

# Joon Yong Kim

## List of Publications by Year in descending order

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57  
papers

1,645  
citations

489802

18  
h-index

371746

37  
g-index

61  
all docs

61  
docs citations

61  
times ranked

2170  
citing authors

#	ARTICLE	IF	CITATIONS
1	Corrigendum to “Effects of the main ingredients of the fermented food, kimchi, on bacterial composition and metabolite profile” [Food Res. Int. 149 (2021) 110668]. Food Research International, 2022, 156, 110946.	2.9	0
2	Long-term population dynamics of viable microbes in a closed ecosystem of fermented vegetables. Food Research International, 2022, 154, 111044.	2.9	7
3	Safety assessment of white colony-forming yeasts in kimchi. Food Microbiology, 2022, 106, 104057.	2.1	4
4	Genomic analysis of facultatively oligotrophic haloarchaea of the genera Halarchaeum, Halorubrum, and Halolamina, isolated from solar salt. Archives of Microbiology, 2021, 203, 261-268.	1.0	2
5	Omics in gut microbiome analysis. Journal of Microbiology, 2021, 59, 292-297.	1.3	30
6	Anaerocolumna sedimenticola sp. nov., isolated from fresh water sediment. Antonie Van Leeuwenhoek, 2021, 114, 507-513.	0.7	8
7	ODFM, an omics data resource from microorganisms associated with fermented foods. Scientific Data, 2021, 8, 113.	2.4	11
8	Calf Diarrhea Caused by Prolonged Expansion of Autochthonous Gut Enterobacteriaceae and Their Lytic Bacteriophages. MSystems, 2021, 6, .	1.7	15
9	Salicibacter cibarius sp. nov. and Salicibacter cibi sp. nov., two novel species of the family Bacillaceae isolated from kimchi. Journal of Microbiology, 2021, 59, 460-466.	1.3	8
10	Aminipila terrae sp. nov., a strictly anaerobic bacterium isolated from river sediment. Archives of Microbiology, 2021, 203, 3163-3169.	1.0	9
11	Host habitat is the major determinant of the gut microbiome of fish. Microbiome, 2021, 9, 166.	4.9	100
12	Effects of the main ingredients of the fermented food, kimchi, on bacterial composition and metabolite profile. Food Research International, 2021, 149, 110668.	2.9	26
13	Complete genome sequence of Lactobacillus amylovorus 1394N20, a potential probiotic strain, isolated from a Hanwoo calf. Journal of Animal Science and Technology, 2021, 63, 1207-1210.	0.8	2
14	The human gut archaeome: identification of diverse haloarchaea in Korean subjects. Microbiome, 2020, 8, 114.	4.9	65
15	Haloplanus rubicundus sp. nov., an extremely halophilic archaeon isolated from solar salt. Systematic and Applied Microbiology, 2020, 43, 126085.	1.2	6
16	Impact of fermentation conditions on the diversity of white colony-forming yeast and analysis of metabolite changes by white colony-forming yeast in kimchi. Food Research International, 2020, 136, 109315.	2.9	19
17	Microbial niches in raw ingredients determine microbial community assembly during kimchi fermentation. Food Chemistry, 2020, 318, 126481.	4.2	66
18	Lentibacillus cibarius sp. nov., isolated from kimchi, a Korean fermented food. Journal of Microbiology, 2020, 58, 387-394.	1.3	16

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19	Raineyella fluvialis sp. nov., an actinobacterium isolated from freshwater sediment. International Journal of Systematic and Evolutionary Microbiology, 2020, 70, 4298-4304.	0.8	7
20	Salicibacter halophilus sp. nov., a moderately halophilic bacterium isolated from kimchi. Journal of Microbiology, 2019, 57, 997-1002.	1.3	7
21	Effects of an auxin-producing symbiotic bacterium on cell growth of the microalga Haematococcus pluvialis: Elevation of cell density and prolongation of exponential stage. Algal Research, 2019, 41, 101547.	2.4	31
22	Paracoccus jeotgali sp. nov., isolated from Korean salted and fermented shrimp. Journal of Microbiology, 2019, 57, 444-449.	1.3	6
23	Community structures and genomic features of undesirable white colony-forming yeasts on fermented vegetables. Journal of Microbiology, 2019, 57, 30-37.	1.3	20
24	Salicibacter kimchii gen. nov., sp. nov., a moderately halophilic and alkalitolerant bacterium in the family Bacillaceae, isolated from kimchi. Journal of Microbiology, 2018, 56, 880-885.	1.3	19
25	Role of jeotgal, a Korean traditional fermented fish sauce, in microbial dynamics and metabolite profiles during kimchi fermentation. Food Chemistry, 2018, 265, 135-143.	4.2	75
26	Novel haloarchaeon Natrinema thermophila having the highest growth temperature among haloarchaea with a large genome size. Scientific Reports, 2018, 8, 7777.	1.6	13
27	Halorubrum aethiopicum sp. nov., an extremely halophilic archaeon isolated from commercial rock salt. International Journal of Systematic and Evolutionary Microbiology, 2018, 68, 416-422.	0.8	10
28	Aquimarina seongsanensis sp. nov., isolated from sea water. Antonie Van Leeuwenhoek, 2017, 110, 1019-1025.	0.7	9
29	Complete genome sequence of Clostridium perfringens CBA7123 isolated from a faecal sample from Korea. Gut Pathogens, 2017, 9, 32.	1.6	4
30	Genomic Analysis of Bacillus licheniformis CBA7126 Isolated from a Human Fecal Sample. Frontiers in Pharmacology, 2017, 8, 724.	1.6	10
31	Genomic Analysis of a Pathogenic Bacterium, Paenicostridium sordellii CBA7122 Containing the Highest Number of rRNA Operons, Isolated from a Human Stool Sample. Frontiers in Pharmacology, 2017, 8, 840.	1.6	27
32	Complete genome sequence of a commensal bacterium, Hafnia alvei CBA7124, isolated from human feces. Gut Pathogens, 2017, 9, 41.	1.6	10
33	Aquimarina versatilis sp. nov., isolated from seashore sand, and emended description of the genus Aquimarina. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 411-416.	0.8	13
34	Maribacter pelagius sp. nov., isolated from seawater. International Journal of Systematic and Evolutionary Microbiology, 2017, 67, 3834-3839.	0.8	13
35	Genomic Analysis of Vulcanisaeta thermophila Type Strain CBA1501T Isolated from Solfataric Soil. Frontiers in Microbiology, 2016, 7, 1639.	1.5	0
36	Genome sequence of a commensal bacterium, Enterococcus faecalis CBA7120, isolated from a Korean fecal sample. Gut Pathogens, 2016, 8, 62.	1.6	9

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37	Genomic Analysis of the Extremely Halophilic Archaeon <i>Halobacterium noricense</i> CBA1132 Isolated from Solar Salt That Is an Essential Material for Fermented Foods. <i>Journal of Microbiology and Biotechnology</i> , 2016, 26, 1375-1382.	0.9	5
38	<i>Actibacter haliotis</i> sp. nov., isolated from the gut of an abalone, <i>Haliotis discus hannai</i> , and emended description of the genus <i>Actibacter</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 49-55.	0.8	25
39	<i>Corynebacterium atrinae</i> sp. nov., isolated from the gastrointestinal tract of a pen shell, <i>Atrina pectinata</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 531-536.	0.8	8
40	Genomic Analysis of the Moderately Haloalkaliphilic Bacterium <i>Oceanobacillus kimchii</i> Strain X50T with Improved High-Quality Draft Genome Sequences. <i>Journal of Microbiology and Biotechnology</i> , 2015, 25, 1971-1976.	0.9	1
41	<i>Polaribacter atrinae</i> sp. nov., isolated from the intestine of a comb pen shell, <i>Atrina pectinata</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2014, 64, 1654-1661.	0.8	26
42	<i>Cloacibacterium haliotis</i> sp. nov., isolated from the gut of an abalone, <i>Haliotis discus hannai</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2014, 64, 72-77.	0.8	21
43	<i>Acinetobacter apis</i> sp. nov., isolated from the intestinal tract of a honey bee, <i>Apis mellifera</i> . <i>Journal of Microbiology</i> , 2014, 52, 639-645.	1.3	44
44	Insect Gut Bacterial Diversity Determined by Environmental Habitat, Diet, Developmental Stage, and Phylogeny of Host. <i>Applied and Environmental Microbiology</i> , 2014, 80, 5254-5264.	1.4	591
45	<i>Dyella jejuensis</i> sp. nov., isolated from soil of Hallasan Mountain in Jeju Island. <i>Journal of Microbiology</i> , 2014, 52, 373-377.	1.3	23
46	<i>Actinomyces haliotis</i> sp. nov., a bacterium isolated from the gut of an abalone, <i>Haliotis discus hannai</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2014, 64, 456-461.	0.8	19
47	<i>Paenibacillus marinisediminis</i> sp. nov., a bacterium isolated from marine sediment. <i>Journal of Microbiology</i> , 2013, 51, 312-317.	1.3	18
48	<i>Weissella diestrammenae</i> sp. nov., isolated from the gut of a camel cricket ( <i>Diestrammena coreana</i> ). <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013, 63, 2951-2956.	0.8	29
49	<i>Gibbsiella papilionis</i> sp. nov., isolated from the intestinal tract of the butterfly <i>Mycalesis gotama</i> , and emended description of the genus <i>Gibbsiella</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013, 63, 2607-2611.	0.8	16
50	<i>Lactobacillus kimchiensis</i> sp. nov., isolated from a fermented food. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013, 63, 1355-1359.	0.8	20
51	<i>Shimia haliotis</i> sp. nov., a bacterium isolated from the gut of an abalone, <i>Haliotis discus hannai</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013, 63, 4248-4253.	0.8	33
52	<i>Pseudoruegeria haliotis</i> sp. nov., isolated from the gut of the abalone <i>Haliotis discus hannai</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013, 63, 4626-4632.	0.8	19
53	<i>Orbus sasakiae</i> sp. nov., a bacterium isolated from the gut of the butterfly <i>Sasakia charonda</i> , and emended description of the genus <i>Orbus</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013, 63, 1766-1770.	0.8	30
54	<i>Gillisia marina</i> sp. nov., from seashore sand, and emended description of the genus <i>Gillisia</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013, 63, 3640-3645.	0.8	16

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55	Genome sequence of the moderately halophilic bacterium <i>Salinicoccus carniancri</i> type strain CrmT (= DSM 23852T). <i>Standards in Genomic Sciences</i> , 2013, 8, 255-263.	1.5	11
56	<i>Enterococcus diestrammenae</i> sp. nov., isolated from the gut of <i>Diestrammena coreana</i> . <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2013, 63, 4540-4545.	0.8	11
57	Safety Assessment of White Colony-Forming Yeasts in Kimchi. <i>SSRN Electronic Journal</i> , 0, , .	0.4	0