Min Yuan

List of Publications by Year in descending order

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		1684188	1372567
19	120	5	10
papers	citations	h-index	g-index
10	1.0	1.0	106
19	19	19	136
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Stratification of population in NHANES 2009–2014 based on exposure pattern of lead, cadmium, mercury, and arsenic and their association with cardiovascular, renal and respiratory outcomes. Environment International, 2021, 149, 106410.	10.0	51
2	Full covariate modelling approach in population pharmacokinetics: understanding the underlying hypothesis tests and implications of multiplicity. British Journal of Clinical Pharmacology, 2018, 84, 1525-1534.	2.4	17
3	Further Evaluation of Covariate Analysis using Empirical Bayes Estimates in Population Pharmacokinetics: the Perception of Shrinkage and Likelihood Ratio Test. AAPS Journal, 2017, 19, 264-273.	4.4	11
4	Basic Reproduction Number of the 2019 Novel Coronavirus Disease in the Major Endemic Areas of China: A Latent Profile Analysis. Frontiers in Public Health, 2021, 9, 575315.	2.7	6
5	Superiority of empirical Bayes estimator of the mean vector in multivariate normal distribution. Science China Mathematics, 2016, 59, 1175-1186.	1.7	5
6	A quick and accurate method for the estimation of covariate effects based on empirical Bayes estimates in mixed-effects modeling: Correction of bias due to shrinkage. Statistical Methods in Medical Research, 2019, 28, 3568-3578.	1.5	5
7	SCEBE: an efficient and scalable algorithm for genome-wide association studies on longitudinal outcomes with mixed-effects modeling. Briefings in Bioinformatics, 2021, 22, .	6.5	5
8	High-throughput and efficient multilocus genome-wide association study on longitudinal outcomes. Bioinformatics, 2020, 36, 3004-3010.	4.1	4
9	A forward selection algorithm to identify mutually exclusive alterations in cancer studies. Journal of Human Genetics, 2021, 66, 509-518.	2.3	4
10	Correcting for biases in affected sib-pair linkage analysis caused by uncertainty in sibling relationship. Science China Mathematics, 2012, 55, 1127-1135.	1.7	2
11	Correcting for differential genotyping error in genetic association analysis. Journal of Human Genetics, 2013, 58, 657-666.	2.3	2
12	Two-sided empirical Bayes test for location parameter in the gamma distribution. Communications in Statistics - Theory and Methods, 2017, 46, 4215-4225.	1.0	2
13	One-sided empirical Bayes test for location parameter in Gamma distribution. Applied Mathematics, 2018, 33, 287-297.	1.0	2
14	A novel quantification of information for longitudinal data analyzed by mixedâ€effects modeling. Pharmaceutical Statistics, 2020, 19, 388-398.	1.3	2
15	Efficient algorithms for covariate analysis with dynamic data using nonlinear mixed-effects model. Statistical Methods in Medical Research, 2021, 30, 233-243.	1.5	1
16	Fast and efficient correction for population stratification in multi-locus genome-wide association studies. Genetica, 2021, 149, 313-325.	1.1	1
17	Bayes factors based on robust TDT-type tests for family trio design. Statistical Applications in Genetics and Molecular Biology, 2015, 14, 253-64.	0.6	0
18	Quantitative Trait Loci Identification by Estimating the Genetic Model based on the Extremal Samples. Current Genomics, 2021, 22, 363-372.	1.6	0

#	Article	IF	CITATIONS
19	Bias correction for multiple covariate analysis using empirical bayesian estimation in mixed-effects models for longitudinal data. Computational Biology and Chemistry, 2022, 99, 107697.	2.3	O