## Huijun Qian

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6305666/publications.pdf

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1163117 1474206 1,077 9 8 9 citations h-index g-index papers 14 14 14 3007 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	SMNN: batch effect correction for single-cell RNA-seq data via supervised mutual nearest neighbor detection. Briefings in Bioinformatics, 2021, 22, .	6.5	17
2	Whole genome sequence analyses of eGFR in 23,732 people representing multiple ancestries in the NHLBI trans-omics for precision medicine (TOPMed) consortium. EBioMedicine, 2021, 63, 103157.	6.1	14
3	Transcriptome-Wide Association Study of Blood Cell Traits in African Ancestry and Hispanic/Latino Populations. Genes, 2021, 12, 1049.	2.4	11
4	The Polygenic and Monogenic Basis of Blood Traits and Diseases. Cell, 2020, 182, 1214-1231.e11.	28.9	388
5	Common genetic risk variants identified in the SPARK cohort support DDHD2 as a candidate risk gene for autism. Translational Psychiatry, 2020, 10, 265.	4.8	56
6	Trans-ethnic and Ancestry-Specific Blood-Cell Genetics in 746,667 Individuals from 5 Global Populations. Cell, 2020, 182, 1198-1213.e14.	28.9	353
7	Genome-Wide Association of Kidney Traits in Hispanics/Latinos Using Dense Imputed Whole-Genome Sequencing Data. Circulation Genomic and Precision Medicine, 2020, 13, e002891.	3.6	6
8	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. PLoS Genetics, 2019, 15, e1008500.	3.5	203
9	Association of <i>APOL1</i> With Heart Failure With Preserved Ejection Fraction in Postmenopausal African American Women. JAMA Cardiology, 2018, 3, 712.	6.1	17