

# Ying Chen

## List of Publications by Year in descending order

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Version: 2024-02-01

23  
papers

915  
citations

759233

12  
h-index

713466

21  
g-index

23  
all docs

23  
docs citations

23  
times ranked

2515  
citing authors

#	ARTICLE	IF	CITATIONS
1	Serum cholinesterase may independently predict prognosis in non-small-cell lung cancer. <i>BMC Cancer</i> , 2022, 22, 93.	2.6	3
2	Multiple small pneumatoceles as a complication of a SARS-CoV-2 infection in a child from Myanmar. <i>Journal of Travel Medicine</i> , 2022, , .	3.0	1
3	A Multi-Omics Study of Familial Lung Cancer: Microbiome and Host Gene Expression Patterns. <i>Frontiers in Immunology</i> , 2022, 13, 827953.	4.8	7
4	Large pulmonary cavity in COVID-19 cured patient case report. <i>Annals of Palliative Medicine</i> , 2021, 10, 5786-5791.	1.2	22
5	Dysregulation of ferroptosis may involve in the development of non-small cell lung cancer in Xuanwei area. <i>Journal of Cellular and Molecular Medicine</i> , 2021, 25, 2872-2884.	3.6	22
6	Effects of lipid emulsions on the formation of <i>Escherichia coli</i> – <i>Candida albicans</i> mixed-species biofilms on PVC. <i>Scientific Reports</i> , 2021, 11, 16929.	3.3	2
7	Household air pollution from, and fuel efficiency of, different coal types following local cooking practices in Xuanwei, China. <i>Environmental Pollution</i> , 2021, 290, 117949.	7.5	1
8	The effects of TGF- $\beta$ 1 on staphylococcus epidermidis biofilm formation in a tree shrew biomaterial-centered infection model. <i>Annals of Translational Medicine</i> , 2021, 9, 57-57.	1.7	0
9	ERAP2 Is Associated With Immune Infiltration and Predicts Favorable Prognosis in SqCLC. <i>Frontiers in Immunology</i> , 2021, 12, 788985.	4.8	10
10	Tuberculosis infection and lung adenocarcinoma: Mendelian randomization and pathway analysis of genome-wide association study data from never-smoking Asian women. <i>Genomics</i> , 2020, 112, 1223-1232.	2.9	15
11	Dynamic serum alkaline phosphatase is an indicator of overall survival in pancreatic cancer. <i>BMC Cancer</i> , 2019, 19, 785.	2.6	20
12	MUC16 in non-small cell lung cancer patients affected by familial lung cancer and indoor air pollution: clinical characteristics and cell behaviors. <i>Translational Lung Cancer Research</i> , 2019, 8, 476-488.	2.8	9
13	Lung cancer family history and exposure to occupational/domestic coal combustion contribute to variations in clinicopathologic features and gene fusion patterns in non-small cell lung cancer. <i>Thoracic Cancer</i> , 2019, 10, 695-707.	1.9	7
14	&lt;p&gt;Nomogram For The Prediction Of Malignancy In Small (8&–20 mm) Indeterminate Solid Solitary Pulmonary Nodules In Chinese Populations&lt;/p&gt;. <i>Cancer Management and Research</i> , 2019, Volume 11, 9439-9448.	1.9	19
15	Clinic-pathologic features and gene fusion pattern of ALK and ROS1 in non-small cell lung cancer show association with household coal combustion. <i>Translational Cancer Research</i> , 2019, 8, 2164-2174.	1.0	2
16	Characterization of germline mutations in familial lung cancer from the Chinese population. <i>Gene</i> , 2018, 641, 94-104.	2.2	21
17	Characteristics of Familial Lung Cancer in Yunnan-Guizhou Plateau of China. <i>Frontiers in Oncology</i> , 2018, 8, 637.	2.8	8
18	Association between GWAS-identified lung adenocarcinoma susceptibility loci and EGFR mutations in never-smoking Asian women, and comparison with findings from Western populations. <i>Human Molecular Genetics</i> , 2016, 26, ddw414.	2.9	50

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19	Meta-analysis of genome-wide association studies identifies multiple lung cancer susceptibility loci in never-smoking Asian women. <i>Human Molecular Genetics</i> , 2016, 25, 620-629.	2.9	50
20	Analysis of Heritability and Shared Heritability Based on Genome-Wide Association Studies for Thirteen Cancer Types. <i>Journal of the National Cancer Institute</i> , 2015, 107, djv279.	6.3	152
21	Study on the Structure of <i>Candida Albicans</i> – <i>Staphylococcus Epidermidis</i> Mixed Species Biofilm on Polyvinyl Chloride Biomaterial. <i>Cell Biochemistry and Biophysics</i> , 2015, 73, 461-468.	1.8	12
22	Influence of common genetic variation on lung cancer risk: meta-analysis of 14 900 cases and 29 485 controls. <i>Human Molecular Genetics</i> , 2012, 21, 4980-4995.	2.9	196
23	Genome-wide association analysis identifies new lung cancer susceptibility loci in never-smoking women in Asia. <i>Nature Genetics</i> , 2012, 44, 1330-1335.	21.4	286