

# Xuemei Zhang

## List of Publications by Year in descending order

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29  
papers

574  
citations

840776

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610901

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31  
docs citations

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times ranked

921  
citing authors

#	ARTICLE	IF	CITATIONS
1	Right Ventricular Strain Is Associated with Increased Length of Stay after Tetralogy of Fallot Repair. <i>Journal of Cardiovascular Imaging</i> , 2022, 29, 50-58.	0.7	1
2	Functional genetic variants in complement component 7 confer susceptibility to gastric cancer. <i>PeerJ</i> , 2022, 10, e12816.	2.0	2
3	TLR4 regulatory region variants reduce the susceptibility of small-cell lung cancer in Chinese population. <i>European Journal of Cancer Prevention</i> , 2022, 31, 363-368.	1.3	2
4	Genetic variants in the regulation region of TLR4 reduce the gastric cancer susceptibility. <i>Gene</i> , 2021, 767, 145181.	2.2	11
5	TLR4 promoter rs1927914 variant contributes to the susceptibility of esophageal squamous cell carcinoma in the Chinese population. <i>PeerJ</i> , 2021, 9, e10754.	2.0	3
6	XAB2 TagSNP Is Associated with the Risk of Gastric Cancer in Chinese Population: A Case-€“Control Study. <i>International Journal of Environmental Research and Public Health</i> , 2021, 18, 1494.	2.6	1
7	Methylation Regulation of TLR3 on Immune Parameters in Lung Adenocarcinoma. <i>Frontiers in Oncology</i> , 2021, 11, 620200.	2.8	3
8	Phlorizin from sweet tea inhibits the progress of esophageal cancer by antagonizing the JAK2/STAT3 signaling pathway. <i>Oncology Reports</i> , 2021, 46, .	2.6	13
9	Integration of transcriptomics and metabolomics reveals anlotinib-induced cytotoxicity in colon cancer cells. <i>Gene</i> , 2021, 786, 145625.	2.2	8
10	XPF -673C&gt;T variation is associated with the susceptibility to breast cancer. <i>Cancer Epidemiology</i> , 2021, 74, 102007.	1.9	0
11	Impact of Genetic Variation in <i>TLR4</i> 3â€²UTR on NSCLC Genetic Susceptibility. <i>Journal of Oncology</i> , 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. <i>BMC Cancer</i> , 2019, 19, 948.	2.6	11
13	TNFSF15 promoter polymorphisms increase the susceptibility to small cell lung cancer: a case-control study. <i>BMC Medical Genetics</i> , 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. <i>Nature Genetics</i> , 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. <i>Nature Communications</i> , 2018, 9, 3688.	12.8	32
16	A common CD55 rs2564978 variant is associated with the susceptibility of non-small cell lung cancer. <i>Oncotarget</i> , 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. <i>PLoS ONE</i> , 2016, 11, e0164358.	2.5	13
18	Complement Receptor 1 Genetic Variants Contribute to the Susceptibility to Gastric Cancer in Chinese Population. <i>Journal of Cancer</i> , 2015, 6, 525-530.	2.5	11

#	ARTICLE	IF	CITATIONS
19	Tag SNPs of CFI contributed to the susceptibility for non-small cell lung cancer in Chinese population. <i>Tumor Biology</i> , 2015, 36, 1955-1962.	1.8	2
20	Two Novel Variants on 13q22.1 Are Associated with Risk of Esophageal Squamous Cell Carcinoma. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 1774-1780.	2.5	12
21	XAB2 tagSNPs contribute to non-small cell lung cancer susceptibility in Chinese population. <i>BMC Cancer</i> , 2015, 15, 560.	2.6	11
22	The tag SNP rs10746463 in decay-accelerating factor is associated with the susceptibility to gastric cancer. <i>Molecular Immunology</i> , 2015, 63, 473-478.	2.2	10
23	XPF-673C>T Polymorphism Effect on the Susceptibility to Esophageal Cancer in Chinese Population. <i>PLoS ONE</i> , 2014, 9, e94136.	2.5	7
24	Tag SNPs in complement receptor-1 contribute to the susceptibility to non-small cell lung cancer. <i>Molecular Cancer</i> , 2014, 13, 56.	19.2	14
25	A regulatory variant in CYP2E1 affects the risk of lung squamous cell carcinoma. <i>Tumor Biology</i> , 2014, 35, 455-462.	1.8	11
26	Functional Genetic Variants of TNFSF15 and Their Association with Gastric Adenocarcinoma: A Case-Control Study. <i>PLoS ONE</i> , 2014, 9, e108321.	2.5	14
27	Cigarette smoking strongly modifies the association of complement factor H variant and the risk of lung cancer. <i>Cancer Epidemiology</i> , 2012, 36, e111-e115.	1.9	11
28	Genetic polymorphisms in cell cycle regulatory genes MDM2 and TP53 are associated with susceptibility to lung cancer. <i>Human Mutation</i> , 2006, 27, 110-117.	2.5	142
29	Polymorphisms in DNA base excision repair genes ADPRT and XRCC1 and risk of lung cancer. <i>Cancer Research</i> , 2005, 65, 722-6.	0.9	127