Wonil Chung

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5155968/publications.pdf

Version: 2024-02-01

20 papers 2,878 citations

933447 10 h-index 18 g-index

22 all docs 22 docs citations

22 times ranked 7720 citing authors

#	Article	IF	CITATIONS
1	Grid-based Gaussian process models for longitudinal genetic data. Communications for Statistical Applications and Methods, 2022, 29, 745-763.	0.3	O
2	Grid-based Gaussian process models for longitudinal genetic data. Communications for Statistical Applications and Methods, 2022, 29, 65-83.	0.3	1
3	Bayesian mixed models for longitudinal genetic data: theory, concepts, and simulation studies. Genomics and Informatics, 2022, 20, e8.	0.8	1
4	Polygenic scores, diet quality, and type 2 diabetes risk: An observational study among 35,759 adults from 3 US cohorts. PLoS Medicine, 2022, 19, e1003972.	8.4	17
5	TwinEQTL: ultrafast and powerful association analysis for eQTL and GWAS in twin studies. Genetics, 2022, 221, .	2.9	O
6	Bayesian analysis of longitudinal traits in the Korea Association Resource (KARE) cohort. Genomics and Informatics, 2022, 20, e16.	0.8	0
7	Genomic-Metabolomic Associations Support the Role of LIPC and Glycerophospholipids in Age-Related Macular Degeneration. Ophthalmology Science, 2021, 1, 100017.	2.5	7
8	A transdisciplinary approach to understand the epigenetic basis of race/ethnicity health disparities. Epigenomics, 2021, 13, 1761-1770.	2.1	19
9	Statistical models and computational tools for predicting complex traits and diseases. Genomics and Informatics, 2021, 19, e36.	0.8	8
10	Shared genetic and experimental links between obesity-related traits and asthma subtypes in UK Biobank. Journal of Allergy and Clinical Immunology, 2020, 145, 537-549.	2.9	240
11	The Mediterranean diet, plasma metabolome, and cardiovascular disease risk. European Heart Journal, 2020, 41, 2645-2656.	2.2	138
12	Human Plasma Metabolomics in Age-Related Macular Degeneration: Meta-Analysis of Two Cohorts. Metabolites, 2019, 9, 127.	2.9	38
13	The nasal methylome as a biomarker of asthma and airway inflammation in children. Nature Communications, 2019, 10, 3095.	12.8	129
14	Genome-Wide Assessment for RestingÂHeart Rate and Shared Genetics WithÂCardiometabolic Traits and Type 2 Diabetes. Journal of the American College of Cardiology, 2019, 74, 2162-2174.	2.8	28
15	Efficient cross-trait penalized regression increases prediction accuracy in large cohorts using secondary phenotypes. Nature Communications, 2019, 10, 569.	12.8	50
16	A genome-wide cross-trait analysis from UK Biobank highlights the shared genetic architecture of asthma and allergic diseases. Nature Genetics, 2018, 50, 857-864.	21.4	191
17	Integrative approaches for large-scale transcriptome-wide association studies. Nature Genetics, 2016, 48, 245-252.	21.4	1,618
18	Fast eQTL Analysis for Twin Studies. Genetic Epidemiology, 2015, 39, 357-365.	1.3	9

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#	Article	IF	CITATIONS
19	Heritability and genomics of gene expression in peripheral blood. Nature Genetics, 2014, 46, 430-437.	21.4	370
20	Mixed-effects models for GAW18 longitudinal blood pressure data. BMC Proceedings, 2014, 8, S87.	1.6	10