

Xinyuan Zhang

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6647119/publications.pdf>

Version: 2024-02-01

8
papers

195
citations

1478505

6
h-index

1720034

7
g-index

10
all docs

10
docs citations

10
times ranked

567
citing authors

#	ARTICLE	IF	CITATIONS
1	Genomics-First Evaluation of Heart Disease Associated With Titin-Truncating Variants. <i>Circulation</i> , 2019, 140, 42-54.	1.6	97
2	Exome-wide evaluation of rare coding variants using electronic health records identifies new gene-phenotype associations. <i>Nature Medicine</i> , 2021, 27, 66-72.	30.7	44
3	Real world scenarios in rare variant association analysis: the impact of imbalance and sample size on the power in silico. <i>BMC Bioinformatics</i> , 2019, 20, 46.	2.6	20
4	Rare variants in drug target genes contributing to complex diseases, phenome-wide. <i>Scientific Reports</i> , 2018, 8, 4624.	3.3	13
5	Mendelian pathway analysis of laboratory traits reveals distinct roles for ciliary subcompartments in common disease pathogenesis. <i>American Journal of Human Genetics</i> , 2021, 108, 482-501.	6.2	7
6	Detecting potential pleiotropy across cardiovascular and neurological diseases using univariate, bivariate, and multivariate methods on 43,870 individuals from the eMERGE network. , 2018, , .		6
7	Detecting potential pleiotropy across cardiovascular and neurological diseases using univariate, bivariate, and multivariate methods on 43,870 individuals from the eMERGE network. <i>Pacific Symposium on Biocomputing Pacific Symposium on Biocomputing</i> , 2019, 24, 272-283.	0.7	6
8	Statistical Impact of Sample Size and Imbalance on Multivariate Analysis and A Case Study in the UK Biobank. <i>AMIA ... Annual Symposium proceedings</i> , 2020, 2020, 1383-1391.	0.2	0