Na Jiao

List of Publications by Year in descending order

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28 papers	1,178 citations	933447 10 h-index	794594 19 g-index
1 1			
32 all docs	32 docs citations	32 times ranked	1933 citing authors

#	Article	IF	CITATIONS
1	Multi-kingdom microbiota analyses identify bacterial–fungal interactions and biomarkers of colorectal cancer across cohorts. Nature Microbiology, 2022, 7, 238-250.	13.3	99
2	IBD Subtype-Regulators IFNG and GBP5 Identified by Causal Inference Drive More Intense Innate Immunity and Inflammatory Responses in CD Than Those in UC. Frontiers in Pharmacology, 2022, 13, 869200.	3.5	2
3	Microbial multidimensional signature assessment reveals microbial SNVs as the superior nonâ€invasive biomarkers for early detection of colorectal cancer. FASEB Journal, 2022, 36, .	0.5	O
4	Adverse Outcomes Associated With Corticosteroid Use in Critical COVID-19: A Retrospective Multicenter Cohort Study. Frontiers in Medicine, 2021, 8, 604263.	2.6	10
5	Altered gut microbiome in FUT2 lossâ€ofâ€function mutants in support of personalized medicine for inflammatory bowel diseases. FASEB Journal, 2021, 35, .	0.5	O
6	Fusobacterium nucleatum , a reproducible microbial marker for CRC prescreening. FASEB Journal, 2021, 35, .	0.5	0
7	Identification of microbial markers across populations in early detection of colorectal cancer. Nature Communications, 2021, 12, 3063.	12.8	109
8	Role of Guanylateâ€binding Protein 5 in Colonic Inflammation. FASEB Journal, 2021, 35, .	0.5	0
9	Comprehensive microbiota alterations in IBD and improved diagnostic accuracy for IBD using multiâ€kingdom microbial features. FASEB Journal, 2021, 35, .	0.5	O
10	Gastrointestinal sequelae 90 days after discharge for COVID-19. The Lancet Gastroenterology and Hepatology, 2021, 6, 344-346.	8.1	80
11	Altered gut microbiome in FUT2 loss-of-function mutants in support of personalized medicine for inflammatory bowel diseases. Journal of Genetics and Genomics, 2021, 48, 771-780.	3.9	21
12	Alterations in bile acid metabolizing gut microbiota and specific bile acid genes as a precision medicine to subclassify NAFLD. Physiological Genomics, 2021, 53, 336-348.	2.3	17
13	Putative Familial Transmissible Bacteria of Various Body Niches Link with Home Environment and Children's Immune Health. Microbiology Spectrum, 2021, 9, e0087221.	3.0	O
14	The roles of Qishen granules recipes, Qingre Jiedu, Wenyang Yiqi and Huo Xue, in the treatment of heart failure. Journal of Ethnopharmacology, 2020, 249, 112372.	4.1	13
15	Risk factors for the critical illness in SARS-CoV-2 infection: a multicenter retrospective cohort study. Respiratory Research, 2020, 21, 277.	3.6	8
16	Non-febrile COVID-19 patients were common and often became critically ill: a retrospective multicenter cohort study. Critical Care, 2020, 24, 314.	5.8	3
17	Enteric involvement in hospitalised patients with COVID-19 outside Wuhan. The Lancet Gastroenterology and Hepatology, 2020, 5, 534-535.	8.1	128
18	Identification of Key Factors in Cardiomyocyte Development by Singleâ€Cell Transcriptome Analysis. FASEB Journal, 2020, 34, 1-1.	0.5	0

#	Article	IF	Citations
19	FUT2 deficiency may influence the pathogenesis of inflammatory bowel diseases through gut microbiome. FASEB Journal, 2020, 34, 1-1.	0.5	O
20	NMI and IFP35 are key DAMP molecules in inflammatory bowel diseases. FASEB Journal, 2020, 34, 1-1.	0.5	0
21	In silico design of novel proton-pump inhibitors with reduced adverse effects. Frontiers of Medicine, 2019, 13, 277-284.	3.4	1
22	Keystone Species in the Pathogenic Process of NAFLD. FASEB Journal, 2019, 33, 496.40.	0.5	2
23	Gut microbiota contributes to the distinction between two traditional Chinese medicine syndromes of ulcerative colitis. World Journal of Gastroenterology, 2019, 25, 3242-3255.	3.3	37
24	Gut bacteria contributes to NAFLD pathogenesis by promoting secondary bile acids biosynthesis. FASEB Journal, 2019, 33, 126.4.	0.5	2
25	The Roles of Qishen Granules Recipes, Qingre Jiedu, Wenyang Yiqi and Huo Xue, in the Treatment of Heart Failure. FASEB Journal, 2019, 33, 818.3.	0.5	1
26	Suppressed hepatic bile acid signalling despite elevated production of primary and secondary bile acids in NAFLD. Gut, 2018, 67, 1881-1891.	12.1	438
27	Gut microbiome may contribute to insulin resistance and systemic inflammation in obese rodents: a meta-analysis. Physiological Genomics, 2018, 50, 244-254.	2.3	198
28	The response of archaeal species to seasonal variables in a subtropical aerated soil: insight into the low abundant methanogens. Applied Microbiology and Biotechnology, 2017, 101, 6505-6515.	3.6	7