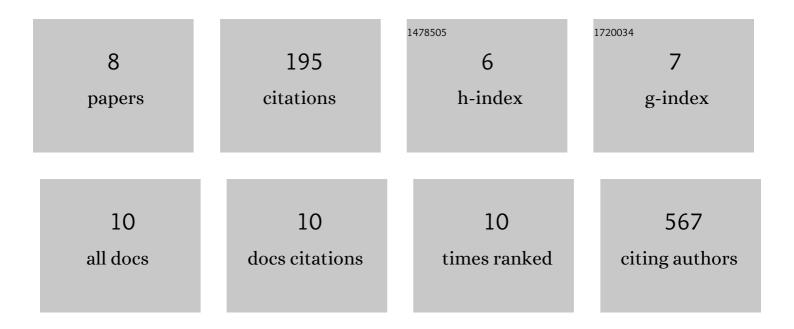
## Xinyuan Zhang

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/6647119/publications.pdf Version: 2024-02-01



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#	Article	IF	CITATIONS
1	Exome-wide evaluation of rare coding variants using electronic health records identifies new gene–phenotype associations. Nature Medicine, 2021, 27, 66-72.	30.7	44
2	Mendelian pathway analysis of laboratory traits reveals distinct roles for ciliary subcompartments in common disease pathogenesis. American Journal of Human Genetics, 2021, 108, 482-501.	6.2	7
3	Statistical Impact of Sample Size and Imbalance on Multivariate Analysis and A Case Study in the UK Biobank. AMIA Annual Symposium proceedings, 2020, 2020, 1383-1391.	0.2	Ο
4	Real world scenarios in rare variant association analysis: the impact of imbalance and sample size on the power in silico. BMC Bioinformatics, 2019, 20, 46.	2.6	20
5	Genomics-First Evaluation of Heart Disease Associated With Titin-Truncating Variants. Circulation, 2019, 140, 42-54.	1.6	97
6	Detecting potential pleiotropy across cardiovascular and neurological diseases using univariate, bivariate, and multivariate methods on 43,870 individuals from the eMERGE network. Pacific Symposium on Biocomputing Pacific Symposium on Biocomputing, 2019, 24, 272-283.	0.7	6
7	Rare variants in drug target genes contributing to complex diseases, phenome-wide. Scientific Reports, 2018, 8, 4624.	3.3	13
8	Detecting potential pleiotropy across cardiovascular and neurological diseases using univariate, bivariate, and multivariate methods on 43,870 individuals from the eMERGE network. , 2018, , .		6