

Tao Wang

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7459028/publications.pdf>

Version: 2024-02-01

19
papers

23,011
citations

932766

10
h-index

794141

19
g-index

19
all docs

19
docs citations

19
times ranked

51332
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|------|-----------|
| 1 | Classical monocyte transcriptomes reveal significant anti-inflammatory statin effect in women with chronic HIV. <i>Cardiovascular Research</i> , 2021, 117, 1166-1177. | 1.8 | 8 |
| 2 | Treatment of Severe COVID-19 with Convalescent Plasma in Bronx, NYC. <i>JCI Insight</i> , 2021, 6, . | 2.3 | 36 |
| 3 | Differences in Vaginal Microbiota, Host Transcriptome, and Proteins in Women With Bacterial Vaginosis Are Associated With Metronidazole Treatment Response. <i>Journal of Infectious Diseases</i> , 2021, 224, 2094-2104. | 1.9 | 10 |
| 4 | Transcriptome-Wide Association Study of Blood Cell Traits in African Ancestry and Hispanic/Latino Populations. <i>Genes</i> , 2021, 12, 1049. | 1.0 | 11 |
| 5 | Molecular markers of risk of subsequent invasive breast cancer in women with ductal carcinoma in situ: protocol for a population-based cohort study. <i>BMJ Open</i> , 2021, 11, e053397. | 0.8 | 1 |
| 6 | Clinical Characteristics of Coronavirus Disease 2019 in China. <i>New England Journal of Medicine</i> , 2020, 382, 1708-1720. | 13.9 | 22,372 |
| 7 | Use of >100,000 NHLBI Trans-Omics for Precision Medicine (TOPMed) Consortium whole genome sequences improves imputation quality and detection of rare variant associations in admixed African and Hispanic/Latino populations. <i>PLoS Genetics</i> , 2019, 15, e1008500. | 1.5 | 203 |
| 8 | Adjustment for covariates using summary statistics of genome-wide association studies. <i>Genetic Epidemiology</i> , 2018, 42, 812-825. | 0.6 | 5 |
| 9 | Opposite Genetic Effects of CMIP Polymorphisms on the Risk of Type 2 Diabetes and Obesity: A Family-Based Study in China. <i>International Journal of Molecular Sciences</i> , 2018, 19, 1011. | 1.8 | 12 |
| 10 | Evaluation and Adaptation of a Laboratory-Based cDNA Library Preparation Protocol for Retrospective Sequencing of Archived MicroRNAs from up to 35-Year-Old Clinical FFPE Specimens. <i>International Journal of Molecular Sciences</i> , 2017, 18, 627. | 1.8 | 15 |
| 11 | Differential expression of circulating microRNAs according to severity of colorectal neoplasia. <i>Translational Research</i> , 2015, 166, 225-232. | 2.2 | 18 |
| 12 | Circulating Adipokines and Inflammatory Markers and Postmenopausal Breast Cancer Risk. <i>Journal of the National Cancer Institute</i> , 2015, 107, . | 3.0 | 83 |
| 13 | A Robust Method for Genome-Wide Association Meta-Analysis With the Application to Circulating Insulin-Like Growth Factor I Concentrations. <i>Genetic Epidemiology</i> , 2014, 38, 162-171. | 0.6 | 5 |
| 14 | The clinical significance of inflammatory cytokines in primary cell culture in endometrial carcinoma. <i>Molecular Oncology</i> , 2013, 7, 41-54. | 2.1 | 49 |
| 15 | Design and Statistical Analysis of Pooled Next Generation Sequencing for Rare Variants. <i>Journal of Probability and Statistics</i> , 2012, 2012, 1-19. | 0.3 | 4 |
| 16 | Hsa-miR-375 is differentially expressed during breast lobular neoplasia and promotes loss of mammary acinar polarity. <i>Journal of Pathology</i> , 2012, 226, 108-119. | 2.1 | 64 |
| 17 | Estimating Allele Frequency from Next-Generation Sequencing of Pooled Mitochondrial DNA Samples. <i>Frontiers in Genetics</i> , 2011, 2, 51. | 1.1 | 9 |
| 18 | A Prospective Study of Inflammation Markers and Endometrial Cancer Risk in Postmenopausal Hormone Nonusers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2011, 20, 971-977. | 1.1 | 83 |

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 19 | Resequencing of pooled DNA for detecting disease associations with rare variants. Genetic Epidemiology, 2010, 34, 492-501. | 0.6 | 23 |