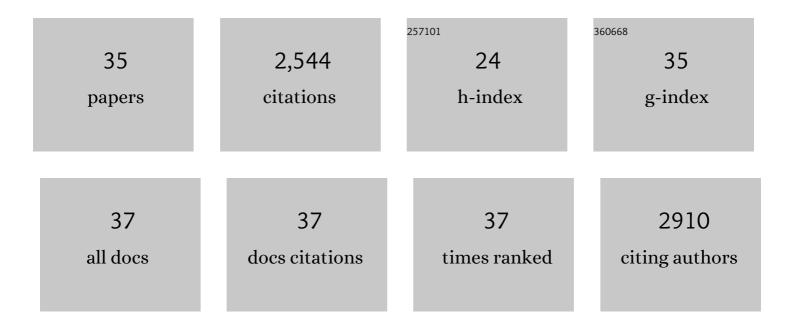
Wen Huang

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/855894/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	MetazExp: a database for gene expression and alternative splicing profiles and their analyses based on 53 615 public RNA-seq samples in 72 metazoan species. Nucleic Acids Research, 2022, 50, D1046-D1054.	6.5	7
2	Nitrogen removal characteristics and potential application of the heterotrophic nitrifying-aerobic denitrifying bacteria Pseudomonas mendocina S16 and Enterobacter cloacae DS'5 isolated from aquaculture wastewater ponds. Bioresource Technology, 2022, 345, 126541.	4.8	29
3	Identify known and novel candidate genes associated with backfat thickness in Duroc pigs by large-scale genome-wide association analysis. Journal of Animal Science, 2022, 100, .	0.2	16
4	Genome-Wide Analysis in Drosophila Reveals the Genetic Basis of Variation in Age-Specific Physical Performance and Response to ACE Inhibition. Genes, 2022, 13, 143.	1.0	5
5	Genetic basis of variation in cocaine and methamphetamine consumption in outbred populations of <i>Drosophila melanogaster</i> . Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	12
6	Epistasis for head morphology in <i>Drosophila melanogaster</i> . G3: Genes, Genomes, Genetics, 2021, 11, .	0.8	2
7	Rapid and Predictable Evolution of Admixed Populations Between Two <i>Drosophila</i> Species Pairs. Genetics, 2020, 214, 211-230.	1.2	42
8	Influence of Genetic Interactions on Polygenic Prediction. G3: Genes, Genomes, Genetics, 2020, 10, 109-115.	0.8	19
9	Genotype by environment interaction for gene expression in Drosophila melanogaster. Nature Communications, 2020, 11, 5451.	5.8	30
10	Leveraging Multiple Layers of Data To Predict <i>Drosophila</i> Complex Traits. G3: Genes, Genomes, Genetics, 2020, 10, 4599-4613.	0.8	21
11	Genome-wide association analyses identify known and novel loci for teat number in Duroc pigs using single-locus and multi-locus models. BMC Genomics, 2020, 21, 344.	1.2	43
12	Context-dependent genetic architecture of Drosophila life span. PLoS Biology, 2020, 18, e3000645.	2.6	47
13	Cene expression networks in the <i>Drosophila</i> Genetic Reference Panel. Genome Research, 2020, 30, 485-496.	2.4	55
14	Genetic Basis of Increased Lifespan and Postponed Senescence in Drosophila melanogaster. G3: Genes, Genomes, Genetics, 2020, 10, 1087-1098.	0.8	8
15	ASlive: a database for alternative splicing atlas in livestock animals. BMC Genomics, 2020, 21, 97.	1.2	8
16	Genetic Basis of Natural Variation in Spontaneous Grooming in <i>Drosophila melanogaster</i> . G3: Genes, Genomes, Genetics, 2020, 10, 3453-3460.	0.8	5
17	Effect of genetic architecture on the prediction accuracy of quantitative traits in samples of unrelated individuals. Heredity, 2018, 120, 500-514.	1.2	59
18	Charting the genotype–phenotype map: lessons from the <i>Drosophila melanogaster</i> Genetic Reference Panel. Wiley Interdisciplinary Reviews: Developmental Biology, 2018, 7, e289.	5.9	121

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19	Genetic and Genomic Response to Selection for Food Consumption in Drosophila melanogaster. Behavior Genetics, 2017, 47, 227-243.	1.4	20
20	The Genetic Architecture of Natural Variation in Recombination Rate in Drosophila melanogaster. PLoS Genetics, 2016, 12, e1005951.	1.5	102
21	The Genetic Architecture of Quantitative Traits Cannot Be Inferred from Variance Component Analysis. PLoS Genetics, 2016, 12, e1006421.	1.5	158
22	Genetic architecture of natural variation in visual senescence in <i>Drosophila</i> . Proceedings of the United States of America, 2016, 113, E6620-E6629.	3.3	46
23	Spontaneous mutations and the origin and maintenance of quantitative genetic variation. ELife, 2016, 5, .	2.8	63
24	Polymorphisms in early neurodevelopmental genes affect natural variation in alcohol sensitivity in adult drosophila. BMC Genomics, 2015, 16, 865.	1.2	54
25	Accounting for Genetic Architecture Improves Sequence Based Genomic Prediction for a Drosophila Fitness Trait. PLoS ONE, 2015, 10, e0126880.	1.1	50
26	Quantitative Genetics of Food Intake in Drosophila melanogaster. PLoS ONE, 2015, 10, e0138129.	1.1	84
27	The Genomic Basis of Postponed Senescence in Drosophila melanogaster. PLoS ONE, 2015, 10, e0138569.	1.1	40
28	The Genetic Basis for Variation in Olfactory Behavior in Drosophila melanogaster. Chemical Senses, 2015, 40, 233-243.	1.1	71
29	Genetic architecture of natural variation in <i>Drosophila melanogaster</i> aggressive behavior. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, E3555-63.	3.3	115
30	Genetic Architecture of Abdominal Pigmentation in Drosophila melanogaster. PLoS Genetics, 2015, 11, e1005163.	1.5	89
31	Genetic basis of transcriptome diversity in <i>Drosophila melanogaster</i> . Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, E6010-9.	3.3	134
32	Natural variation in genome architecture among 205 <i>Drosophila melanogaster</i> Genetic Reference Panel lines. Genome Research, 2014, 24, 1193-1208.	2.4	565
33	Genome-Wide Association Analysis of Tolerance to Methylmercury Toxicity in Drosophila Implicates Myogenic and Neuromuscular Developmental Pathways. PLoS ONE, 2014, 9, e110375.	1.1	42
34	Precipitation in the Yellow River drainage basin and East Asian monsoon strength on a decadal time scale. Quaternary Research, 2012, 78, 486-491.	1.0	31
35	Epistasis dominates the genetic architecture of <i>Drosophila</i> quantitative traits. Proceedings of the United States of America, 2012, 109, 15553-15559.	3.3	348