

You-Tae Kim

List of Publications by Year in descending order

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Version: 2024-02-01

23
papers

717
citations

758635

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h-index

713013

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docs citations

23
times ranked

942
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 1 | Molecular interaction between methicillin-resistant <i>Staphylococcus aureus</i> (MRSA) and chicken breast reveals enhancement of pathogenesis and toxicity for food-borne outbreak. <i>Food Microbiology</i> , 2021, 93, 103602. | 2.1 | 15 |
| 2 | Characterization of bacteriophage VVP001 and its application for the inhibition of <i>Vibrio vulnificus</i> causing seafood-borne diseases. <i>Food Microbiology</i> , 2021, 94, 103630. | 2.1 | 25 |
| 3 | Cancer-protective effect of a synbiotic combination between <i>Lactobacillus gasseri</i> 505 and a <i>Cudrania tricuspidata</i> leaf extract on colitis-associated colorectal cancer. <i>Gut Microbes</i> , 2020, 12, 1785803. | 4.3 | 85 |
| 4 | Effects of Probiotics on the Symptoms and Surgical Outcomes after Anterior Resection of Colon Cancer (POSTCARE): A Randomized, Double-Blind, Placebo-Controlled Trial. <i>Journal of Clinical Medicine</i> , 2020, 9, 2181. | 1.0 | 26 |
| 5 | Microbiome Study of Initial Gut Microbiota from Newborn Infants to Children Reveals that Diet Determines Its Compositional Development. <i>Journal of Microbiology and Biotechnology</i> , 2020, 30, 1067-1071. | 0.9 | 14 |
| 6 | Role of Probiotics in Human Gut Microbiome-Associated Diseases. <i>Journal of Microbiology and Biotechnology</i> , 2019, 29, 1335-1340. | 0.9 | 221 |
| 7 | Oral consumption of cinnamon enhances the expression of immunity and lipid absorption genes in the small intestinal epithelium and alters the gut microbiota in normal mice. <i>Journal of Functional Foods</i> , 2018, 49, 96-104. | 1.6 | 3 |
| 8 | <i>Lelliottia jeotgali</i> sp. nov., isolated from a traditional Korean fermented clam. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2018, 68, 1725-1731. | 0.8 | 27 |
| 9 | Metagenomic Approach to Identifying Foodborne Pathogens on Chinese Cabbage. <i>Journal of Microbiology and Biotechnology</i> , 2018, 28, 227-235. | 0.9 | 32 |
| 10 | Analysis of Microbiota in Bellflower Root, <i>Platycodon grandiflorum</i> , Obtained from South Korea. <i>Journal of Microbiology and Biotechnology</i> , 2018, 28, 551-560. | 0.9 | 2 |
| 11 | Isolation and Characterization of <i>Bifidobacterium longum</i> subsp. <i>longum</i> BCBL-583 for Probiotic Applications in Fermented Foods. <i>Journal of Microbiology and Biotechnology</i> , 2018, 28, 1846-1849. | 0.9 | 1 |
| 12 | The complete genome sequence of <i>Cronobacter sakazakii</i> ATCC 29544T, a food-borne pathogen, isolated from a child's throat. <i>Gut Pathogens</i> , 2017, 9, 2. | 1.6 | 12 |
| 13 | Characterization and genome analysis of novel bacteriophages infecting the opportunistic human pathogens <i>Klebsiella oxytoca</i> and <i>K. pneumoniae</i> . <i>Archives of Virology</i> , 2017, 162, 1129-1139. | 0.9 | 18 |
| 14 | Genomic Insights and Its Comparative Analysis with <i>Yersinia enterocolitica</i> Reveals the Potential Virulence Determinants and Further Pathogenicity for Foodborne Outbreaks. <i>Journal of Microbiology and Biotechnology</i> , 2017, 27, 262-270. | 0.9 | 5 |
| 15 | Genomic Insights of <i>Weissella jogaejeotgali</i> FOL01T Reveals Its Food Fermentation Ability and Human Gut Adaptive Potential for Probiotic Applications in Food Industries. <i>Journal of Microbiology and Biotechnology</i> , 2017, 27, 943-946. | 0.9 | 6 |
| 16 | Biocontrol and Rapid Detection of Food-Borne Pathogens Using Bacteriophages and Endolysins. <i>Frontiers in Microbiology</i> , 2016, 7, 474. | 1.5 | 99 |
| 17 | Characterization and Genomic Study of the Novel Bacteriophage HY01 Infecting Both <i>Escherichia coli</i> O157:H7 and <i>Shigella flexneri</i> : Potential as a Biocontrol Agent in Food. <i>PLoS ONE</i> , 2016, 11, e0168985. | 1.1 | 59 |
| 18 | Complete genome sequence of <i>Vibrio parahaemolyticus</i> strain FORC_008, a foodborne pathogen from a flounder fish in South Korea. <i>Pathogens and Disease</i> , 2016, 74, ftw044. | 0.8 | 5 |

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|----|---|-----|-----------|
| 19 | Complete genome sequence of <i>Vibrio vulnificus</i> FORC_017 isolated from a patient with a hemorrhagic rash after consuming raw dotted gizzard shad. <i>Gut Pathogens</i> , 2016, 8, 22. | 1.6 | 14 |
| 20 | Complete genome sequence of the hyperthermophilic methanogen <i>Methanocaldococcus bathoardescens</i> JH146T isolated from the basalt seafloor. <i>Marine Genomics</i> , 2015, 24, 229-230. | 0.4 | 2 |
| 21 | <i>Methanocaldococcus bathoardescens</i> sp. nov., a hyperthermophilic methanogen isolated from a volcanically active deep-sea hydrothermal vent. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2015, 65, 1280-1283. | 0.8 | 37 |
| 22 | Complete genome sequence of hyperthermophilic archaeon <i>Thermococcus</i> sp. ES1. <i>Journal of Biotechnology</i> , 2014, 174, 14-15. | 1.9 | 7 |
| 23 | In vivo Trial of <i>Bifidobacterium longum</i> Revealed the Complex Network Correlations Between Gut Microbiota and Health Promotional Effects. <i>Frontiers in Microbiology</i> , 0, 13, . | 1.5 | 2 |