

Tao Wang

List of Publications by Year in descending order

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

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19
papers

23,011
citations

933447

10
h-index

794594

19
g-index

19
all docs

19
docs citations

19
times ranked

51332
citing authors

#	ARTICLE	IF	CITATIONS
1	Clinical Characteristics of Coronavirus Disease 2019 in China. <i>New England Journal of Medicine</i> , 2020, 382, 1708-1720.	27.0	22,372
2	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.	3.5	203
3	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.	2.5	83
4	Circulating Adipokines and Inflammatory Markers and Postmenopausal Breast Cancer Risk. <i>Journal of the National Cancer Institute</i> , 2015, 107, .	6.3	83
5	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.	4.5	64
6	The clinical significance of inflammatory cytokines in primary cell culture in endometrial carcinoma. <i>Molecular Oncology</i> , 2013, 7, 41-54.	4.6	49
7	Treatment of Severe COVID-19 with Convalescent Plasma in Bronx, NYC. <i>JCI Insight</i> , 2021, 6, .	5.0	36
8	Resequencing of pooled DNA for detecting disease associations with rare variants. <i>Genetic Epidemiology</i> , 2010, 34, 492-501.	1.3	23
9	Differential expression of circulating microRNAs according to severity of colorectal neoplasia. <i>Translational Research</i> , 2015, 166, 225-232.	5.0	18
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.	4.1	15
11	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.	4.1	12
12	Transcriptome-Wide Association Study of Blood Cell Traits in African Ancestry and Hispanic/Latino Populations. <i>Genes</i> , 2021, 12, 1049.	2.4	11
13	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.	4.0	10
14	Estimating Allele Frequency from Next-Generation Sequencing of Pooled Mitochondrial DNA Samples. <i>Frontiers in Genetics</i> , 2011, 2, 51.	2.3	9
15	Classical monocyte transcriptomes reveal significant anti-inflammatory statin effect in women with chronic HIV. <i>Cardiovascular Research</i> , 2021, 117, 1166-1177.	3.8	8
16	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.	1.3	5
17	Adjustment for covariates using summary statistics of genome-wide association studies. <i>Genetic Epidemiology</i> , 2018, 42, 812-825.	1.3	5
18	Design and Statistical Analysis of Pooled Next Generation Sequencing for Rare Variants. <i>Journal of Probability and Statistics</i> , 2012, 2012, 1-19.	0.7	4

#	ARTICLE	IF	CITATIONS
19	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.	1.9	1