Ying Zhou

List of Publications by Year
in descending order

15	39	1937685	1872680
papers	citations	4	6
		h-index	g-index

15	15	15	25
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Association detection between ordinal trait and rare variants based on adaptive combination of P values. Journal of Human Genetics, 2018, 63, 37-45.	2.3	10
2	Multiple Interval Mapping for Quantitative Trait Loci via EM Algorithm in the Presence of Crossover Interference. Communications in Statistics - Theory and Methods, 2010, 39, 3041-3057.	1.0	5
3	Simultaneous estimation of QTL effects and positions when using genotype data with errors. Journal of Genetics, 2015, 94, 27-34.	0.7	5
4	A nonparametric method to test for associations between rare variants and multiple traits. Genetical Research, 2016, 98, e1.	0.9	5
5	A two-step method for estimating QTL effects and positions in multi-marker analysis. Genetical Research, 2011, 93, 115-124.	0.9	4
6	A modified association test for rare and common variants based on affected sib-pair design. Journal of Theoretical Biology, 2019, 467, 1-6.	1.7	3
7	A modified two-stage approach for family-based genome-wide association studies. European Journal of Human Genetics, 2014, 22, 148-151.	2.8	2
8	Simultaneous estimation of QTL parameters for mapping multiple traits. Journal of Genetics, 2018, 97, 267-274.	0.7	2
9	An estimating function approach to linkage heterogeneity. Journal of Genetics, 2013, 92, 413-421.	0.7	1
10	Joint parameter estimation in the QTL mapping of ordinal traits. Journal of Theoretical Biology, 2017, 432, 100-108.	1.7	1
11	Simultaneous estimation of QTL parameters for mapping multiple traits. Journal of Genetics, 2018, 97, 267-274.	0.7	1
12	Gene association detection via local linear regression method. Journal of Human Genetics, 2020, 65, 115-123.	2.3	0
13	Bayesian interval mapping of count trait loci based on zero€inflated generalized Poisson regression model. Biometrical Journal, 2020, 62, 1428-1442.	1.0	0
14	A two-stage testing strategy for detecting genesÃ—environment interactions in association studies. G3: Genes, Genomes, Genetics, 2021, 11, .	1.8	0
15	Quantitative trait locus mapping analysis of multiple traits when using genotype data with potential errors. PeerJ, 2021, 9, e12187.	2.0	0