

Lang Wu

List of Publications by Year in descending order

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43
papers

1,409
citations

394421

19
h-index

377865

34
g-index

43
all docs

43
docs citations

43
times ranked

3060
citing authors

#	ARTICLE	IF	CITATIONS
1	A transcriptome-wide association study identifies novel candidate susceptibility genes for prostate cancer risk. <i>International Journal of Cancer</i> , 2022, 150, 80-90.	5.1	9
2	Novel role of prostate cancer risk variant rs7247241 on <i>PPP1R14A</i> isoform transition through allelic TF binding and CpG methylation. <i>Human Molecular Genetics</i> , 2022, 31, 1610-1621.	2.9	5
3	Recommended Definitions of Aggressive Prostate Cancer for Etiologic Epidemiologic Research. <i>Journal of the National Cancer Institute</i> , 2021, 113, 727-734.	6.3	36
4	Associations Between Genetically Predicted Protein Levels and COVID-19 Severity. <i>Journal of Infectious Diseases</i> , 2021, 223, 19-22.	4.0	10
5	Red meat consumption, cooking mutagens, <i>NAT1</i> genotypes and pancreatic cancer risk in two ethnically diverse prospective cohorts. <i>International Journal of Cancer</i> , 2021, 149, 811-819.	5.1	12
6	Associations between Genetically Predicted Circulating Protein Concentrations and Endometrial Cancer Risk. <i>Cancers</i> , 2021, 13, 2088.	3.7	10
7	An integrative multiomics analysis identifies putative causal genes for COVID-19 severity. <i>Genetics in Medicine</i> , 2021, 23, 2076-2086.	2.4	25
8	InTACT: An adaptive and powerful framework for joint-tissue transcriptome-wide association studies. <i>Genetic Epidemiology</i> , 2021, 45, 848-859.	1.3	4
9	A transcriptome-wide association study identifies novel blood-based gene biomarker candidates for Alzheimer's disease risk. <i>Human Molecular Genetics</i> , 2021, 31, 289-299.	2.9	7
10	A transcriptome-wide association study of Alzheimer's disease using prediction models of relevant tissues identifies novel candidate susceptibility genes. <i>Genome Medicine</i> , 2021, 13, 141.	8.2	25
11	Associations Between Genetically Predicted Plasma N-Glycans and Prostate Cancer Risk: Analysis of Over 140,000 European Descendants. <i>Pharmacogenomics and Personalized Medicine</i> , 2021, Volume 14, 1211-1220.	0.7	1
12	Novel strategy for disease risk prediction incorporating predicted gene expression and DNA methylation data: a multi-phased study of prostate cancer. <i>Cancer Communications</i> , 2021, 41, 1387-1397.	9.2	6
13	Integrating Genome and Methylome Data to Identify Candidate DNA Methylation Biomarkers for Pancreatic Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 2079-2087.	2.5	10
14	Genetically Predicted Levels of DNA Methylation Biomarkers and Breast Cancer Risk: Data From 228,951 Women of European Descent. <i>Journal of the National Cancer Institute</i> , 2020, 112, 295-304.	6.3	35
15	Evaluation of associations between genetically predicted circulating protein biomarkers and breast cancer risk. <i>International Journal of Cancer</i> , 2020, 146, 2130-2138.	5.1	13
16	A Transcriptome-Wide Association Study Identifies Novel Candidate Susceptibility Genes for Pancreatic Cancer. <i>Journal of the National Cancer Institute</i> , 2020, 112, 1003-1012.	6.3	59
17	Comprehensive Analysis of RNA-Seq Gene Expression Profiling of Brain Transcriptomes Reveals Novel Genes, Regulators, and Pathways in Autism Spectrum Disorder. <i>Brain Sciences</i> , 2020, 10, 747.	2.3	45
18	An integrative multi-omics analysis to identify candidate DNA methylation biomarkers related to prostate cancer risk. <i>Nature Communications</i> , 2020, 11, 3905.	12.8	28

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19	Mendelian Randomization Analysis of n-6 Polyunsaturated Fatty Acid Levels and Pancreatic Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 2735-2739.	2.5	6
20	A Transcriptome-Wide Association Study Identifies Candidate Susceptibility Genes for Pancreatic Cancer Risk. <i>Cancer Research</i> , 2020, 80, 4346-4354.	0.9	28
21	Transcriptome-wide association study of breast cancer risk by estrogen receptor status. <i>Genetic Epidemiology</i> , 2020, 44, 442-468.	1.3	32
22	Associations between Genetically Predicted Blood Protein Biomarkers and Pancreatic Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2020, 29, 1501-1508.	2.5	18
23	Genetic Data from Nearly 63,000 Women of European Descent Predicts DNA Methylation Biomarkers and Epithelial Ovarian Cancer Risk. <i>Cancer Research</i> , 2019, 79, 505-517.	0.9	49
24	Transcriptome-Wide Association Study Identifies Susceptibility Loci and Genes for Age at Natural Menopause. <i>Reproductive Sciences</i> , 2019, 26, 496-502.	2.5	13
25	Analysis of Over 140,000 European Descendants Identifies Genetically Predicted Blood Protein Biomarkers Associated with Prostate Cancer Risk. <i>Cancer Research</i> , 2019, 79, 4592-4598.	0.9	16
26	Authors' response: Associations of obesity and circulating insulin and glucose with breast cancer risk. <i>International Journal of Epidemiology</i> , 2019, 48, 1016-1017.	1.9	1
27	Identification of Novel Susceptibility Loci and Genes for Prostate Cancer Risk: A Transcriptome-Wide Association Study in Over 140,000 European Descendants. <i>Cancer Research</i> , 2019, 79, 3192-3204.	0.9	43
28	Associations of obesity and circulating insulin and glucose with breast cancer risk: a Mendelian randomization analysis. <i>International Journal of Epidemiology</i> , 2019, 48, 795-806.	1.9	81
29	Physical Activity and Pancreatic Cancer Risk among Urban Chinese: Results from Two Prospective Cohort Studies. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2018, 27, 479-487.	2.5	16
30	Transcriptome-wide association studies accounting for colocalization using Egger regression. <i>Genetic Epidemiology</i> , 2018, 42, 418-433.	1.3	59
31	<i>CDKN2A</i> Germline Rare Coding Variants and Risk of Pancreatic Cancer in Minority Populations. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2018, 27, 1364-1370.	2.5	23
32	A Transcriptome-Wide Association Study Among 97,898 Women to Identify Candidate Susceptibility Genes for Epithelial Ovarian Cancer Risk. <i>Cancer Research</i> , 2018, 78, 5419-5430.	0.9	54
33	Is There Causal Relationship of Smoking and Alcohol Consumption with Bone Mineral Density? A Mendelian Randomization Study. <i>Calcified Tissue International</i> , 2018, 103, 546-553.	3.1	20
34	A transcriptome-wide association study of 229,000 women identifies new candidate susceptibility genes for breast cancer. <i>Nature Genetics</i> , 2018, 50, 968-978.	21.4	184
35	Genetic Evidence for the Association between Schizophrenia and Breast Cancer. <i>Journal of Psychiatry and Brain Science</i> , 2018, 3, .	0.5	10
36	Body mass index and persistent pain after breast cancer surgery: findings from the women's healthy eating and living study and a meta-analysis. <i>Oncotarget</i> , 2017, 8, 43332-43343.	1.8	20

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37	Dietary n-3 polyunsaturated fatty acids, fish consumption, and endometrial cancer risk: a meta-analysis of epidemiological studies. <i>Oncotarget</i> , 2017, 8, 91684-91693.	1.8	13
38	Different extent in decline of infant mortality by region and cause in Shenyang, China. <i>Scientific Reports</i> , 2016, 6, 24527.	3.3	10
39	Parity and thyroid cancer risk: a meta-analysis of epidemiological studies. <i>Cancer Medicine</i> , 2016, 5, 739-752.	2.8	29
40	Clopidogrel Resistance Increases Rate of Recurrent Stroke and Other Vascular Events in Chinese Population. <i>Journal of Stroke and Cerebrovascular Diseases</i> , 2016, 25, 1222-1228.	1.6	42
41	Breastfeeding and thyroid cancer risk in women: A dose-response meta-analysis of epidemiological studies. <i>Clinical Nutrition</i> , 2016, 35, 1039-1046.	5.0	15
42	Dose-response association of screen time-based sedentary behaviour in children and adolescents and depression: a meta-analysis of observational studies. <i>British Journal of Sports Medicine</i> , 2016, 50, 1252-1258.	6.7	231
43	Statin use and breast cancer survival and risk: a systematic review and meta-analysis. <i>Oncotarget</i> , 2015, 6, 42988-43004.	1.8	56