

Wenting Wu

List of Publications by Year in descending order

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Version: 2024-02-01

56
papers

3,027
citations

257450

24
h-index

175258

52
g-index

56
all docs

56
docs citations

56
times ranked

7225
citing authors

#	ARTICLE	IF	CITATIONS
1	Impact of Proinflammatory Cytokines on Alternative Splicing Patterns in Human Islets. <i>Diabetes</i> , 2022, 71, 116-127.	0.6	4
2	Genome-wide association study in almost 195,000 individuals identifies 50 previously unidentified genetic loci for eye color. <i>Science Advances</i> , 2021, 7, .	10.3	36
3	Common genetic polymorphisms contribute to the association between chronic lymphocytic leukaemia and non-melanoma skin cancer. <i>International Journal of Epidemiology</i> , 2021, 50, 1325-1334.	1.9	4
4	Deoxyhypusine synthase, an essential enzyme for hypusine biosynthesis, is required for proper exocrine pancreas development. <i>FASEB Journal</i> , 2021, 35, e21473.	0.5	13
5	Deoxyhypusine synthase promotes a pro-inflammatory macrophage phenotype. <i>Cell Metabolism</i> , 2021, 33, 1883-1893.e7.	16.2	24
6	Cutaneous nevi and internal cancer risk: Results from two large prospective cohorts of US women. <i>International Journal of Cancer</i> , 2020, 147, 14-20.	5.1	2
7	STAT5 promotes accessibility and is required for BATF-mediated plasticity at the Il9 locus. <i>Nature Communications</i> , 2020, 11, 4882.	12.8	29
8	Expression of SARS-CoV-2 Entry Factors in the Pancreas of Normal Organ Donors and Individuals with COVID-19. <i>Cell Metabolism</i> , 2020, 32, 1041-1051.e6.	16.2	135
9	Single-Cell Transcriptional Profiling of Mouse Islets Following Short-Term Obesogenic Dietary Intervention. <i>Metabolites</i> , 2020, 10, 513.	2.9	14
10	Bcl6 and Blimp1 reciprocally regulate ST2+ Treg cell development in the context of allergic airway inflammation. <i>Journal of Allergy and Clinical Immunology</i> , 2020, 146, 1121-1136.e9.	2.9	35
11	Genome-wide meta-analysis identifies eight new susceptibility loci for cutaneous squamous cell carcinoma. <i>Nature Communications</i> , 2020, 11, 820.	12.8	30
12	The Il9 CNS-25 Regulatory Element Controls Mast Cell and Basophil IL-9 Production. <i>Journal of Immunology</i> , 2019, 203, 1111-1121.	0.8	23
13	Hypusine biosynthesis in Î² cells links polyamine metabolism to facultative cellular proliferation to maintain glucose homeostasis. <i>Science Signaling</i> , 2019, 12, .	3.6	37
14	USP9X deubiquitinates ALDH1A3 and maintains mesenchymal identity in glioblastoma stem cells. <i>Journal of Clinical Investigation</i> , 2019, 129, 2043-2055.	8.2	45
15	Inverse Relationship between Vitiligo-Related Genes and Skin Cancer Risk. <i>Journal of Investigative Dermatology</i> , 2018, 138, 2072-2075.	0.7	20
16	Genome-wide association study in 176,678 Europeans reveals genetic loci for tanning response to sun exposure. <i>Nature Communications</i> , 2018, 9, 1684.	12.8	80
17	Association between genetic variation within vitamin D receptor DNA binding sites and risk of basal cell carcinoma. <i>International Journal of Cancer</i> , 2017, 140, 2085-2091.	5.1	11
18	Genetic Variants in WNT2B and BTRC Predict Melanoma Survival. <i>Journal of Investigative Dermatology</i> , 2017, 137, 1749-1756.	0.7	5

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19	Association study of genetic variation in <sc>DNA</sc> repair pathway genes and risk of basal cell carcinoma. International Journal of Cancer, 2017, 141, 952-957.	5.1	14
20	Genetic variants in the integrin signaling pathway genes predict cutaneous melanoma survival. International Journal of Cancer, 2017, 140, 1270-1279.	5.1	4
21	Genetic variants in the genes encoding rho GTPases and related regulators predict cutaneous melanoma-specific survival. International Journal of Cancer, 2017, 141, 721-730.	5.1	8
22	Phosphodiesterase type 5 inhibitors and risk of melanoma: A meta-analysis. Journal of the American Academy of Dermatology, 2017, 77, 480-488.e9.	1.2	13
23	Contribution of copy number variants to schizophrenia from a genome-wide study of 41,321 subjects. Nature Genetics, 2017, 49, 27-35.	21.4	838
24	Genetic variants of PDGF signaling pathway genes predict cutaneous melanoma survival. Oncotarget, 2017, 8, 74595-74606.	1.8	3
25	Two-stage genome-wide association study identifies a novel susceptibility locus associated with melanoma. Oncotarget, 2017, 8, 17586-17592.	1.8	61
26	Genetic variants in the vitamin <sc>D</sc> pathway genes <i><sc>VDBP</sc></i> and <i><sc>RXRA</sc></i> modulate cutaneous melanoma disease-specific survival. Pigment Cell and Melanoma Research, 2016, 29, 176-185.	3.3	19
27	Genetic variants in the PIWI-piRNA pathway gene <i>DCP1A</i> predict melanoma disease-specific survival. International Journal of Cancer, 2016, 139, 2730-2737.	5.1	21
28	Genome-wide association study identifies novel susceptibility loci for cutaneous squamous cell carcinoma. Nature Communications, 2016, 7, 12048.	12.8	117
29	Tissue-specific Co-expression of Long Non-coding and Coding RNAs Associated with Breast Cancer. Scientific Reports, 2016, 6, 32731.	3.3	35
30	Genome-wide association study identifies 14 novel risk alleles associated with basal cell carcinoma. Nature Communications, 2016, 7, 12510.	12.8	94
31	Longitudinal genome-wide methylation study of Roux-en-Y gastric bypass patients reveals novel CpG sites associated with essential hypertension. BMC Medical Genomics, 2016, 9, 20.	1.5	18
32	Associations between smoking behavior-related alleles and the risk of melanoma. Oncotarget, 2016, 7, 47366-47375.	1.8	15
33	<i>SMAD7</i> polymorphisms and colorectal cancer risk: a meta-analysis of case-control studies. Oncotarget, 2016, 7, 75561-75570.	1.8	17
34	Association of TERT Polymorphisms with Clinical Outcome of Non-Small Cell Lung Cancer Patients. PLoS ONE, 2015, 10, e0129232.	2.5	11
35	Genetic Variants of the MDM2 Gene Are Predictive of Treatment-Related Toxicities and Overall Survival in Patients With Advanced NSCLC. Clinical Lung Cancer, 2015, 16, e37-e53.	2.6	13
36	c-Myc-miR-29-REV3L signalling pathway drives the acquisition of temozolomide resistance in glioblastoma. Brain, 2015, 138, 3654-3672.	7.6	55

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37	MicroRNA-377 inhibited proliferation and invasion of human glioblastoma cells by directly targeting specificity protein 1. <i>Neuro-Oncology</i> , 2014, 16, 1510-1522.	1.2	59
38	Association Between CASP8 and CASP10 Polymorphisms and Toxicity Outcomes With Platinum-Based Chemotherapy in Chinese Patients With Non-small Cell Lung Cancer. <i>Oncologist</i> , 2012, 17, 1551-1561.	3.7	20
39	Contactin-1 (CNTN-1) Overexpression is Correlated with Advanced Clinical Stage and Lymph Node Metastasis in Oesophageal Squamous Cell Carcinomas. <i>Japanese Journal of Clinical Oncology</i> , 2012, 42, 612-618.	1.3	24
40	Association of CASP7 Polymorphisms and Survival of Patients With Non-small Cell Lung Cancer With Platinum-Based Chemotherapy Treatment. <i>Chest</i> , 2012, 142, 680-689.	0.8	8
41	Whole-Genome Sequencing in Autism Identifies Hot Spots for De Novo Germline Mutation. <i>Cell</i> , 2012, 151, 1431-1442.	28.9	501
42	<i>hsa-miR-196a2</i> Functional SNP is Associated With Severe Toxicity After Platinum-Based Chemotherapy of Advanced Nonsmall Cell Lung Cancer Patients in a Chinese Population. <i>Journal of Clinical Laboratory Analysis</i> , 2012, 26, 441-446.	2.1	24
43	Effect of Polymorphisms in XPD on Clinical Outcomes of Platinum-Based Chemotherapy for Chinese Non-Small Cell Lung Cancer Patients. <i>PLoS ONE</i> , 2012, 7, e33200.	2.5	32
44	Matrix metalloproteinase-2 polymorphisms and clinical outcome of Chinese patients with nonsmall cell lung cancer treated with first-line, platinum-based chemotherapy. <i>Cancer</i> , 2012, 118, 3587-3598.	4.1	12
45	Association of <i>CASP3</i> polymorphism with hematologic toxicity in patients with advanced non-small cell lung carcinoma treated with platinum-based chemotherapy. <i>Cancer Science</i> , 2012, 103, 1451-1459.	3.9	25
46	Association of ABCC2 polymorphisms with platinum-based chemotherapy response and severe toxicity in non-small cell lung cancer patients. <i>Lung Cancer</i> , 2011, 72, 238-243.	2.0	47
47	Polymorphisms of the vascular endothelial growth factor A gene and susceptibility to sporadic brain arteriovenous malformation in a Chinese population. <i>Journal of Clinical Neuroscience</i> , 2011, 18, 549-553.	1.5	26
48	Possible association between genetic variants in the H2AFX promoter region and risk of adult glioma in a Chinese Han population. <i>Journal of Neuro-Oncology</i> , 2011, 105, 211-218.	2.9	12
49	A variant in the CHEK2 promoter at a methylation site relieves transcriptional repression and confers reduced risk of lung cancer. <i>Carcinogenesis</i> , 2010, 31, 1251-1258.	2.8	26
50	Analysis of specialized DNA polymerases expression in human gliomas: association with prognostic significance. <i>Neuro-Oncology</i> , 2010, 12, 679-686.	1.2	75
51	Functional characterization of a promoter polymorphism in APE1/Ref-1 that contributes to reduced lung cancer susceptibility. <i>FASEB Journal</i> , 2009, 23, 3459-3469.	0.5	65
52	REV3L confers chemoresistance to cisplatin in human gliomas: The potential of its RNAi for synergistic therapy. <i>Neuro-Oncology</i> , 2009, 11, 790-802.	1.2	65
53	Association of <i>XPD</i> Polymorphisms with Severe Toxicity in Non-small Cell Lung Cancer Patients in a Chinese Population. <i>Clinical Cancer Research</i> , 2009, 15, 3889-3895.	7.0	43
54	Genetic variants in GTF2H1 and risk of lung cancer: A case-control analysis in a Chinese population. <i>Lung Cancer</i> , 2009, 63, 180-186.	2.0	10

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55	Association of polymorphisms in one-carbon metabolizing genes and lung cancer risk: a case-control study in Chinese population. <i>Lung Cancer</i> , 2008, 61, 21-29.	2.0	44
56	Methyl-CpG binding domain 1 gene polymorphisms and lung cancer risk in a Chinese population. <i>Biomarkers</i> , 2008, 13, 607-617.	1.9	6