Hana Yi

List of Publications by Year in descending order

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36 papers	2,876 citations	12 h-index	345118 36 g-index
36	36	36	2846
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Proposed minimal standards for the use of genome data for the taxonomy of prokaryotes. International Journal of Systematic and Evolutionary Microbiology, 2018, 68, 461-466.	0.8	2,359
2	Metagenomic Insights into the Bioaerosols in the Indoor and Outdoor Environments of Childcare Facilities. PLoS ONE, 2015, 10, e0126960.	1.1	75
3	Difference in the Gut Microbiome between Ovariectomy-Induced Obesity and Diet-Induced Obesity. Journal of Microbiology and Biotechnology, 2017, 27, 2228-2236.	0.9	56
4	Lung virome: New potential biomarkers for asthma severity and exacerbation. Journal of Allergy and Clinical Immunology, 2021, 148, 1007-1015.e9.	1.5	30
5	Epidemiology and Factors Related to Clinical Severity of Acute Gastroenteritis in Hospitalized Children after the Introduction of Rotavirus Vaccination. Journal of Korean Medical Science, 2017, 32, 465.	1.1	28
6	Cochleicola gelatinilyticus gen. nov., sp. nov., Isolated from a Marine Gastropod, Reichia luteostoma. Journal of Microbiology and Biotechnology, 2016, 26, 1439-1445.	0.9	25
7	Genomic insights into the taxonomic status of the three subspecies of Bacillus subtilis. Systematic and Applied Microbiology, 2014, 37, 95-99.	1.2	24
8	Real-time selective monitoring of allergenic Aspergillus molds using pentameric antibody-immobilized single-walled carbon nanotube-field effect transistors. RSC Advances, 2015, 5, 15728-15735.	1.7	21
9	Reclassification of Serpens flexibilis Hespell 1977 as Pseudomonas flexibilis comb. nov., with Pseudomonas tuomuerensis Xin et al. 2009 as a later heterotypic synonym. Systematic and Applied Microbiology, 2015, 38, 563-566.	1.2	19
10	Flavobacterium magnum sp. nov., Flavobacterium pallidum sp. nov., Flavobacterium crocinum sp. nov. and Flavobacterium album sp. nov. International Journal of Systematic and Evolutionary Microbiology, 2018, 68, 3837-3843.	0.8	19
11	Fully Automated Field-Deployable Bioaerosol Monitoring System Using Carbon Nanotube-Based Biosensors. Environmental Science &	4.6	18
12	Gemmobacter aquarius sp. nov., Runella rosea sp. nov. and Flavobacterium fluviale sp. nov., isolated from the Namhangang River system. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 5640-5647.	0.8	16
13	Alteration of Lung and Gut Microbiota in IL-13-Transgenic Mice Simulating Chronic Asthma. Journal of Microbiology and Biotechnology, 2020, 30, 1819-1826.	0.9	15
14	Distribution of bacteriocin genes in the lineages of Lactiplantibacillus plantarum. Scientific Reports, 2021, 11, 20063.	1.6	15
15	Mucilaginibacter celer sp. nov. and Aquirhabdus parva gen. nov., sp. nov., isolated from freshwater. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 5479-5487.	0.8	15
16	Burkholderia megalochromosomata sp. nov., isolated from grassland soil. International Journal of Systematic and Evolutionary Microbiology, 2015, 65, 959-964.	0.8	14
17	Burkholderia monticola sp. nov., isolated from mountain soil. International Journal of Systematic and Evolutionary Microbiology, 2015, 65, 504-509.	0.8	14
18	Flavobacterium gilvum sp. nov., isolated from stream water. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 153-157.	0.8	11

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19	Thalassotalea crassostreae sp. nov., isolated from Pacific oyster. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 2195-2198.	0.8	10
20	Non-contiguous finished genome sequence and description of the gliding bacterium Flavobacterium seoulense sp. nov Standards in Genomic Sciences, 2014, 9, 34.	1.5	9
21	Neisseria weaveri Andersen et al. 1993 is a later heterotypic synonym of Neisseria weaveri Holmes et al. 1993. International Journal of Systematic and Evolutionary Microbiology, 2015, 65, 463-464.	0.8	9
22	Paenibacillus crassostreae sp. nov., isolated from the Pacific oyster Crassostrea gigas. International Journal of Systematic and Evolutionary Microbiology, 2018, 68, 58-63.	0.8	9
23	Phenylobacterium parvum sp. nov., isolated from lake water. International Journal of Systematic and Evolutionary Microbiology, 2019, 69, 1169-1172.	0.8	9
24	Gramella salexigens sp. nov., isolated from seawater. International Journal of Systematic and Evolutionary Microbiology, 2018, 68, 2381-2385.	0.8	8
25	Microbiome profiling of uncinate tissue and nasal polyps in patients with chronic rhinosinusitis using swab and tissue biopsy. PLoS ONE, 2021, 16, e0249688.	1.1	6
26	Sphingorhabdus lutea sp. nov., isolated from sea water. International Journal of Systematic and Evolutionary Microbiology, 2019, 69, 3593-3598.	0.8	6
27	Cohnella candidum sp. nov., radiation-resistant bacterium from soil. Antonie Van Leeuwenhoek, 2019, 112, 1029-1037.	0.7	5
28	Interactions between NCR ⁺ ILC3s and the Microbiome in the Airways Shape Asthma Severity. Immune Network, 2021, 21, e25.	1.6	5
29	Human microbiome studies in Korea. Allergy Asthma & Respiratory Disease, 2016, 4, 311.	0.3	4
30	A Half-Day Genome Sequencing Protocol for Middle East Respiratory Syndrome Coronavirus. Frontiers in Microbiology, 2021, 12, 602754.	1.5	4
31	<i>Selenomonas</i> : A marker of asthma severity with the potential therapeutic effect. Allergy: European Journal of Allergy and Clinical Immunology, 2022, 77, 317-320.	2.7	4
32	Flavisolibacter galbus sp. nov., isolated from soil in Jeju Island. Antonie Van Leeuwenhoek, 2019, 112, 1559-1565.	0.7	3
33	Wolbachia Sequence Typing in Butterflies Using Pyrosequencing. Journal of Microbiology and Biotechnology, 2015, 25, 1410-1416.	0.9	3
34	Human Alphacoronavirus Universal Primers for Genome Amplification and Sequencing. Frontiers in Microbiology, 2022, 13, 789665.	1.5	3
35	Effect of luxS encoding a synthase of quorum-sensing signal molecule Al-2 of Vibrio vulnificus on mouse gut microbiome. Applied Microbiology and Biotechnology, 2022, 106, 3721-3734.	1.7	3
36	Synonymy of Micropolyspora internatus and Saccharomonospora viridis and Emended Description of Saccharomonospora viridis. Journal of Microbiology and Biotechnology, 2017, 27, 149-154.	0.9	2