

Na Jiao

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9084929/publications.pdf>

Version: 2024-02-01

28
papers

1,178
citations

933447

10
h-index

794594

19
g-index

32
all docs

32
docs citations

32
times ranked

1933
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Suppressed hepatic bile acid signalling despite elevated production of primary and secondary bile acids in NAFLD. <i>Gut</i> , 2018, 67, 1881-1891. | 12.1 | 438 |
| 2 | Gut microbiome may contribute to insulin resistance and systemic inflammation in obese rodents: a meta-analysis. <i>Physiological Genomics</i> , 2018, 50, 244-254. | 2.3 | 198 |
| 3 | Enteric involvement in hospitalised patients with COVID-19 outside Wuhan. <i>The Lancet Gastroenterology and Hepatology</i> , 2020, 5, 534-535. | 8.1 | 128 |
| 4 | Identification of microbial markers across populations in early detection of colorectal cancer. <i>Nature Communications</i> , 2021, 12, 3063. | 12.8 | 109 |
| 5 | Multi-kingdom microbiota analyses identify bacterial–fungal interactions and biomarkers of colorectal cancer across cohorts. <i>Nature Microbiology</i> , 2022, 7, 238-250. | 13.3 | 99 |
| 6 | Gastrointestinal sequelae 90 days after discharge for COVID-19. <i>The Lancet Gastroenterology and Hepatology</i> , 2021, 6, 344-346. | 8.1 | 80 |
| 7 | Gut microbiota contributes to the distinction between two traditional Chinese medicine syndromes of ulcerative colitis. <i>World Journal of Gastroenterology</i> , 2019, 25, 3242-3255. | 3.3 | 37 |
| 8 | Altered gut microbiome in FUT2 loss-of-function mutants in support of personalized medicine for inflammatory bowel diseases. <i>Journal of Genetics and Genomics</i> , 2021, 48, 771-780. | 3.9 | 21 |
| 9 | Alterations in bile acid metabolizing gut microbiota and specific bile acid genes as a precision medicine to subclassify NAFLD. <i>Physiological Genomics</i> , 2021, 53, 336-348. | 2.3 | 17 |
| 10 | The roles of Qishen granules recipes, Qingre Jiedu, Wenyang Yiqi and Huo Xue, in the treatment of heart failure. <i>Journal of Ethnopharmacology</i> , 2020, 249, 112372. | 4.1 | 13 |
| 11 | Adverse Outcomes Associated With Corticosteroid Use in Critical COVID-19: A Retrospective Multicenter Cohort Study. <i>Frontiers in Medicine</i> , 2021, 8, 604263. | 2.6 | 10 |
| 12 | Risk factors for the critical illness in SARS-CoV-2 infection: a multicenter retrospective cohort study. <i>Respiratory Research</i> , 2020, 21, 277. | 3.6 | 8 |
| 13 | The response of archaeal species to seasonal variables in a subtropical aerated soil: insight into the low abundant methanogens. <i>Applied Microbiology and Biotechnology</i> , 2017, 101, 6505-6515. | 3.6 | 7 |
| 14 | Non-febrile COVID-19 patients were common and often became critically ill: a retrospective multicenter cohort study. <i>Critical Care</i> , 2020, 24, 314. | 5.8 | 3 |
| 15 | Keystone Species in the Pathogenic Process of NAFLD. <i>FASEB Journal</i> , 2019, 33, 496.40. | 0.5 | 2 |
| 16 | Gut bacteria contributes to NAFLD pathogenesis by promoting secondary bile acids biosynthesis. <i>FASEB Journal</i> , 2019, 33, 126.4. | 0.5 | 2 |
| 17 | IBD Subtype-Regulators IFNG and GBP5 Identified by Causal Inference Drive More Intense Innate Immunity and Inflammatory Responses in CD Than Those in UC. <i>Frontiers in Pharmacology</i> , 2022, 13, 869200. | 3.5 | 2 |
| 18 | In silico design of novel proton-pump inhibitors with reduced adverse effects. <i>Frontiers of Medicine</i> , 2019, 13, 277-284. | 3.4 | 1 |

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|----|---|-----|-----------|
| 19 | 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 |
| 20 | Altered gut microbiome in FUT2 loss-of-function mutants in support of personalized medicine for inflammatory bowel diseases. FASEB Journal, 2021, 35, . | 0.5 | 0 |
| 21 | Fusobacterium nucleatum , a reproducible microbial marker for CRC prescreening. FASEB Journal, 2021, 35, . | 0.5 | 0 |
| 22 | Role of Guanylate-binding Protein 5 in Colonic Inflammation. FASEB Journal, 2021, 35, . | 0.5 | 0 |
| 23 | Comprehensive microbiota alterations in IBD and improved diagnostic accuracy for IBD using multi-kingdom microbial features. FASEB Journal, 2021, 35, . | 0.5 | 0 |
| 24 | Identification of Key Factors in Cardiomyocyte Development by Single-Cell Transcriptome Analysis. FASEB Journal, 2020, 34, 1-1. | 0.5 | 0 |
| 25 | FUT2 deficiency may influence the pathogenesis of inflammatory bowel diseases through gut microbiome. FASEB Journal, 2020, 34, 1-1. | 0.5 | 0 |
| 26 | NMI and IFP35 are key DAMP molecules in inflammatory bowel diseases. FASEB Journal, 2020, 34, 1-1. | 0.5 | 0 |
| 27 | Putative Familial Transmissible Bacteria of Various Body Niches Link with Home Environment and Children's Immune Health. Microbiology Spectrum, 2021, 9, e0087221. | 3.0 | 0 |
| 28 | Microbial multidimensional signature assessment reveals microbial SNVs as the superior non-invasive biomarkers for early detection of colorectal cancer. FASEB Journal, 2022, 36, . | 0.5 | 0 |