

Qiang Feng

List of Publications by Year in descending order

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24
papers

7,904
citations

567281

15
h-index

610901

24
g-index

24
all docs

24
docs citations

24
times ranked

12836
citing authors

#	ARTICLE	IF	CITATIONS
1	Using metagenomic data to boost protein structure prediction and discovery. Computational and Structural Biotechnology Journal, 2022, 20, 434-442.	4.1	3
2	Age-Related Cancer-Associated Microbiota Potentially Promotes Oral Squamous Cell Cancer Tumorigenesis by Distinct Mechanisms. Frontiers in Microbiology, 2022, 13, 852566.	3.5	6
3	Chitosan and Chitooligosaccharide: The Promising Non-Plant-Derived Prebiotics with Multiple Biological Activities. International Journal of Molecular Sciences, 2022, 23, 6761.	4.1	38
4	Chitosan Oligosaccharide Ameliorates Metabolic Syndrome Induced by Overnutrition via Altering Intestinal Microbiota. Frontiers in Nutrition, 2021, 8, 743492.	3.7	10
5	Upregulation of CPNE7 in mesenchymal stromal cells promotes oral squamous cell carcinoma metastasis through the NF- κ B pathway. Cell Death Discovery, 2021, 7, 294.	4.7	7
6	Soluble Dietary Fiber, One of the Most Important Nutrients for the Gut Microbiota. Molecules, 2021, 26, 6802.	3.8	81
7	Chitosan attenuates obesity by modifying the intestinal microbiota and increasing serum leptin levels in mice. Journal of Functional Foods, 2020, 64, 103659.	3.4	35
8	The Pathogenic Effects of Fusobacterium nucleatum on the Proliferation, Osteogenic Differentiation, and Transcriptome of Osteoblasts. Frontiers in Cell and Developmental Biology, 2020, 8, 807.	3.7	10
9	Alteration of salivary microbiome in periodontitis with or without type-2 diabetes mellitus and metformin treatment. Scientific Reports, 2020, 10, 15363.	3.3	46
10	Pathogenic and antimicrobial resistance genes in Streptococcus oralis strains revealed by comparative genome analysis. Genomics, 2020, 112, 3783-3793.	2.9	6
11	Changes of saliva microbiota in the onset and after the treatment of diabetes in patients with periodontitis. Aging, 2020, 12, 13090-13114.	3.1	29
12	Time-Course Transcriptome Analysis for Drug Repositioning in Fusobacterium nucleatum-Infected Human Gingival Fibroblasts. Frontiers in Cell and Developmental Biology, 2019, 7, 204.	3.7	10
13	<i>Fusobacterium nucleatum</i> Facilitates Apoptosis, ROS Generation, and Inflammatory Cytokine Production by Activating AKT/MAPK and NF- κ B Signaling Pathways in Human Gingival Fibroblasts. Oxidative Medicine and Cellular Longevity, 2019, 2019, 1-22.	4.0	69
14	Whole Body Vibration Triggers a Change in the Mutual Shaping State of Intestinal Microbiota and Body's Immunity. Frontiers in Bioengineering and Biotechnology, 2019, 7, 377.	4.1	6
15	Time-Course Transcriptome Analysis of Gingiva-Derived Mesenchymal Stem Cells Reveals That Fusobacterium nucleatum Triggers Oncogene Expression in the Process of Cell Differentiation. Frontiers in Cell and Developmental Biology, 2019, 7, 359.	3.7	27
16	Gut microbiome and serum metabolome alterations in obesity and after weight-loss intervention. Nature Medicine, 2017, 23, 859-868.	30.7	1,074
17	An LC-MS based untargeted metabolomics study identified novel biomarkers for coronary heart disease. Molecular BioSystems, 2016, 12, 3425-3434.	2.9	17
18	De novo assembly of a haplotype-resolved human genome. Nature Biotechnology, 2015, 33, 617-622.	17.5	73

#	ARTICLE	IF	CITATIONS
19	Gut microbiome development along the colorectal adenoma→carcinoma sequence. Nature Communications, 2015, 6, 6528.	12.8	1,062
20	The oral and gut microbiomes are perturbed in rheumatoid arthritis and partly normalized after treatment. Nature Medicine, 2015, 21, 895-905.	30.7	1,306
21	Dynamics and Stabilization of the Human Gut Microbiome during the First Year of Life. Cell Host and Microbe, 2015, 17, 690-703.	11.0	2,276
22	The Genome of a Mongolian Individual Reveals the Genetic Imprints of Mongolians on Modern Human Populations. Genome Biology and Evolution, 2014, 6, 3122-3136.	2.5	24
23	sapFinder: an R/Bioconductor package for detection of variant peptides in shotgun proteomics experiments. Bioinformatics, 2014, 30, 3136-3138.	4.1	25
24	An integrated catalog of reference genes in the human gut microbiome. Nature Biotechnology, 2014, 32, 834-841.	17.5	1,664