Wenhua Wei

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/370597/publications.pdf

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		304743	276875
56	1,939	22	41
papers	citations	h-index	g-index
59	59	59	3713
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Detecting epistasis in human complex traits. Nature Reviews Genetics, 2014, 15, 722-733.	16.3	372
2	Meta-Analysis of Genome-Wide Association Studies for Abdominal Aortic Aneurysm Identifies Four New Disease-Specific Risk Loci. Circulation Research, 2017, 120, 341-353.	4.5	166
3	Selenium-containing green tea has higher antioxidant and prebiotic activities than regular green tea. Food Chemistry, 2009, 114, 829-835.	8.2	102
4	Mouse models for human hyperuricaemia: a critical review. Nature Reviews Rheumatology, 2019, 15, 413-426.	8.0	99
5	Association of urinary metals levels with type 2 diabetes risk in coke oven workers. Environmental Pollution, 2016, 210, 1-8.	7.5	95
6	EpiGPU: exhaustive pairwise epistasis scans parallelized on consumer level graphics cards. Bioinformatics, 2011, 27, 1462-1465.	4.1	77
7	Exposure to the Chinese Famine in Childhood Increases Type 2 Diabetes Risk in Adults. Journal of Nutrition, 2016, 146, 2289-2295.	2.9	70
8	Biochar is superior to lime in improving acidic soil properties and fruit quality of Satsuma mandarin. Science of the Total Environment, 2020, 714, 136722.	8.0	56
9	High-throughput analysis of epistasis in genome-wide association studies with BiForce. Bioinformatics, 2012, 28, 1957-1964.	4.1	49
10	Bidirectional association between nonalcoholic fatty liver disease and type 2 diabetes in Chinese population: Evidence from the Dongfeng-Tongji cohort study. PLoS ONE, 2017, 12, e0174291.	2.5	48
11	<scp><i>Helicobacter pylori</i></scp> infection is associated with type 2 diabetes among a middle―and oldâ€age Chinese population. Diabetes/Metabolism Research and Reviews, 2016, 32, 95-101.	4.0	43
12	Association between exposure to arsenic, nickel, cadmium, selenium, and zinc and fasting blood glucose levels. Environmental Pollution, 2019, 255, 113325.	7.5	41
13	Using different anthropometric indices to assess prediction ability of type 2 diabetes in elderly population: a 5Âyear prospective study. BMC Geriatrics, 2018, 18, 218.	2.7	38
14	Long sleep duration and afternoon napping are associated with higher risk of incident diabetes in middle-aged and older Chinese: the Dongfeng-Tongji cohort study. Annals of Medicine, 2016, 48, 216-223.	3.8	34
15	Current Progress on MicroRNA-Based Gene Delivery in the Treatment of Osteoporosis and Osteoporotic Fracture. International Journal of Endocrinology, 2019, 2019, 1-17.	1.5	34
16	Genome-wide analysis of epistasis in body mass index using multiple human populations. European Journal of Human Genetics, 2012, 20, 857-862.	2.8	33
17	QTL modulating ear size and erectness in pigs. Animal Genetics, 2007, 38, 222-226.	1.7	30
18	Abundant local interactions in the 4p16.1 region suggest functional mechanisms underlying SLC2A9 associations with human serum uric acid. Human Molecular Genetics, 2014, 23, 5061-5068.	2.9	29

#	Article	IF	CITATIONS
19	Controlling false positives in the mapping of epistatic QTL. Heredity, 2010, 104, 401-409.	2.6	27
20	Mendelian randomization study of serum uric acid levels and diabetes risk: evidence from the Dongfeng-Tongji cohort. BMJ Open Diabetes Research and Care, 2020, 8, e000834.	2.8	26
21	Quantitative trait loci affecting eggshell traits in an F ₂ population. Animal Genetics, 2011, 42, 293-299.	1.7	25
22	BiForce Toolbox: powerful high-throughput computational analysis of gene-gene interactions in genome-wide association studies. Nucleic Acids Research, 2012, 40, W628-W632.	14.5	25
23	Differential regulation of two <i>FLNA</i> transcripts explains some of the phenotypic heterogeneity in the loss-of-function filaminopathies. Human Mutation, 2018, 39, 103-113.	2.5	24
24	Mapping quantitative trait loci regulating chicken body composition traits. Animal Genetics, 2009, 40, 952-954.	1.7	21
25	Serum bilirubin levels and risk of type 2 diabetes: results from two independent cohorts in middle-aged and elderly Chinese. Scientific Reports, 2017, 7, 41338.	3.3	20
26	Genotypic variability based association identifies novel non-additive loci DHCR7 and IRF4 in sero-negative rheumatoid arthritis. Scientific Reports, 2017, 7, 5261.	3.3	20
27	Victims of Chinese famine in early life have increased risk of metabolic syndrome in adulthood. Nutrition, 2018, 53, 20-25.	2.4	18
28	Recessive Spondylocarpotarsal Synostosis Syndrome Due to Compound Heterozygosity for Variants in MYH3. American Journal of Human Genetics, 2018, 102, 1115-1125.	6.2	18
29	Functional Urate-Associated Genetic Variants Influence Expression of lincRNAs LINC01229 and MAFTRR. Frontiers in Genetics, 2018, 9, 733.	2.3	18
30	Associations of plasma metal levels with type 2 diabetes and the mediating effects of microRNAs. Environmental Pollution, 2022, 292, 118452.	7.5	18
31	Major histocompatibility complex harbors widespread genotypic variability of non-additive risk of rheumatoid arthritis including epistasis. Scientific Reports, 2016, 6, 25014.	3.3	17
32	Metabolic syndrome is associated with hearing loss among a middle-aged and older Chinese population: a cross-sectional study. Annals of Medicine, 2018, 50, 587-595.	3.8	17
33	Genetic Risk, a Healthy Lifestyle, and Type 2 Diabetes: the Dongfeng-Tongji Cohort Study. Journal of Clinical Endocrinology and Metabolism, 2020, 105, 1242-1250.	3.6	17
34	Association between serum bilirubin levels and decline in estimated glomerular filtration rate among patients with type 2 diabetes. Journal of Diabetes and Its Complications, 2016, 30, 1255-1260.	2.3	16
35	Properties of Local Interactions and Their Potential Value in Complementing Genome-Wide Association Studies. PLoS ONE, 2013, 8, e71203.	2.5	15
36	Population genetic structure of Diaphorina citri Kuwayama (Hemiptera: Liviidae): host-driven genetic differentiation in China. Scientific Reports, 2018, 8, 1473.	3.3	15

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37	Serum calprotectin correlates with risk and disease severity of ankylosing spondylitis and its change during first month might predict favorable response to treatment. Modern Rheumatology, 2019, 29, 836-842.	1.8	15
38	Characterisation of Genome-Wide Association Epistasis Signals for Serum Uric Acid in Human Population Isolates. PLoS ONE, 2011, 6, e23836.	2.5	15
39	Serum bilirubin concentrations and incident coronary heart disease risk among patients with type 2 diabetes: the Dongfeng–Tongji cohort. Acta Diabetologica, 2017, 54, 257-264.	2.5	14
40	Association between resting heart rate and incident diabetes risk: a Mendelian randomization study. Acta Diabetologica, 2019, 56, 1037-1044.	2.5	12
41	Plasma metals and cancer incidence in patients with type 2 diabetes. Science of the Total Environment, 2021, 758, 143616.	8.0	11
42	A combined strategy for quantitative trait loci detection by genome-wide association. BMC Proceedings, 2009, 3, S6.	1.6	10
43	Serum alanine transaminase levels predict type 2 diabetes risk among a middle-aged and elderly Chinese population. Annals of Hepatology, 2019, 18, 298-303.	1.5	10
44	Prediction of 5-year risk of diabetes mellitus in relatively low risk middle-aged and elderly adults. Acta Diabetologica, 2020, 57, 63-70.	2.5	10
45	Mapping QTL in the porcine MHC region affecting fatness and growth traits in a Meishan/Large White composite population. Animal Genetics, 2011, 42, 83-85.	1.7	9
46	Genotypic variability-based genome-wide association study identifies non-additive loci HLA-C and IL12B for psoriasis. Journal of Human Genetics, 2018, 63, 289-296.	2.3	9
47	Development of a new scoring system to predict 5-year incident diabetes risk in middle-aged and older Chinese. Acta Diabetologica, 2018, 55, 13-19.	2.5	9
48	Immunochip Analyses of Epistasis in Rheumatoid Arthritis Confirm Multiple Interactions within MHC and Suggest Novel Non-MHC Epistatic Signals. Journal of Rheumatology, 2016, 43, 839-845.	2.0	8
49	Dyslexia associated functional variants in Europeans are not associated with dyslexia in Chinese. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2019, 180, 488-495.	1.7	8
50	High throughput analyses of epistasis for swine body dimensions and organ weights. Animal Genetics, 2011, 42, 15-21.	1.7	7
51	Trans-ancestral dissection of urate- and gout-associated major loci SLC2A9 and ABCG2 reveals primate-specific regulatory effects. Journal of Human Genetics, 2021, 66, 161-169.	2.3	6
52	Association between cancer antigen $19\hat{a} \in 9$ and diabetes risk: A prospective and Mendelian randomization study. Journal of Diabetes Investigation, 2020, 11, 585-593.	2.4	3
53	Pinyin Is an Effective Proxy for Early Screening for Mandarin-Speaking Children at Risk of Reading Disorders. Frontiers in Psychology, 2020, 11, 327.	2.1	3
54	The protective mechanism of action of plantamajoside on a rat model of acute spinal cord injury. Experimental and Therapeutic Medicine, 2021, 21, 378.	1.8	3

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#	Article	IF	CITATIONS
55	Reply to comment: Serum bilirubin concentrations, type 2 diabetes, and incident coronary heart disease. Acta Diabetologica, 2019, 56, 383-384.	2.5	2
56	257. Incorporating Genotypic Variability Mapping Enhances Discovery of Risk Loci for Rheumatoid Arthritis. Rheumatology, 2015, , .	1.9	0