity, biofilm and virulence characteristics of Listeria monocytogenes in salmon sushi.	8
416 415	Capsicum annuum L. 2021, 28, 11323-11332 Microbial fuel cell-upflow biofilter coupling system for deep denitrification and power recovery: Efficiencies, bacterial succession and interactions. 2021, 196, 110331 Genetic diversity, biofilm and virulence characteristics of Listeria monocytogenes in salmon sushi. 2021, 140, 109871 Bacteriophages as drivers of bacterial virulence and their potential for biotechnological	1 8 3
416 415 414	Capsicum annuum L. 2021, 28, 11323-11332 Microbial fuel cell-upflow biofilter coupling system for deep denitrification and power recovery: Efficiencies, bacterial succession and interactions. 2021, 196, 110331 Genetic diversity, biofilm and virulence characteristics of Listeria monocytogenes in salmon sushi. 2021, 140, 109871 Bacteriophages as drivers of bacterial virulence and their potential for biotechnological exploitation. 2021, 45, A review on anammox process for the treatment of antibiotic-containing wastewater: Linking	1 8 3 16

410	Plant-Microbiome Interactions for Bacterial Wilt Suppression in Modern Tobacco Production. 2021 , 2-10	1
409	SLTRiP induces long lasting and protective T-cell memory response.	O
408	Linking plasmid-based beta-lactamases to their bacterial hosts using single-cell fusion PCR.	1
407	Microbial electrochemical technologies for wastewater treatment. 2021 , 287-306	O
406	Vibrio areninigrae as a pathogenic bacterium in a crustacean. 2021 , 178, 107517	2
405	Random Mutagenesis of Filamentous Fungi Strains for High-Yield Production of Secondary Metabolites: The Role of Polyamines.	
404	A novel high-content screening approach for the elucidation of C. jejuni biofilm composition and integrity. 2021 , 21, 2	1
403	The use of probes and bacteriophages for the detection of bacteria. 2021 , 49-93	
402	Stenotrophomonas maltophilia bacteremia in children: risk factors and mortality rate. 2021 , 10, 19	5
401	Insights into the microbial autotrophic potential of a shallow oligotrophic alpine pond. 2021 , 72, 899	
400	Structured environments foster competitor coexistence by manipulating interspecies interfaces. 2021 , 17, e1007762	2
399	The basic biological resources for the production of microbial pesticides. 2021 , 1-134	O
398	Strain dynamics of specific contaminant bacteria modulate the performance of ethanol biorefineries.	1
397	Evolutionary priming and transition to the ectomycorrhizal habit in an iconic lineage of mushroom-forming fungi: is preadaptation a requirement?.	2
396	Soil microbial inoculation during flood events shapes headwater stream microbial communities and diversity. 2021 , 82, 591-601	1
395	Surfactin stimulated by pectin molecular patterns and root exudates acts as a key driver of Bacillus-plant mutualistic interaction.	
394	H I I I I I I I I I I I I I I I I I I I	
393	Biological stoichiometry and growth dynamics of a diazotrophic cyanobacteria in nitrogen sufficient and deficient conditions. 2021 , 103, 102011	3

392	A Broadly Conserved Deoxycytidine Deaminase Protects Bacteria from Phage Infection.	6
391	The velvet protein Vel1 controls initial plant root colonization and conidia formation for xylem distribution in Verticillium wilt. 2021 , 17, e1009434	3
390	(+)-Terpinen-4-ol Inhibits Biofilm Formation by Upregulating the Interspecies Quorum Sensing Signals Diketopiperazines and Diffusing Signaling Factors. 2021 , 69, 3496-3510	3
389	Antimicrobial activities evaluation and phytochemical screening of some selected medicinal plants: A possible alternative in the treatment of multidrug-resistant microbes. 2021 , 16, e0249253	13
388	Legionella pneumophila CRISPR-Cas suggests recurrent encounters with Gokushovirinae.	
387	Spatiotemporal variation of bacterial communities in three cascade reservoirs in a southern city of China. 2021 , 21, 2532-2542	
386	Comparative assessment of immune evasion mechanisms in human whole-blood infection assays by a systems biology approach. 2021 , 16, e0249372	0
385	Nisin- and ripcin-derived hybrid lanthipeptides display selective antimicrobial activity against Staphylococcus aureus.	
384	Eden Benumbed: A Critique of Panqualityism and the Disclosure View of Consciousness. 1	O
383	Comparative efficacy of resorbable fiber wraps loaded with gentamicin sulfate or gallium maltolate in the treatment of osteomyelitis. 2021 , 109, 2255-2268	2
382	Determination of antibiotic resistance genes in a WWTP-impacted river in surface water, sediment, and biofilm: Influence of seasonality and water quality. 2021 , 768, 144526	12
381	Density-dependent effects are the main determinants of variation in growth dynamics between closely related bacterial strains.	
380	Differential surface competition and biofilm invasion strategies of Pseudomonas aeruginosa PA14 and PA01.	
379	Identification and differentiation of Pseudomonas species in field samples using an rpoD amplicon sequencing methodology.	
378	Monoclonal antibodies against cell wall chitooligomers as accessory tools for the control of cryptococcosis.	
377	High-throughput screening and selection of PCB-bioelectrocholeaching, electrogenic microbial communities using single chamber microbial fuel cells based on 96-well plate array.	1
376	Prognostic accuracy of MALDI-TOF mass spectrometric analysis of plasma in COVID-19. 2021 , 4,	9
375	Fungal biosurfactants, from nature to biotechnological product: bioprospection, production and potential applications. 2021 , 44, 2003-2034	11

374 Elevational changes in bacterial microbiota structure and diversity in an arthropod-disease vector.

373	Production of surfactin by Bacillus subtilis LAMI005 and evaluation of its potential as tensoactive and emulsifier.	1
372	The exposure of gadolinium at environmental relevant levels induced genotoxic effects in Arabidopsis thaliana (L.). 2021 , 215, 112138	3
371	Display of the self-sufficient CYP102A1 on the surface of E. coli-derived Outer Membrane Vesicles.	
370	Delivering the goods: Fungal secretion modulates virulence during hostpathogen interactions. 2021 , 36, 76-86	3
369	Functional Redundancy in Ocean Microbiomes Controls Trait Stability.	1
368	Fate and tissue distribution of Mycobacterium fortuitum through immersion challenge as a model of natural infection in Osphronemus goramy. 2021 , 29, 1979-1989	1
367	New Biocomposite Electrospun Fiber/Alginate Hydrogel for Probiotic Bacteria Immobilization. 2021 , 14,	4
366	Effectiveness of antimicrobial photodynamic therapy with indocyanine green against the standard and fluconazole-resistant Candida albicans. 2021 , 36, 1971-1977	2
365	Development and validation of an extended predictive model for the effect of pH and water activity on the growth kinetics of Geobacillus stearothermophilus in plant-based milk alternatives. 2021 , 145, 110407	2
364	Combined application of zinc-lysine chelate and zinc-solubilizing bacteria improves yield and grain biofortification of maize (Zea mays L.). 2021 , 16, e0254647	5
363	Essential gene acquisition destabilizes plasmid inheritance. 2021 , 17, e1009656	6
362	First historical genome of a crop bacterial pathogen from herbarium specimen: Insights into citrus canker emergence. 2021 , 17, e1009714	O
361	ASSEMBLY AND DYNAMICS OF MICROBIAL COMMUNITIES IN GRANULAR, FIXED-BIOFILM AND PLANKTONIC METHANOGENIC MICROBIOMES VALORIZING LONG CHAIN FATTY ACID (LCFA)-RICH WASTEWATER.	
360	Kinetic Properties of Microbial Exoenzymes Vary with Soil Depth but Have Similar Temperature Sensitivities Through the Soil Profile.	
359	Mechanisms of phosphatidylserine influence on viral production: a computational model of Ebola virus matrix protein assembly.	
358	HO Induces Major Phosphorylation Changes in Critical Regulators of Signal Transduction, Gene Expression, Metabolism and Developmental Networks in. 2021 , 7,	3
357	Inoculation of grape musts with single strains of Saccharomyces cerevisiae yeast reduces the diversity of chemical profiles of wines. 2021 , 16, e0254919	3

356	Silver nanoparticles-chitosan composites activity against resistant bacteria: tolerance and biofilm inhibition. 2021 , 23, 196	3
355	CRISPR-Cas9-mediated Large Cluster Deletion and Multiplex Genome Editing in Paenibacillus polymyxa.	O
354	Local meteorological conditions, shape and desiccation influence dispersal capabilities for airborne microorganisms. 2021 , 780, 146653	4
353	Unraveling Fe(II)-oxidizing mechanisms in a facultative Fe(II) oxidizer Sideroxydans lithotrophicus ES-1 via culturing, transcriptomics, and RT-qPCR.	1
352	Biosurfactant production by native marine bacteria (Acinetobacter calcoaceticus P1-1A) using waste carbon sources: Impact of process conditions. 2021 , 99, 2386	3
351	Evolution and origin of sliding clamp in bacteria, archaea and eukarya. 2021 , 16, e0241093	O
350	sp. nov. and sp. nov., two flavobacteria from the epiphytic microbiota of the brown alga, and emended description of the genus. 2021 , 71,	2
349	Genomic insights into an andean multiresistant soil actinobacterium of biotechnological interest. 2021 , 37, 166	2
348	Revealing the role of Plant Growth Promoting Rhizobacteria in suppressive soils against f.sp. based on metagenomic analysis. 2021 , 7, e07636	3
347	Recurrent micropollutant exposure leads to divergent degradation abilities and community succession in a freshwater microbiome.	
346	Microbial dark carbon fixation fueled by nitrate enrichment.	
345	Bacterial supergroup specific Costlof Wolbachia infections in Nasonia vitripennis.	
344	Bioprospecting thermotolerant yeasts from distillery effluent and molasses for high-temperature ethanol production. 2021 ,	O
343	Evolving a mitigation of the stress response pathway to change the basic chemistry of life.	
342	Classification of antimicrobial mechanism of action using bacterial time-lapse imaging.	
341	Bacterial diversity dynamics in microbial consortia selected for lignin utilization. 2021 , 16, e0255083	O
340	Prokaryotic diversity and composition within equatorial lakes Olbolosat and Oloiden in Kenya (Africa). 2021 , 2, 100066	2
339	Control of Methanogenesis in Dairy Animals. 2022 , 65-78	O

338	Response of soil protozoa to acid mine drainage in a contaminated terrace. 2022 , 421, 126790	9
337	A fluorescence-based yeast sensor for monitoring acetic acid. 2021 , 21, 303-313	2
336	Taxonomic and functional analyses reveal existence of virulence and antibiotic resistance genes in beach sand bacterial populations. 2021 , 203, 1753-1766	0
335	High-Throughput Inertial Focusing of Micrometer- and Sub-Micrometer-Sized Particles Separation. 2017 , 4, 1700153	33
334	Bioremediation: New Prospects for Environmental Cleaning by Fungal Enzymes. 2019, 17-52	2
333	Nanotechnology in Wastewater and the Capacity of Nanotechnology for Sustainability. 2020 , 1-45	3
332	Coordination of Fungal Secondary Metabolism and Development. 2020, 173-205	2
331	An Updated View of the Microbial Diversity in Deep Hypersaline Anoxic Basins. 2017 , 23-40	2
330	Ectomycorrhizal Diversity and Tree Sustainability. 2019 , 145-166	2
329	Restricting mycotoxins without killing the producers: a new paradigm in nano-fungal interactions. 2020 , 104, 2803-2813	8
328	Cultivable microscopic fungi from an underground chemosynthesis-based ecosystem: a preliminary study. 2018 , 63, 43-55	18
327	Biomethane: The energy storage, platform chemical and greenhouse gas mitigation target. 2017 , 46, 13-22	21
326	Ciliates as model organisms for the ecotoxicological risk assessment of heavy metals: A meta-analysis. 2020 , 199, 110669	10
325	Biostimulation of sulfate-reducing bacteria and metallic ions removal from coal mine-impacted water (MIW) using shrimp shell as treatment agent. 2020 , 398, 122893	15
324	Agricultural potential of rhizospheric Bacillus subtilis strains exhibiting varied efficiency of surfactin production. 2017 , 225, 802-809	16
323	Diverse coral reef invertebrates exhibit patterns of phylosymbiosis. 2020 , 14, 2211-2222	12
322	Adaptive engineering of a hyperthermophilic archaeon on CO and discovering the underlying mechanism by multi-omics analysis. 2016 , 6, 22896	19
321	Metal nanoparticles synthesis through natural phenolic acids. 2019 , 13, 771-777	25

320	Do more mechanistic models increase accuracy of prediction of metabolisable protein supply in ruminants?. 2019 , 59, 1991	2
319	Assessing the comparability of different DNA extraction and amplification methods in gut microbial community profiling. 2019 , 1, e000060	9
318	Ecogenomics and biogeochemical impacts of uncultivated globally abundant ocean viruses.	5
317	Higher stress and immunity responses are associated with higher mortality in reef-building coral exposed to a bacterial challenge.	3
316	The evolution of no-cost resistance at sub-MIC concentrations of streptomycin in Streptomyces coelicolor.	1
315	Synthetic Photosynthetic Consortia Define Interactions Leading to Robustness and Photoproduction.	2
314	Pherotype polymorphism inStreptococcus pneumoniaeand its effects on population structure and recombination.	1
313	Identification ofKlebsiellacapsule synthesis loci from whole genome data.	3
312	Nutrient dynamics and stream order influence microbial community patterns along a 2914 km transect of the Mississippi River.	1
311	How to name and classify your phage: an informal guide.	8
310	Bacterial lipopolysaccharide reduces the stability of avian and human influenza viruses.	
		1
309	The urinary microbiota composition remains stable over time and under various storage conditions.	1
309	The urinary microbiota composition remains stable over time and under various storage conditions. Epiphytic Microbiome of Grapes Berries Varies Between Phenological Timepoints, Growing Seasons and Regions.	
	Epiphytic Microbiome of Grapes Berries Varies Between Phenological Timepoints, Growing Seasons	1
308	Epiphytic Microbiome of Grapes Berries Varies Between Phenological Timepoints, Growing Seasons and Regions. Ecological theory applied to environmental metabolomes reveals compositional divergence despite	1
308	Epiphytic Microbiome of Grapes Berries Varies Between Phenological Timepoints, Growing Seasons and Regions. Ecological theory applied to environmental metabolomes reveals compositional divergence despite conserved molecular properties.	1 3
308 307 306	Epiphytic Microbiome of Grapes Berries Varies Between Phenological Timepoints, Growing Seasons and Regions. Ecological theory applied to environmental metabolomes reveals compositional divergence despite conserved molecular properties. Temperature regulation as a tool to program synthetic microbial community composition. Phage-Antibiotic Synergy is Driven by a Unique Combination of Antibacterial Mechanism of Action	1 1 3

302	Reverse transcriptase enzyme and priming strategy affect quantification and diversity of environmental transcripts.	1
301	Evidence for a growth zone for deep subsurface microbial clades in near-surface anoxic sediments.	2
300	In-host Modelling of COVID-19 in Humans.	20
299	Ancestral absence of electron transport chains in Patescibacteria and DPANN.	3
298	Seed biopriming with P- and K-solubilizing Enterobacter hormaechei sp. improves the early vegetative growth and the P and K uptake of okra (Abelmoschus esculentus) seedling.	2
297	Comparative co-occurrence network analyses of the cichlid fish gut microbiota: community conservation and diet-associated shifts.	1
296	Temperature restriction in entomopathogenic bacteria.	1
295	Multilevel community assembly of the tadpole gut microbiome.	2
294	Mechanisms and heterogeneity of mineral use by natural colonies of the cyanobacterium Trichodesmium.	1
293	Genomic features underlying the evolutionary transitions of Apibacter to honey bee gut symbionts.	Ο
292	Prognostic accuracy of MALDI mass spectrometric analysis of plasma in COVID-19.	1
291	Quantitative assessment of NCLDVBost interactions predicted by co-occurrence analyses.	3
290	Salicylate, Bile Acids and Extreme Acid Cause Fitness Tradeoffs for Multidrug Pumps in Escherichia coli K-12.	1
289	Systems-informed genome mining for electroautotrophic microbial production.	3
288	PlantBoil feedbacks between arbuscular- and ecto-mycorrhizal communities.	2
287	An orthogonal and pH-tunable sensor-selector for muconic acid biosynthesis in yeast.	1
286	Microbial diversity and metabolic potential in cyanotoxin producing cyanobacterial mats throughout a river network.	4
285	Convergent gain and loss of genomic islands drive lifestyle changes in plant-associated Pseudomonas.	2

284	Integrated soybean transcriptomics, metabolomics, and chemical genomics reveal the importance of the phenylpropanoid pathway and antifungal activity in resistance to the broad host range pathogenSclerotinia sclerotiorum.	2
283	The capsule regulatory network ofKlebsiella pneumoniaedefined by density-TraDISort.	1
282	Shrinkage improves estimation of microbial associations under different normalization methods.	6
281	Trait-based approach to bacterial growth efficiency.	5
280	Processing of grassland soil C-N compounds into soluble and volatile molecules is depth stratified and mediated by genomically novel bacteria and archaea.	4
279	Extracellular Electron Uptake by TwoMethanosarcinaSpecies.	4
278	Bacterial diversity and population dynamics during the fermentation of palm wine from Guerrero Mexico.	1
277	Dynamics of microbial contaminants is driven by selection during ethanol production.	1
276	Automation for the identification of Pseudomonas aeruginosa: Comparison of TDR-300B, VITEK $\!$	1
275	BalticMethanosarcinaandClostridiumcompete for electrons from metallic iron.	2
274	Emergence of stable coexistence in a complex microbial community through metabolic cooperation and spatio-temporal niche partitioning.	8
273	Cryptic inoviruses are pervasive in bacteria and archaea across Earth biomes.	2
272	In Situ Electrochemical Studies of the Terrestrial Deep Subsurface Biosphere at the Sanford Underground Research Facility, South Dakota, USA.	1
271	ANME-2d anaerobic methanotrophic archaea differ from other ANME archaea in lipid composition and carbon source.	1
270	A global perspective on microbial diversity in the terrestrial deep subsurface.	2
269	Community structure Lecosystem function relationships in the Congo Basin methane cycle depend on the physiological scale of function.	3
268	Genome-wide transposon screen of aPseudomonas syringae mexBmutant reveals the substrates of efflux transporters.	1
267	Microbial diversity of the Arabian Sea in the Oxygen minimum zones by metagenomics approach.	2

266	Direct observation of independently moving replisomes in Escherichia coli.	1
265	Dinoroseobacter shibaeouter membrane vesicles are enriched for the chromosome dimer resolution sitedif.	1
264	MreC and MreD balance the interaction between the elongasome proteins PBP2 and RodA.	1
263	Diversity patterns of marine heterotrophic culturable bacteria along vertical and latitudinal gradients.	1
262	Selection-based model of prokaryote pangenomes.	6
261	Large-scale network analysis captures biological features of bacterial plasmids.	1
260	Extracellular electron transfer-dependent anaerobic oxidation of ammonium by anammox bacteria.	2
259	LaeA Controls Citric Acid Production through Regulation of the Citrate Exporter-Encoding Gene in Aspergillus luchuensis mut 2020 , 86,	9
258	Pseudomonas aeruginosa RRALC3 Enhances the Biomass, Nutrient and Carbon Contents of Pongamia pinnata Seedlings in Degraded Forest Soil. 2015 , 10, e0139881	21
257	The Metagenome of Utricularia gibba's Traps: Into the Microbial Input to a Carnivorous Plant. 2016 , 11, e0148979	12
256	RmtA, a Putative Arginine Methyltransferase, Regulates Secondary Metabolism and Development in Aspergillus flavus. 2016 , 11, e0155575	24
255	A multi-objective constraint-based approach for modeling genome-scale microbial ecosystems. 2017 , 12, e0171744	26
254	Decomposition of recalcitrant carbon under experimental warming in boreal forest. 2017 , 12, e0179674	21
253	Assigning ecological roles to the populations belonging to a phenanthrene-degrading bacterial consortium using omic approaches. 2017 , 12, e0184505	26
252	Zika virus infection modulates the bacterial diversity associated with Aedes aegypti as revealed by metagenomic analysis. 2018 , 13, e0190352	24
251	Long-term use of cover crops and no-till shift soil microbial community life strategies in agricultural soil. 2018 , 13, e0192953	89
250	Development of an innovative methodology combining chemical fractionation and in vivo analysis to investigate the biological properties of cheese. 2020 , 15, e0242370	0
249	Identification of the Clostridial cellulose synthase and characterization of the cognate glycosyl hydrolase, CcsZ. 2020 , 15, e0242686	4

248	Global Analysis of the Fungal Microbiome in Cystic Fibrosis Patients Reveals Loss of Function of the Transcriptional Repressor Nrg1 as a Mechanism of Pathogen Adaptation. 2015 , 11, e1005308	54
247	Microwave-assisted green synthesis of silver nanoparticles using dried extracts of Chlorella vulgaris and antibacterial activity studies. 2020 , 9, 283-293	4
246	Phytosynthesis of selenium nanoparticles using the costus extract for bactericidal application against foodborne pathogens. 2020 , 9, 477-487	12
245	Regulation of Secondary Metabolism by the Velvet Complex Is Temperature-Responsive in Aspergillus. 2016 , 6, 4023-4033	33
244	Application of silver nanoparticles in food packages: a review. 2019 , 39, 793-802	53
243	Drug-Resistant Infections. 2017 , 433-448	6
242	Bacteriological, Clinical and Virulence Aspects of Aeromonas-associated Diseases in Humans. 2018 , 67, 137-149	22
241	Structure and Functions of Hydrocarbon-Degrading Microbial Communities in Bioelectrochemical Systems. 2020 , 12, 343	11
240	Coping with oxidative stress in extreme environments: the distinctive roles played by Acinetobacter sp. Ver3 superoxide dismutases.	
239	Secondary metabolites of HIle cells mediate protection of fungal reproductive and overwintering structures against fungivorous animals. 2021 , 10,	2
238	In vitro study on the potential fungicidal effects of atorvastatin in combination with some azole drugs against multidrug resistant Candida albicans. 2021 , 37, 191	1
237	Recent speciation and hybridization in Icelandic deep-sea isopods: An integrative approach using genomics and proteomics. 2021 , 31, 313	2
236	Fishmeal replacement by periphyton reduces the fish in fish out ratio and alimentation cost in gilthead sea bream Sparus aurata. 2021 , 11, 20990	2
235	Utilization of drought-tolerant bacterial strains isolated from harsh soils as a plant growth-promoting rhizobacteria (PGPR) 2022 , 29, 1760-1769	3
234	LAMP assays for the simple and rapid detection of clinically important urinary pathogens including the detection of resistance to 3rd generation cephalosporins. 2021 , 21, 1037	2
233	Isolation and Characterization of Two Methyltransferase Genes, AfuvipB and AfuvipC in Aspergillus fumigatus. 2015 , 43, 33-39	
232	Hierarchical effects of pro-inflammatory cytokines on the post-influenza susceptibility to pneumococcal coinfection.	1
231	Benzoate and Salicylate Tolerant Strains Lose Antibiotic Resistance during Laboratory Evolution of Escherichia coliK-12.	

230	Microbial Communities are Well Adapted to Disturbances in Energy Input.	
229	Single cell transcriptomics, mega-phylogeny and the genetic basis of morphological innovations in Rhizaria.	
228	Network analysis links genome-wide phenotypic and transcriptional stress responses in a bacterial pathogen with a large pan-genome.	ſ
227	Aerobic H2respiration enhances metabolic flexibility of methanotrophic bacteria.	
226	Growth-independent cross-feeding modifies boundaries for coexistence in a bacterial mutualism.)
225	Ubiquity and quantitative significance of bacterioplankton lineages inhabiting the oxygenated hypolimnion of deep freshwater lakes.	
224	Bacteria. 2017 , 175-183	
223	Post-translational thioamidation of methyl-coenzyme M reductase, a key enzyme in methanogenic and methanotrophic Archaea.	
222	Tracking electron uptake from a cathode intoShewanellacells: implications for generating maintenance energy from solid substrates.	ſ
221	Impact of Wolbachia on oxidative stress sensitivity in the parasitic wasp Asobara japonica. 2017 , 12, e0175974	Ĺ
220	Microbial network, phylogenetic diversity and community membership in the active layer across a permafrost thaw gradient.	Ĺ
219	A thermostable Cas9 with increased lifetime in human plasma.	2
218	CarrierSeq: a sequence analysis workflow for low-input nanopore sequencing.)
217	Fine scale diversification of endolithic microbial communities in the hyper-arid Atacama Desert.	
216	Genomes from uncultivated prokaryotes: a comparison of metagenome-assembled and single-amplified genomes.)
215	Comprehensive analysis of Verticillium nonalfalfae in silico secretome uncovers putative effector proteins expressed during hop invasion.	
214	Research Tools and Methods for the Analysis of Microbiota in Dairy Products. 2018 , 23-53	2
213	Phylogenetic distribution of roseobacticides in the Roseobacter group and their effect on microalgae.	

212	AnEscherichia colinitrogen starvation response is important for mutualistic coexistence withRhodopseudomonas palustris.	
211	Particle-templated emulsification for microfluidics-free digital biology.	
210	Taxa-driven functional shifts associated with stormflow in an urban stream microbial community.	
209	Environmental sources of bacteria and genetic variation in behavior influence host-associated microbiota.	1
208	Antagonistic Pleiotropy Is Unexpectedly Rare In New Mutations.	0
207	Long-term seasonality of marine photoheterotrophic bacteria reveals low cohesiveness within the different phylogroups.	O
206	Novel antimicrobial peptide discovery using machine learning and biophysical selection of minimal bacteriocin domains.	
205	First report on Bacterial Diversity of Potable Spring water of Indian Himalayan Region.	
204	Functional and genetic characterization of an incF-type multidrug resistance plasmid isolated from fresh spinach.	
203	In vitrocommunity synergy between bacterial soil isolates can be facilitated by pH stabilisation of the environment.	
202	Structured environments fundamentally alter dynamics and stability of ecological communities.	
201	Transcriptome analysis of mycorrhizal and non-mycorrhizal soybean plantlets upon infection with Fusarium virguliforme, one causal agent of sudden death syndrome.	
200	Survival of selected lactic acid bacteria isolated from pineapple waste in refrigerated pineapple juices.	
199	Functional and Comparative Genomics of Niche-Specific Adapted ActinomycetesKocuria rhizophilaStrain D2 Isolated from Healthy Human Gut.	1
198	Coculturing bacteria leads to reduced phenotypic heterogeneities.	0
197	16S rRNA:rDNA ratios and cell activity staining reveal consistent patterns of soil microbial activity.	O
196	Soil microbial communities with greater investment in resource acquisition have lower growth yield.	
195	Nomenclature Errors in Public 16S rRNA Gene Reference Databases.	1

194	RefSoil+: A reference for antimicrobial resistance genes on soil plasmids.	
193	Artificial selection of microbial communities to enhance degradation of recalcitrant polymers.	Ο
192	Charting the diversity of Uncultured Viruses of Archaea and Bacteria.	
191	Evolutionary dynamics of carbapenem-resistant Acinetobacter baumannii circulating in Chilean hospitals.	
190	Metabolite composition of sinking particles reflects a changing microbial community and differential metabolite degradation.	
189	Shifts in microbial transcription patterns prior to onset of Necrotizing enterocolitis may indicate low oxygen levels in the gut.	
188	A real-time multiplexed microbial growth intervalometer for capturing high resolution growth curves.	
187	Geographic Patterns of Bacterioplankton among Lakes of the Middle and Lower Reaches of the Yangtze River Basin, China.	
186	New Insights Into Wine Yeast Diversities. 2019 , 117-163	
185	Analysis of Vibrio cholerae genomes using a novel bioinformatic tool identifies new, active Type VI Secretion System gene clusters.	
184	Impact of Plant-Associated Microbial Communities on Host Plants Under Abiotic Stresses. 2019 , 303-340	Ο
183	Diversity decoupled from sulfur isotope fractionation in a sulfate reducing microbial community.	
182	Futuristic Methods in Virus Genome Evolution Using the Third-Generation DNA Sequencing and Artificial Neural Networks. 2019 , 485-513	
181	Distribution Pattern and Diversity Maintenance Mechanism of the Denitrifier Community in Alkaline Copper Mine Drainage.	
180	Bacteriophages as Biocontrol Agents of Biofilm Infections Associated with Abiotic Prosthetic Devices. 2019 , 81-99	
179	Microbial Biological Control of Diseases and Pests by PGPR and PGPF. 2019 , 75-122	1
178	Marine deep biosphere microbial communities assemble in near-surface sediments in Aarhus Bay.	2
177	Evaluating the stoichiometric trait distributions of cultured bacterial populations and uncultured microbial communities.	

The distribution of fitness effects among synonymous mutations in a gene under selection. 176 7 Lactobacillus rhamnosusLcr35[] as an effective treatment for preventingCandida albicansinfection 175 in preclinical models: first mechanistical insights. Manganese/iron-supported sulfate-dependent anaerobic oxidation of methane by archaea in lake 174 sediments. Defining the root endosphere and rhizosphere microbiomes from the World Olive Germplasm Collection. Temporal dynamics of Bacteria, Archaea and protists in equatorial coastal waters. 172 Deciphering the metabolic capabilities of Bifidobacteria using genome-scale metabolic models. 171 Depth and location influence prokaryotic and eukaryotic microbial community structure in New 170 Zealand fjords. Growth and rapid succession of methanotrophs effectively limit methane release during lake 169 overturn. Population structure determined by comparative amplicon gene sequencing in biofilm and 168 planktonic bacterial communities from pesticide-contaminated water. Gut microbiome shifts with urbanization and potentially facilitates a zoonotic pathogen in a wading 167 Substantial near-infrared radiation-driven photosynthesis of chlorophyllf-containing cyanobacteria 166 in a natural habitat. LaeA controls citric acid production through regulation of a citrate exporter 165 encodingcexAinAspergillus luchuensismut.kawachii. Gram-positive and Gram-negative Bacteria Share Common Principles to Coordinate Growth and the 164 2 Cell Cycle at the Single-cell Level. Dormancy dynamics and dispersal contribute to soil microbiome resilience. 163 Using null models to compare bacterial and microeukaryotic metacommunity assembly under 162 \circ shifting environmental conditions. PseudomonasmRNA 2.0: Boosting Gene Expression Through Enhanced mRNA Stability and 161 Translational Efficiency. Trophic Selective Pressures Organize the Composition of Endolithic Microbial Communities from 160 Global Deserts. Diversity analysis and metagenomic insights into the Antibiotic Resistance and Metal Resistances among Hot Spring Bacteriobiome-insinuating inherent environmental baseline levels of Antibiotic 159 and Metal tolerance.

158	Understanding the evolution of interspecies interactions in microbial communities.	О
157	Virus Replication in Multicellular Photosynthetic Life Forms. 1-17	
156	Identification of the Clostridial cellulose synthase and characterization of the cognate glycosyl hydrolase, CcsZ.	
155	Effects of spatial heterogeneity on bacterial genetic circuits.	
154	Environmental Biosensors: A Microbiological View. 2020 , 1-22	
153	Abiotic and biotic drivers underly short- and long-term soil respiration responses to experimental warming in a dryland ecosystem.	1
152	Contribution Evenness: A functional redundancy metric sensitive to trait stability in microbial communities.	1
151	The heart microbiome of insectivorous bats from Central and South Eastern Europe.	
150	Prevalence of Diversified Antibiotic Resistant Bacteria within Sanitation Related Facilities of Human Populated Workplaces in Abbottabad.	
149	The household resistome Ifrequency of beta-lactamases, class 1 integron and antibiotic resistant bacteria in the domestic environment.	
148	Boosting innate immunity: Asaia bacteria expressing a protein from Wolbachia determine macrophage activation and killing of Leishmania.	
147	Portiera gives new clues on the evolutionary history of whiteflies.	1
146	Constraints on the Efficiency of Electromicrobial Production.	О
145	Evolutionary conserved nitric oxide synthesis proteins responding to bacterial MAMPs are located to the endoplasmic reticulum and are also involved in secondary metabolite synthesis and sterol production.	
144	Treated wastewater irrigation promotes the spread of antibiotic resistance into subsoil pore-water.	
143	Curcumin-Sophorolipid nano-conjugate inhibits Candida albicans filamentation and biofilm development.	
142	An inside look at a biofilm: Pseudomonas aeruginosa flagella bio-tracking.	
141	The biogeography of Streptomyces in New Zealand enabled by high-throughput sequencing of genus-specific rpoB amplicons.	

140	Fecal metabolomics reveals products of dysregulated proteolysis and altered microbial metabolism in obesity-related osteoarthritis. 2021 ,	5
139	Unearthing the genes of plant-beneficial marine yeast - Wickerhamomyces anomalus strain MSD1.	O
138	Occurrence, identification, and antibiogram signatures of selected Enterobacteriaceae from Tsomo and Tyhume rivers in the Eastern Cape Province, Republic of South Africa. 2020 , 15, e0238084	4
137	Use of null models to compare the assembly of northeast Atlantic bacterial community in the presence of crude oil with either chemical dispersant or biosurfactant.	
136	RNA Interference (RNAi) in Tomato Crop Research. 2021, 2264, 163-175	
135	Environmental patterns of brown moss- and Sphagnum-associated microbial communities. 2020 , 10, 22412	2
134	Adaptation measures (AMs) and mitigation policies (MPs) to climate change and sustainable blue economy: a global perspective. 2021 , 12, 1344-1369	О
133	Molecular Epidemiology, Risk Factors and Clinical Outcomes of Carbapenem- and Polymyxin-Resistant Gram-negative Bacterial Infections in Pregnant Women and Infants: A Systematic Review.	
132	Press disturbance unveils different community structure, function and assembly of bacterial taxa and functional genes in mesocosm-scale bioreactors.	
131	Competitive exclusion of uropathogenic E. coli biofilm by Salmonella through matrix inhibition.	
130	Omics Approaches in Chickpea Fusarium Wilt Disease Management. 2020 , 57-72	О
129	Trichoderma as Biostimulant: Factors Responsible for Plant Growth Promotion. 2020 , 287-309	
128	Microbial Characterization of Andean Peatland⊠ Soil. 2020 , 87-93	О
127	Mathematical investigation of microbial quorum sensing under various flow conditions.	
126	Translation and Post-translational Modifications in Protein Biosynthesis. 2020, 595-665	1
125	Identification of Listeria monocytogenes in cattle meat using biochemical methods and amplification of the hemolysin gene. 2020 , 92 Suppl´1, e20180557	0
124	Metagenomic Insights into Microbial Metabolisms of a Sulfur-Influenced Glacial Ecosystem.	
123	Microbial community assembly in a multi-layer dendritic metacommunity.	

122	Unification of Environmental Metabolomics with Metacommunity Ecology.	0
121	Competitive ecosystems are robustly stabilized by structured environments.	
120	Investigation of a night soil compost psychrotrophic bacterium Glutamicibacter arilaitensis LJH19 for its safety, efficient hydrolytic and plant growth-promoting potential.	O
119	Polyamines Upregulate Cephalosporin C Production and Expression of Lactam Biosynthetic Genes in High-Yielding Strain. 2021 , 26,	3
118	Exploration of bacteria associated with Nudibranchs to control Vibrio spp 2020 , 530, 012031	
117	Ex vivo Immune Profiling in Patient Blood enables Quantification of innate immune Effector Functions.	
116	Evolution and origin of sliding clamp in bacteria, archaea and eukarya.	1
115	Leaf Endophytes and Their Bioactive Compounds. 2021 , 147-159	3
114	Nitric Oxide Regulates the Ralstonia solanacearum Type 3 Secretion System.	2
113	Annual phytoplankton dynamics in coastal waters from Fildes Bay, Western Antarctic Peninsula.	
112	Pharmacotherapy of Addiction to Amphetamine-Type Stimulants. 2021, 187-195	
111	Overcoming drug´resistance in TB: an update. 2020 , 15, 1607-1609	1
110	Probiotics for Management of Gastrointestinal Cancers. 2021 , 191-209	1
109	The exploration of Thermococcus barophilus lipidome reveals the widest variety of phosphoglycolipids in Thermococcales.	
108	Understanding the transmission of African swine fever in wild boars of South Korea: A simulation study for parameter estimation. 2021 ,	1
107	Profile of sulfur oxidizing bacteria in full-scale Biotrickling filter to remove H2S in biogas from in cassava starch industry. 2021 , 7, 677-685	1
106	Membrane adaptation in the hyperthermophilic archaeon Pyrococcus furiosus relies upon a novel strategy involving glycerol monoalkyl glycerol tetraether lipids.	
105	Health status of free-ranging pure and cross-mixed miniature swine population from Northeast Spain. 2021 ,	1

104	PTEN differentially regulates endocytosis, migration, and proliferation in the enteric protozoan parasite Entamoeba histolytica.		
103	Multiple groups of methanotrophic bacteria mediate methane oxidation in anoxic lake sediments.		Ο
102	Nutrient and salt depletion synergistically boosts glucose metabolism in individual bacteria.		
101	Evidence for quinol oxidase activity of ImoA, a novel NapC/NirT family protein from the neutrophilic Fe(II) oxidizing bacterium Sideroxydans lithotrophicus ES-1.		
100	Ectomycorrhizal Fungi Dominated the Root and Rhizosphere Microbial Communities of Two Willow Cultivars Grown for Six-Years in a Mixed-Contaminated Environment 2022 , 8,		О
99	Bacterial lifestyle switch in response to algal metabolites.		
98	Microbial diversity in intensively farmed lake sediment contaminated by heavy metals and identification of microbial taxa bioindicators of environmental quality 2022 , 12, 80		1
97	Three models of vaccination strategies against cryptococcosis in immunocompromised hosts using heat-killed Cryptococcus neoformans \(\text{Bg} \) l1.		
96	Effect of inoculum sources on autotrophic nitrogen removal in anaerobic hollow fiber membrane reactors. 2022 , 26, 102375		1
95	The Pseudomonas aeruginosa homeostasis enzyme AlgL clears the periplasmic space of accumulated alginate during polymer biosynthesis 2022 , 101560		1
94	Endophytic Fungi: Key Insights, Emerging Prospects, and Challenges in Natural Product Drug Discovery 2022 , 10,		3
93	Ecological Observations Based on Functional Gene Sequencing Are Sensitive to the Amplicon Processing Method.		O
92	Electrochemical profiles of bacteria isolated from crude oil on simple benzene compounds detection. 2021 , 948, 012026		
91	The Solvent Dimethyl Sulfoxide Affects Physiology, Transcriptome and Secondary Metabolism of 2021 , 7,		1
90	Chemistry of petroleum wastewater. 2022 , 7-18		
89	Evolutionary Genomics of Marine Bacteria and Archaea. 2022 , 327-354		Ο
88	Combination of MALDI-TOF Mass Spectrometry and Machine Learning for Rapid Antimicrobial Resistance Screening: The Case of spp <i>Frontiers in Microbiology</i> , 2021 , 12, 804484	5.7	1
87	Key Enzymes in Fatty Acid Synthesis Pathway for Bioactive Lipids Biosynthesis 2022 , 9, 851402		Ο

86	Dysbiosis of skin microbiome and gut microbiome in melanoma progression 2022 , 22, 63	2
85	A novel murine model of post-implantation malaria-induced preterm birth 2022 , 17, e0256060	0
84	Xanthomonas infection transforms the apoplast into an accessible and habitable niche for Salmonella enterica.	
83	and anti- activity of citral in combination with fluconazole 2022 , 14, 2045813	1
82	Mucosal microbiomes and Fusobacterium genomics in Vietnamese colorectal cancer patients.	О
81	Preparation and characterization of PLA-lemon essential oil nanofibrous membranes for the preservation of Mongolian cheese.	
80	The white koji fungus Aspergillus luchuensis mut. kawachii 2022,	
79	Impact of Feeding Postbiotics and Paraprobiotics Produced From on Colon Mucosa Microbiota in Broiler Chickens 2022 , 9, 859284	
78	The effects of antibiotic combination treatments on Pseudomonas aeruginosa tolerance evolution and coexistence with Stenotrophomonas maltophilia.	
77	Enteric Methane Emissions and Animal Performance in Dairy and Beef Cattle Production: Strategies, Opportunities, and Impact of Reducing Emissions 2022 , 12,	2
76	Deposit-feeding worms control subsurface ecosystem functioning in intertidal sediment with strong physical forcing.	
75	Studying the effect of oxygen availability and matrix structure on population density and inter-strain interactions of Listeria monocytogenes in different dairy model systems. 2022 , 156, 111118	
74	Plant growth promoting activity of bacteria isolated from Asian rice (Oryza sativa L.) are plant subspecies dependent.	
73	Characterization of the Gene Encoding S-adenosyl-L-methionine (AdoMet) Synthetase in ; Role in Secondary Metabolism and Penicillin Production 2021 , 10,	1
73 72		0
	Secondary Metabolism and Penicillin Production 2021, 10, Ultrasonic pre-treatment of for improved electrogenic response in a single chambered microbial	
72	Secondary Metabolism and Penicillin Production 2021, 10, Ultrasonic pre-treatment of for improved electrogenic response in a single chambered microbial fuel cell 2022, 12, 18 Simulated colonic fluid replicates the in vivo growth capabilities of Citrobacter rodentium mutants and highlights a critical role for the Cpx envelope stress response in mediating stressors	

68	Indoor green wall affects health-associated commensal skin microbiota and enhances immune regulation: a randomized trial among urban office workers 2022 , 12, 6518	5
67	Faster Growth Enhances Low Carbon Fuel and Chemical Production Through Gas Fermentation 2022 , 10, 879578	Ο
66	Customized low-cost high-throughput amplifier for electro-fluidic detection of cell volume changes in point-of-care applications 2022 , 17, e0267207	
65	Nutrient and salt depletion synergistically boosts glucose metabolism in individual Escherichia coli cells 2022 , 5, 385	1
64	Water Resources in Africa under Global Change: Monitoring Surface Waters from Space 2022 , 1-51	5
63	Detection of Enterotoxigenic Psychrotrophic Presumptive and Cereulide Producers in Food Products and Ingredients 2022 , 14,	Ο
62	Data_Sheet_1.docx. 2019 ,	
61	Data_Sheet_1.docx. 2020 ,	
60	Unveiling Differences in Root Defense Mechanisms Between Tolerant and Susceptible Olive Cultivars to 2022 , 13, 863055	0
59	Density fluctuations, homeostasis, and reproduction effects in bacteria 2022 , 5, 397	Ο
58	Consistency in defence and competitiveness trade-off in a planktonic predator-prey system.	O
57	Natural-Based Solutions for Bioremediation in Water Environment. 2022, 1-93	
56	PTEN differentially regulates endocytosis, migration, and proliferation in the enteric protozoan parasite Entamoeba histolytica 2022 , 18, e1010147	
55	Review on corrosion and corrosion scale formation upon unlined cast iron pipes in drinking water distribution systems. 2022 ,	3
54	Metagenomic insights into surface water microbial communities of a South Asian mangrove ecosystem 2022 , 10, e13169	0
53	Phylogenetic diversity of culturable marine actinobacteria isolated from the Havelock island, the Andamans, India. 2022 , 23, 100123	
52	Response surface methodological approach for optimization of photodynamic therapy of onychomycosis using chlorin e6 loaded nail penetration enhancer vesicles 2022 , 232, 112461	О
51	Punicalagin inhibits biofilm formation and virulence gene expression of Vibrio parahaemolyticus. 2022 , 139, 109045	1

50	Slipchip-based immunomagnetic separation combined with loop-mediated isothermal amplification for rapid detection of Bacillus cereus with tetracycline resistance gene tetL in pasteurized milk. 2022 , 109122	1
49	Cytochrome expression shifts in Geobacter sulfurreducens to maximize energy conservation in response to changes in redox conditions.	O
48	In vitro antioxidant, anticholinesterase, tyrosinase activity studies, and LC-MS/MS simultaneous determination of 37 bioactive compounds in Indigofera heterantha. 2022 , 148, 537-545	O
47	Phylogenomic diversity elucidates mechanistic insights into Lyme borreliae host association.	O
46	Adaptation of cyanobacteria to the endolithic light spectrum in hyper-arid deserts.	
45	The Effects of Lactic Acid Bacteria on Salmonella Biofilms. 2022 , 91, 278-285	1
44	WeCoNET: a hostpathogen interactome database for deciphering crucial molecular networks of wheat-common bunt cross-talk mechanisms. 2022 , 18,	1
43	Functional characterization of novel phosphate solubilizing bacteria, Chryseomicrobium imtechense, for enhanced strawberry growth and yield parameters. 2022 , 15,	
42	The nutritional value and yields of amaranth (Amaranthus hypochondriacus) cultivar silages compared to silage from corn (Zea mays) harvested at the milk stage grown in a hot-humid climate. 2022 , 289, 115336	Ο
41	Recycling of Polyurethanes. 413-427	
40	Molecular mechanisms associated with microbial biostimulant-mediated growth enhancement, priming and drought stress tolerance in maize plants. 2022 , 12,	6
39	Biodeterioration. 2022 , 1-30	O
38	A history of over 40 years of potentially pathogenic free-living amoeba studies in Brazil - a systematic review. 117,	1
37	Identification of high-risk contact areas between feral pigs and outdoor-raised pig operations in California: Implications for disease transmission in the wildlife-livestock interface. 2022 , 17, e0270500	
36	Light regulates the degradation of the regulatory protein VE-1 in the fungus Neurospora crassa. 2022 , 20,	1
35	Mutations in rv0678 confer low-level resistance to benzothiazinone DprE1 inhibitors in M. tuberculosis.	O
34	Variations in microbial diversity affect the stability and function of dark fermentation bioreactors.	
33	Development of zebrafish (Danio rerio) as an in vivo model for Borrelia burgdorferi infection.	

32	Within-community variation of interspecific divergence patterns in passerine gut microbiota. 2022 , 12,	О
31	An Insight into the Mechanisms of Homeostasis in Extremophiles. 2022 , 127115	O
30	Effect of active phenolic acids on properties of PLA-PHBV blend films. 2022, 33, 100894	2
29	Immunity priming uncouples the growth-defense tradeoff in tomato.	
28	Metagenomic Thermometer.	
27	Combined sensor-based monitoring of mycothiol redox potential and DNA-damage response in Corynebacterium glutamicum.	O
26	Geobacter sulfurreducens Linique metabolism results in cells with a high iron and lipid content.	
25	From empirical to data-driven host selection: a broad-host-range expression platform to facilitate chassis screening.	
24	Overexpression of llm1 Affects the Synthesis of Secondary Metabolites of Aspergillus cristatus. 2022 , 10, 1707	0
23	Transient glycolytic complexation of arsenate enhances resistance in the enteropathogen Vibrio cholerae.	
22	Simulated colonic fluid replicates the in vivo growth capabilities of Citrobacter rodentium cpxRA mutants and uncovers additive effects of Cpx regulated genes on fitness.	
21	Abrupt permafrost thaw triggers microbial bloom and increased activity of microbial predators.	
20	Implication of VelB in the development, pathogenicity, and secondary metabolism of Penicillium expansum. 2023 , 195, 112121	О
19	How to Completely Squeeze a FungusAdvanced Genome Mining Tools for Novel Bioactive Substances. 2022 , 14, 1837	1
18	Capsid structure of a metazoan fungal dsRNA megabirnavirus reveals its uniquely acquired structures.	О
17	Influence of metal cations on the viscoelastic properties of Escherichia coli biofilms.	O
16	The maize pathogen Ustilago maydis secretes glycoside hydrolases and carbohydrate oxidases directed towards components of the fungal cell wall.	0
15	Early season soil microbiome best predicts wheat grain quality.	O

14	Global mosquito virome profiling and mosquito spatial diffusion pathways revealed by marker-viruses.	O
13	Cardiolipin synthesis inPseudomonas fluorescensUM270 plays a relevant role in stimulating plant growth under salt stress.	O
12	D ivergent TLR2 and TLR4 activation by fungal spores and species diversity in dust from waste-sorting plants□	О
11	A methyltransferase LaeA regulates ganoderic acid biosynthesis in Ganoderma lingzhi. 13,	O
10	The global groundwater resistome: core ARGs and their dynamics - anin silicore-analysis of publicly available groundwater metagenomes.	O
9	Antibiotic resistance response to sulfamethoxazole from the intracellular and extracellular DNA fractions of activated sludge.	О
8	Predicting environmental stressor levels with machine learning: a comparison between amplicon sequencing, metagenomics, and total RNA sequencing based on taxonomically assigned data.	О
7	Effect of light spectra on cereal fungal pathogens, a review. 2022 ,	O
6	The velvet proteins CsVosA and CsVelB coordinate growth, cell wall integrity, sporulation, conidial viability and pathogenicity in the rubber anthracnose fungus Colletotrichum siamense. 2023 , 268, 127290	O
5	Curli fibers inEscherichia colibiofilms: the influence of water availability on amyloid structure and properties.	O
4	The Role of Chromatin and Transcriptional Control in the Formation of Sexual Fruiting Bodies in Fungi. 2022 , 86,	О
3	Abundant deep ocean heterotrophic bacteria are culturable.	O
2	Metabolic handoffs between multiple symbionts may benefit the deep-sea bathymodioline mussels.	O
1	The global regulator FpLaeB is required for the regulation of growth, development, and virulence in Fusarium pseudograminearum. 14,	O