## Sang-Cheol Park

## List of Publications by Citations

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Version: 2024-04-04

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

5 6,194 5 g-index

5 6,976 3 avg, IF L-index

#	Paper	IF	Citations
5	Introducing EzTaxon-e: a prokaryotic 16S rRNA gene sequence database with phylotypes that represent uncultured species. <i>International Journal of Systematic and Evolutionary Microbiology</i> , <b>2012</b> , 62, 716-721	2.2	4480
4	Towards a taxonomic coherence between average nucleotide identity and 16S rRNA gene sequence similarity for species demarcation of prokaryotes. <i>International Journal of Systematic and Evolutionary Microbiology</i> , <b>2014</b> , 64, 346-351	2.2	1657
3	Improved pipeline for reducing erroneous identification by 16S rRNA sequences using the Illumina MiSeq platform. <i>Journal of Microbiology</i> , <b>2015</b> , 53, 60-9	3	30
2	Large-Scale Genomics Reveals the Genetic Characteristics of Seven Species and Importance of Phylogenetic Distance for Estimating Pan-Genome Size. <i>Frontiers in Microbiology</i> , <b>2019</b> , 10, 834	5.7	16
1	Comparative genomics of Neisseria weaveri clarifies the taxonomy of this species and identifies genetic determinants that may be associated with virulence. <i>FEMS Microbiology Letters</i> , <b>2012</b> , 328, 100	)-5 <sup>2.9</sup>	11