

Li-An Lin

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8488677/publications.pdf>

Version: 2024-02-01

16
papers

1,737
citations

1039406

9
h-index

794141

19
g-index

20
all docs

20
docs citations

20
times ranked

5752
citing authors

#	ARTICLE	IF	CITATIONS
1	Integrative Analysis of Randomized Clinical Trial and Observational Study Data to Inform Post-marketing Safety Decision-Making. <i>Therapeutic Innovation and Regulatory Science</i> , 2022, 56, 423.	0.8	2
2	Estimation of causal effect in integrating randomized clinical trial and observational data – An example application to cardiovascular outcome trial. <i>Contemporary Clinical Trials</i> , 2021, 107, 106492.	0.8	2
3	Meta-analysis of blinded and unblinded studies for ongoing aggregate safety monitoring and evaluation. <i>Contemporary Clinical Trials</i> , 2020, 95, 106068.	0.8	3
4	Bayesian Meta-analysis of Safety Outcomes Using Blinded Clinical Trial Data. <i>Therapeutic Innovation and Regulatory Science</i> , 2020, 54, 1557-1565.	0.8	4
5	Bridging blinded and unblinded analysis for ongoing safety monitoring and evaluation. <i>Contemporary Clinical Trials</i> , 2019, 83, 81-87.	0.8	5
6	Protein-coding variants implicate novel genes related to lipid homeostasis contributing to body-fat distribution. <i>Nature Genetics</i> , 2019, 51, 452-469.	9.4	89
7	Bayesian regression model for recurrent event data with event-varying covariate effects and event effect. <i>Journal of Applied Statistics</i> , 2018, 45, 1260-1276.	0.6	6
8	Protein-altering variants associated with body mass index implicate pathways that control energy intake and expenditure in obesity. <i>Nature Genetics</i> , 2018, 50, 26-41.	9.4	286
9	Bayesian analysis of multi-type recurrent events and dependent termination with nonparametric covariate functions. <i>Statistical Methods in Medical Research</i> , 2017, 26, 2869-2884.	0.7	13
10	Rare and low-frequency coding variants alter human adult height. <i>Nature</i> , 2017, 542, 186-190.	13.7	544
11	An Empirical Comparison of Joint and Stratified Frameworks for Studying G × E Interactions: Systolic Blood Pressure and Smoking in the CHARGE Gene–Lifestyle Interactions Working Group. <i>Genetic Epidemiology</i> , 2016, 40, 404-415.	0.6	18
12	Meta-analysis identifies common and rare variants influencing blood pressure and overlapping with metabolic trait loci. <i>Nature Genetics</i> , 2016, 48, 1162-1170.	9.4	223
13	Fine-mapping of lipid regions in global populations discovers ethnic-specific signals and refines previously identified lipid loci. <i>Human Molecular Genetics</i> , 2016, 25, 5500-5512.	1.4	29
14	Rare and low-frequency variants and their association with plasma levels of fibrinogen, FVII, FVIII, and vWF. <i>Blood</i> , 2015, 126, e19-e29.	0.6	55
15	Epigenome-wide study identifies novel methylation loci associated with body mass index and waist circumference. <i>Obesity</i> , 2015, 23, 1493-1501.	1.5	152
16	Epigenome-wide association study (EWAS) of BMI, BMI change and waist circumference in African American adults identifies multiple replicated loci. <i>Human Molecular Genetics</i> , 2015, 24, 4464-4479.	1.4	289