

Yiqin Wang

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11539848/publications.pdf>

Version: 2024-02-01

18
papers

2,071
citations

623734

14
h-index

677142

22
g-index

23
all docs

23
docs citations

23
times ranked

6749
citing authors

#	ARTICLE	IF	CITATIONS
1	Association of mitochondrial DNA content, heteroplasmies and inter-generational transmission with autism. <i>Nature Communications</i> , 2022, 13, .	12.8	14
2	Accelerated expansion of pathogenic mitochondrial DNA heteroplasmies in Huntington's disease. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	15
3	STAMP: a multiplex sequencing method for simultaneous evaluation of mitochondrial DNA heteroplasmies and content. <i>NAR Genomics and Bioinformatics</i> , 2020, 2, lqaa065.	3.2	6
4	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.	21.4	286
5	Rare and low-frequency coding variants alter human adult height. <i>Nature</i> , 2017, 542, 186-190.	27.8	544
6	Genome-Wide Association Study Meta-Analysis of Long-Term Average Blood Pressure in East Asians. <i>Circulation: Cardiovascular Genetics</i> , 2017, 10, e001527.	5.1	26
7	Exome chip meta-analysis identifies novel loci and East Asian-specific coding variants that contribute to lipid levels and coronary artery disease. <i>Nature Genetics</i> , 2017, 49, 1722-1730.	21.4	129
8	Independent impacts of aging on mitochondrial DNA quantity and quality in humans. <i>BMC Genomics</i> , 2017, 18, 890.	2.8	116
9	Genetic Evidence for Elevated Pathogenicity of Mitochondrial DNA Heteroplasmy in Autism Spectrum Disorder. <i>PLoS Genetics</i> , 2016, 12, e1006391.	3.5	52
10	Age at menarche and age at natural menopause in East Asian women: a genome-wide association study. <i>Age</i> , 2016, 38, 513-523.	3.0	47
11	Coding-sequence variants are associated with blood lipid levels in 14,473 Chinese. <i>Human Molecular Genetics</i> , 2016, 25, 4107-4116.	2.9	14
12	Genome-wide association studies in East Asians identify new loci for waist-hip ratio and waist circumference. <i>Scientific Reports</i> , 2016, 6, 17958.	3.3	58
13	Trans-ancestry genome-wide association study identifies 12 genetic loci influencing blood pressure and implicates a role for DNA methylation. <i>Nature Genetics</i> , 2015, 47, 1282-1293.	21.4	294
14	Meta-analysis of genome-wide association studies of adult height in East Asians identifies 17 novel loci. <i>Human Molecular Genetics</i> , 2015, 24, 1791-1800.	2.9	105
15	Multiple Nonglycemic Genomic Loci Are Newly Associated With Blood Level of Glycated Hemoglobin in East Asians. <i>Diabetes</i> , 2014, 63, 2551-2562.	0.6	61
16	A Genome-Wide Association Study Identifies <i>GRK5</i> and <i>RASGRP1</i> as Type 2 Diabetes Loci in Chinese Hans. <i>Diabetes</i> , 2013, 62, 291-298.	0.6	166
17	Genome-Wide Association Study Meta-Analysis Reveals Transethnic Replication of Mean Arterial and Pulse Pressure Loci. <i>Hypertension</i> , 2013, 62, 853-859.	2.7	63
18	A Genome Wide Association Study Identifies Common Variants Associated with Lipid Levels in the Chinese Population. <i>PLoS ONE</i> , 2013, 8, e82420.	2.5	57