

# Wen Tan

## List of Publications by Year in descending order

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Version: 2024-02-01

50  
papers

2,994  
citations

236925

25  
h-index

182427

51  
g-index

51  
all docs

51  
docs citations

51  
times ranked

5883  
citing authors

#	ARTICLE	IF	CITATIONS
1	Multi-omic characterization of genome-wide abnormal DNA methylation reveals diagnostic and prognostic markers for esophageal squamous-cell carcinoma. <i>Signal Transduction and Targeted Therapy</i> , 2022, 7, 53.	17.1	22
2	Metformin inhibits pancreatic cancer metastasis caused by SMAD4 deficiency and consequent HNF4G upregulation. <i>Protein and Cell</i> , 2021, 12, 128-144.	11.0	41
3	Interleukin-33-Dependent Accumulation of Regulatory T Cells Mediates Pulmonary Epithelial Regeneration During Acute Respiratory Distress Syndrome. <i>Frontiers in Immunology</i> , 2021, 12, 653803.	4.8	9
4	VAV2 is required for DNA repair and implicated in cancer radiotherapy resistance. <i>Signal Transduction and Targeted Therapy</i> , 2021, 6, 322.	17.1	14
5	Dissecting esophageal squamous-cell carcinoma ecosystem by single-cell transcriptomic analysis. <i>Nature Communications</i> , 2021, 12, 5291.	12.8	98
6	CstF64-Induced Shortening of the 3' UTR Promotes Esophageal Squamous Cell Carcinoma Progression by Disrupting ceRNA Cross-talk with ZFP36L2. <i>Cancer Research</i> , 2021, 81, 5638-5651.	0.9	13
7	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
8	Genetic risk of extranodal natural killer T-cell lymphoma: a genome-wide association study in multiple populations. <i>Lancet Oncology</i> , 2020, 21, 306-316.	10.7	49
9	Genomic and transcriptomic alterations associated with drug vulnerabilities and prognosis in adenocarcinoma at the gastroesophageal junction. <i>Nature Communications</i> , 2020, 11, 6091.	12.8	21
10	Serum piRNA-54265 is a New Biomarker for early detection and clinical surveillance of Human Colorectal Cancer. <i>Theranostics</i> , 2020, 10, 8468-8478.	10.0	58
11	Metabolic remodeling by TIGAR overexpression is a therapeutic target in esophageal squamous-cell carcinoma. <i>Theranostics</i> , 2020, 10, 3488-3502.	10.0	27
12	Identification of risk loci and a polygenic risk score for lung cancer: a large-scale prospective cohort study in Chinese populations. <i>Lancet Respiratory Medicine</i> , 2019, 7, 881-891.	10.7	167
13	Genome landscapes of rectal cancer before and after preoperative chemoradiotherapy. <i>Theranostics</i> , 2019, 9, 6856-6866.	10.0	27
14	Excessive miR-25-3p maturation via N6-methyladenosine stimulated by cigarette smoke promotes pancreatic cancer progression. <i>Nature Communications</i> , 2019, 10, 1858.	12.8	242
15	Regulatory T cells promote pulmonary repair by modulating T helper cell immune responses in lipopolysaccharide-induced acute respiratory distress syndrome. <i>Immunology</i> , 2019, 157, 151-162.	4.4	26
16	PIWI-interacting RNA-36712 restrains breast cancer progression and chemoresistance by interaction with SEPWI pseudogene SEPWI1P RNA. <i>Molecular Cancer</i> , 2019, 18, 9.	19.2	139
17	Associations of Genetic Variations in Mismatch Repair Genes MSH3 and PMS1 with Acute Adverse Events and Survival in Patients with Rectal Cancer Receiving Postoperative Chemoradiotherapy. <i>Cancer Research and Treatment</i> , 2019, 51, 1198-1206.	3.0	10
18	Associations of Genetic Variations in MicroRNA Seed Regions With Acute Adverse Events and Survival in Patients With Rectal Cancer Receiving Postoperative Chemoradiation Therapy. <i>International Journal of Radiation Oncology Biology Physics</i> , 2018, 100, 1026-1033.	0.8	3

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19	Genetic variant repressing ADH1A expression confers susceptibility to esophageal squamous-cell carcinoma. <i>Cancer Letters</i> , 2018, 421, 43-50.	7.2	16
20	Functional role of PLCE1 intronic insertion variant associated with susceptibility to esophageal squamous cell carcinoma. <i>Carcinogenesis</i> , 2018, 39, 191-201.	2.8	5
21	Remote Ischemic Preconditioning has a Cardioprotective Effect in Children in the Early Postoperative Phase: A Meta-Analysis of Randomized Controlled Trials. <i>Pediatric Cardiology</i> , 2018, 39, 617-626.	1.3	11
22	Protectin DX Exhibits Protective Effects in Mouse Model of Lipopolysaccharide-Induced Acute Lung Injury. <i>Chinese Medical Journal</i> , 2018, 131, 1167-1173.	2.3	12
23	PIWI-interacting RNA-54265 is oncogenic and a potential therapeutic target in colorectal adenocarcinoma. <i>Theranostics</i> , 2018, 8, 5213-5230.	10.0	115
24	CCGD-ESCC: A Comprehensive Database for Genetic Variants Associated with Esophageal Squamous Cell Carcinoma in Chinese Population. <i>Genomics, Proteomics and Bioinformatics</i> , 2018, 16, 262-268.	6.9	17
25	Functional role of BTB and CNC Homology 1 gene in pancreatic cancer and its association with survival in patients treated with gemcitabine. <i>Theranostics</i> , 2018, 8, 3366-3379.	10.0	19
26	Integrative analysis of gene expression profiles reveals specific signaling pathways associated with pancreatic duct adenocarcinoma. <i>Cancer Communications</i> , 2018, 38, 1-12.	9.2	14
27	Solute Carrier Family 39 Member 6 Gene Promotes Aggressiveness of Esophageal Carcinoma Cells by Increasing Intracellular Levels of Zinc, Activating Phosphatidylinositol 3-Kinase Signaling, and Up-regulating Genes That Regulate Metastasis. <i>Gastroenterology</i> , 2017, 152, 1985-1997.e12.	1.3	40
28	Genomic analysis of oesophageal squamous-cell carcinoma identifies alcohol drinking-related mutation signature and genomic alterations. <i>Nature Communications</i> , 2017, 8, 15290.	12.8	195
29	BRCA1-Associated Protein Increases Invasiveness of Esophageal Squamous Cell Carcinoma. <i>Gastroenterology</i> , 2017, 153, 1304-1319.e5.	1.3	23
30	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
31	Pancreatic cancer risk variant in LINC00673 creates a miR-1231 binding site and interferes with PTPN11 degradation. <i>Nature Genetics</i> , 2016, 48, 747-757.	21.4	237
32	Genetic risk of extranodal natural killer T-cell lymphoma: a genome-wide association study. <i>Lancet Oncology</i> , 2016, 17, 1240-1247.	10.7	84
33	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
34	Genetic variants associated with longer telomere length are associated with increased lung cancer risk among never-smoking women in Asia: a report from the female lung cancer consortium in Asia. <i>International Journal of Cancer</i> , 2015, 137, 311-319.	5.1	72
35	Relatively Small Contribution of Methylation and Genomic Copy Number Aberration to the Aberrant Expression of Inflammation-Related Genes in HBV-Related Hepatocellular Carcinoma. <i>PLoS ONE</i> , 2015, 10, e0126836.	2.5	1
36	Pulmonary expression of <i>CYP2A13</i> and <i>ABCB1</i> is regulated by <i>FOXA2</i> , and their genetic interaction is associated with lung cancer. <i>FASEB Journal</i> , 2015, 29, 1986-1998.	0.5	15

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37	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
38	Associations of ATM Polymorphisms With Survival in Advanced Esophageal Squamous Cell Carcinoma Patients Receiving Radiation Therapy. <i>International Journal of Radiation Oncology Biology Physics</i> , 2015, 93, 181-189.	0.8	4
39	Imputation and subset-based association analysis across different cancer types identifies multiple independent risk loci in the TERT-CLPTM1L region on chromosome 5p15.33. <i>Human Molecular Genetics</i> , 2014, 23, 6616-6633.	2.9	90
40	Joint analysis of three genome-wide association studies of esophageal squamous cell carcinoma in Chinese populations. <i>Nature Genetics</i> , 2014, 46, 1001-1006.	21.4	148
41	Genome-wide association study identifies three susceptibility loci for laryngeal squamous cell carcinoma in the Chinese population. <i>Nature Genetics</i> , 2014, 46, 1110-1114.	21.4	57
42	Genome-wide association study identifies common variants in SLC39A6 associated with length of survival in esophageal squamous-cell carcinoma. <i>Nature Genetics</i> , 2013, 45, 632-638.	21.4	97
43	A Genome Wide Association Study Identifies Common Variants Associated with Lipid Levels in the Chinese Population. <i>PLoS ONE</i> , 2013, 8, e82420.	2.5	57
44	Genome-wide association study identifies five loci associated with susceptibility to pancreatic cancer in Chinese populations. <i>Nature Genetics</i> , 2012, 44, 62-66.	21.4	164
45	Functional XPF polymorphisms associated with lung cancer susceptibility in a Chinese population. <i>Frontiers of Medicine in China</i> , 2010, 4, 82-89.	0.1	4
46	Reply to "A promoter polymorphism in the CASP8 gene is not associated with cancer risk". <i>Nature Genetics</i> , 2008, 40, 260-261.	21.4	14
47	Associations of functional polymorphisms in cyclooxygenase-2 and platelet 12-lipoxygenase with risk of occurrence and advanced disease status of colorectal cancer. <i>Carcinogenesis</i> , 2007, 28, 1197-1201.	2.8	89
48	Significant increase in risk of gastroesophageal cancer is associated with interaction between promoter polymorphisms in thymidylate synthase and serum folate status. <i>Carcinogenesis</i> , 2005, 26, 1430-1435.	2.8	37
49	Frequency of CYP2A6 gene deletion and its relation to risk of lung and esophageal cancer in the Chinese population. <i>International Journal of Cancer</i> , 2001, 95, 96-101.	5.1	115
50	CYP 1A1 polymorphism and risk of lung cancer in relation to tobacco smoking: a case-control study in China. <i>Carcinogenesis</i> , 2001, 22, 11-16.	2.8	131