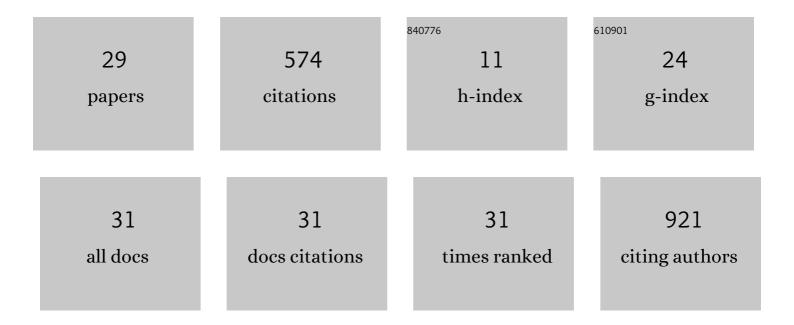
Xuemei Zhang

List of Publications by Year in descending order

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XHEMEL THANC

#	Article	IF	CITATIONS
1	Right Ventricular Strain Is Associated with Increased Length of Stay after Tetralogy of Fallot Repair. Journal of Cardiovascular Imaging, 2022, 29, 50-58.	0.7	1
2	Functional genetic variants in complement component 7 confer susceptibility to gastric cancer. PeerJ, 2022, 10, e12816.	2.0	2
3	TLR4 regulatory region variants reduce the susceptibility of small-cell lung cancer in Chinese population. European Journal of Cancer Prevention, 2022, 31, 363-368.	1.3	2
4	Genetic variants in the regulation region of TLR4 reduce the gastric cancer susceptibility. Gene, 2021, 767, 145181.	2.2	11
5	TLR4 promoter rs1927914 variant contributes to the susceptibility of esophageal squamous cell carcinoma in the Chinese population. PeerJ, 2021, 9, e10754.	2.0	3
6	XAB2 TagSNP Is Associated with the Risk of Gastric Cancer in Chinese Population: A Case–Control Study. International Journal of Environmental Research and Public Health, 2021, 18, 1494.	2.6	1
7	Methylation Regulation of TLR3 on Immune Parameters in Lung Adenocarcinoma. Frontiers in Oncology, 2021, 11, 620200.	2.8	3
8	Phlorizin from sweet tea inhibits the progress of esophageal cancer by antagonizing the JAK2/STAT3 signaling pathway. Oncology Reports, 2021, 46, .	2.6	13
9	Integration of transcriptomics and metabolomics reveals anlotinib-induced cytotoxicity in colon cancer cells. Gene, 2021, 786, 145625.	2.2	8
10	XPF -673C>T variation is associated with the susceptibility to breast cancer. Cancer Epidemiology, 2021, 74, 102007.	1.9	0
11	Impact of Genetic Variation in <i>TLR4</i> 3′UTR on NSCLC Genetic Susceptibility. Journal of Oncology, 2020, 2020, 1-7.	1.3	8
12	CSB affected on the sensitivity of lung cancer cells to platinum-based drugs through the global decrease of let-7 and miR-29. BMC Cancer, 2019, 19, 948.	2.6	11
13	TNFSF15 promoter polymorphisms increase the susceptibility to small cell lung cancer: a case-control study. BMC Medical Genetics, 2019, 20, 29.	2.1	5
14	Exome-wide analyses identify low-frequency variant in CYP26B1 and additional coding variants associated with esophageal squamous cell carcinoma. Nature Genetics, 2018, 50, 338-343.	21.4	75
15	Exome-wide analysis identifies three low-frequency missense variants associated with pancreatic cancer risk in Chinese populations. Nature Communications, 2018, 9, 3688.	12.8	32
16	A common CD55 rs2564978 variant is associated with the susceptibility of non-small cell lung cancer. Oncotarget, 2017, 8, 6216-6221.	1.8	14
17	Genetic Polymorphisms in the Apoptosis-Associated Gene CASP3 and the Risk of Lung Cancer in Chinese Population. PLoS ONE, 2016, 11, e0164358.	2.5	13
18	Complement Receptor 1 Genetic Variants Contribute to the Susceptibility to Gastric Cancer in Chinese Population. Journal of Cancer, 2015, 6, 525-530.	2.5	11

XUEMEI ZHANG

#	Article	IF	CITATIONS
19	Tag SNPs of CFI contributed to the susceptibility for non-small cell lung cancer in Chinese population. Tumor Biology, 2015, 36, 1955-1962.	1.8	2
20	Two Novel Variants on 13q22.1 Are Associated with Risk of Esophageal Squamous Cell Carcinoma. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1774-1780.	2.5	12
21	XAB2 tagSNPs contribute to non-small cell lung cancer susceptibility in Chinese population. BMC Cancer, 2015, 15, 560.	2.6	11
22	The tag SNP rs10746463 in decay-accelerating factor is associated with the susceptibility to gastric cancer. Molecular Immunology, 2015, 63, 473-478.	2.2	10
23	XPF-673C>T Polymorphism Effect on the Susceptibility to Esophageal Cancer in Chinese Population. PLoS ONE, 2014, 9, e94136.	2.5	7
24	Tag SNPs in complement receptor-1 contribute to the susceptibility to non-small cell lung cancer. Molecular Cancer, 2014, 13, 56.	19.2	14
25	A regulatory variant in CYP2E1 affects the risk of lung squamous cell carcinoma. Tumor Biology, 2014, 35, 455-462.	1.8	11
26	Functional Genetic Variants of TNFSF15 and Their Association with Gastric Adenocarcinoma: A Case-Control Study. PLoS ONE, 2014, 9, e108321.	2.5	14
27	Cigarette smoking strongly modifies the association of complement factor H variant and the risk of lung cancer. Cancer Epidemiology, 2012, 36, e111-e115.	1.9	11
28	Genetic polymorphisms in cell cycle regulatory genesMDM2 andTP53 are associated with susceptibility to lung cancer. Human Mutation, 2006, 27, 110-117.	2.5	142
29	Polymorphisms in DNA base excision repair genes ADPRT and XRCC1 and risk of lung cancer. Cancer Research 2005 65 722-6	0.9	127