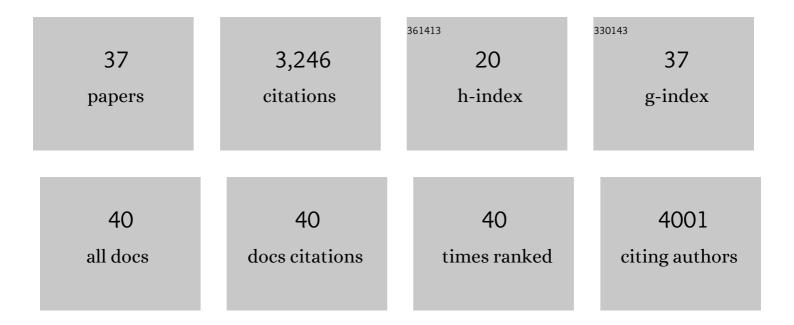
Seil Kim

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	EzTaxon: a web-based tool for the identification of prokaryotes based on 16S ribosomal RNA gene sequences. International Journal of Systematic and Evolutionary Microbiology, 2007, 57, 2259-2261.	1.7	1,960
2	Caproiciproducens galactitolivorans gen. nov., sp. nov., a bacterium capable of producing caproic acid from galactitol, isolated from a wastewater treatment plant. International Journal of Systematic and Evolutionary Microbiology, 2015, 65, 4902-4908.	1.7	153
3	Comparative Analysis of Primer–Probe Sets for RT-qPCR of COVID-19 Causative Virus (SARS-CoV-2). ACS Infectious Diseases, 2020, 6, 2513-2523.	3.8	111
4	Correlation between microbial community structure and biofouling in a laboratory scale membrane bioreactor with synthetic wastewater. Desalination, 2012, 287, 209-215.	8.2	98
5	A pilot scale two-stage anaerobic digester treating food waste leachate (FWL): Performance and microbial structure analysis using pyrosequencing. Process Biochemistry, 2014, 49, 301-308.	3.7	68
6	Comparison of the Bacterial Communities in Anaerobic, Anoxic, and Oxic Chambers of a Pilot A2O Process Using Pyrosequencing Analysis. Current Microbiology, 2013, 66, 555-565.	2.2	62
7	Oscillibacter ruminantium sp. nov., isolated from the rumen of Korean native cattle. International Journal of Systematic and Evolutionary Microbiology, 2013, 63, 1942-1946.	1.7	59
8	Megasphaera hexanoica sp. nov., a medium-chain carboxylic acid-producing bacterium isolated from a cow rumen. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 2114-2120.	1.7	57
9	Microbial population dynamics and proteomics in membrane bioreactors with enzymatic quorum quenching. Applied Microbiology and Biotechnology, 2013, 97, 4665-4675.	3.6	52
10	Co-culturing a novel Bacillus strain with Clostridium tyrobutyricum ATCC 25755 to produce butyric acid from sucrose. Biotechnology for Biofuels, 2013, 6, 35.	6.2	50
11	TTSV1, a new virus-like particle isolated from the hyperthermophilic crenarchaeote Thermoproteus tenax. Virology, 2006, 351, 280-290.	2.4	46
12	Marinifilum fragile gen. nov., sp. nov., isolated from tidal flat sediment. International Journal of Systematic and Evolutionary Microbiology, 2009, 59, 2241-2246.	1.7	42
13	Characterization of the Anti-Cancer Activity of the Probiotic Bacterium Lactobacillus fermentum Using 2D vs. 3D Culture in Colorectal Cancer Cells. Biomolecules, 2019, 9, 557.	4.0	42
14	Nucleic Acid Testing of SARS-CoV-2. International Journal of Molecular Sciences, 2021, 22, 6150.	4.1	42
15	Comparison of Digital PCR and Quantitative PCR with Various SARS-CoV-2 Primer-Probe Sets. Journal of Microbiology and Biotechnology, 2021, 31, 358-367.	2.1	41
16	Conversion of levulinic acid to 2-butanone by acetoacetate decarboxylase from Clostridium acetobutylicum. Applied Microbiology and Biotechnology, 2013, 97, 5627-5634.	3.6	28
17	Complete genome sequence of Bacillus sp. 275, producing extracellular cellulolytic, xylanolytic and ligninolytic enzymes. Journal of Biotechnology, 2017, 254, 59-62.	3.8	24
18	Clostridium aestuarii sp. nov., from tidal flat sediment. International Journal of Systematic and Evolutionary Microbiology, 2007, 57, 1315-1317.	1.7	22

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19	Clostridium ganghwense sp. nov., isolated from tidal flat sediment. International Journal of Systematic and Evolutionary Microbiology, 2006, 56, 691-693.	1.7	21
20	Complete genome of Kangiella geojedonensis KCTC 23420T, putative evidence for recent genome reduction in marine environments. Marine Genomics, 2015, 24, 215-217.	1.1	21
21	Discovery and characterization of a thermostable d-lactate dehydrogenase from Lactobacillus jensenii through genome mining. Process Biochemistry, 2013, 48, 109-117.	3.7	20
22	Anticancer Effects of Propionic Acid Inducing Cell Death in Cervical Cancer Cells. Molecules, 2021, 26, 4951.	3.8	20
23	Asticcacaulis solisilvae sp. nov., isolated from forest soil. International Journal of Systematic and Evolutionary Microbiology, 2013, 63, 3829-3834.	1.7	17
24	Comparison of 16S rRNA Gene Based Microbial Profiling Using Five Next-Generation Sequencers and Various Primers. Frontiers in Microbiology, 2021, 12, 715500.	3.5	16
25	Higher thermostability of l-lactate dehydrogenases is a key factor in decreasing the optical purity of d-lactic acid produced from Lactobacillus coryniformis. Enzyme and Microbial Technology, 2014, 58-59, 29-35.	3.2	15
26	Kitasatospora acidiphila sp. nov., isolated from pine grove soil, exhibiting antimicrobial potential. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 5567-5575.	1.7	15
27	Dynamic shift in community structures of biofilm-forming bacteria by the pre-treatment systems of seawater reverse osmosis processes. Desalination, 2014, 343, 17-25.	8.2	14
28	Burkholderia jirisanensis sp. nov., isolated from forest soil. International Journal of Systematic and Evolutionary Microbiology, 2016, 66, 1260-1267.	1.7	10
29	Development of SARS-CoV-2 packaged RNA reference material for nucleic acid testing. Analytical and Bioanalytical Chemistry, 2022, 414, 1773-1785.	3.7	10
30	Evaluating Cell Death Using Cell-Free Supernatant of Probiotics in Three-Dimensional Spheroid Cultures of Colorectal Cancer Cells. Journal of Visualized Experiments, 2020, , .	0.3	6
31	Complete Genome of Bacillus subtilis subsp. subtilis KCTC 3135T and Variation in Cell Wall Genes of B. subtilis Strains. Journal of Microbiology and Biotechnology, 2018, 28, 1760-1765.	2.1	6
32	Draft Genome Sequence of Bordetella trematum Strain HR18. Genome Announcements, 2015, 3, .	0.8	4
33	Draft Genome Sequence of Acinetobacter sp. HR7, Isolated from Hanwoo, Korean Native Cattle. Genome Announcements, 2015, 3, .	0.8	4
34	A Half-Day Genome Sequencing Protocol for Middle East Respiratory Syndrome Coronavirus. Frontiers in Microbiology, 2021, 12, 602754.	3.5	4
35	Complete genome of subsp. KCTC 3135 and variation in cell wall genes of strains. Journal of Microbiology and Biotechnology, 2018, , .	2.1	4
36	Human Alphacoronavirus Universal Primers for Genome Amplification and Sequencing. Frontiers in Microbiology, 2022, 13, 789665.	3.5	3

#	Article	IF	CITATIONS
37	Complete Genome Sequence of Paenibacillus sp. CAA11: A Promising Microbial Host for Lignocellulosic Biorefinery with Consolidated Processing. Current Microbiology, 2019, 76, 732-737.	2.2	1