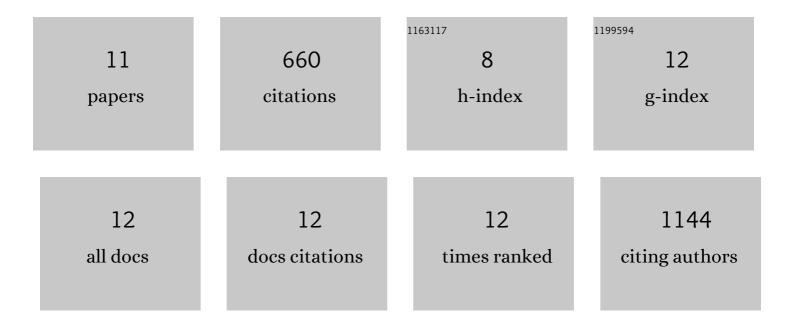
Tiejuan Shao

List of Publications by Year in descending order

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ΤΙΕΙΙΙΑΝ SHAO

#	Article	IF	CITATIONS
1	Alteration of Gut Microbiome and Correlated Amino Acid Metabolism Contribute to Hyperuricemia and Th17-Driven Inflammation in Uox-KO Mice. Frontiers in Immunology, 2022, 13, 804306.	4.8	18
2	Altered oral microbiota composition associated with recurrent aphthous stomatitis in young females. Medicine (United States), 2021, 100, e24742.	1.0	7
3	Combined effects of MSU crystals injection and high fat-diet feeding on the establishment of a gout model in C57BL/6 mice. Advances in Rheumatology, 2020, 60, 52.	1.7	13
4	Simiao Decoction Alleviates Gouty Arthritis by Modulating Proinflammatory Cytokines and the Gut Ecosystem. Frontiers in Pharmacology, 2020, 11, 955.	3.5	40
5	Qu-Zhuo-Tong-Bi Decoction Alleviates Gouty Arthritis by Regulating Butyrate-Producing Bacteria in Mice. Frontiers in Pharmacology, 2020, 11, 610556.	3.5	18
6	Effect of a Traditional Chinese Medicine Formula (CoTOL) on Serum Uric Acid and Intestinal Flora in Obese Hyperuricemic Mice Inoculated with Intestinal Bacteria. Evidence-based Complementary and Alternative Medicine, 2020, 2020, 1-10.	1.2	10
7	Whole Genome Sequencing of Extended-Spectrum Beta-Lactamase (ESBL)-Producing Escherichia coli Isolated From a Wastewater Treatment Plant in China. Frontiers in Microbiology, 2019, 10, 1797.	3.5	26
8	Alterations of the Gut Microbiota Associated With Promoting Efficacy of Prednisone by Bromofuranone in MRL/lpr Mice. Frontiers in Microbiology, 2019, 10, 978.	3.5	34
9	Linderae radix ethanol extract attenuates alcoholic liver injury via attenuating inflammation and regulating gut microbiota in rats. Brazilian Journal of Medical and Biological Research, 2019, 52, e7628.	1.5	20
10	Quantitative metagenomics reveals unique gut microbiome biomarkers in ankylosing spondylitis. Genome Biology, 2017, 18, 142.	8.8	268
11	Alterations of the gut microbiome in Chinese patients with systemic lupus erythematosus. Gut Pathogens, 2016, 8, 64.	3.4	195