

# Yifeng Zhou

## List of Publications by Year in descending order

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63  
papers

3,673  
citations

186265

28  
h-index

133252

59  
g-index

67  
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67  
docs citations

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times ranked

5891  
citing authors

#	ARTICLE	IF	CITATIONS
1	A micropeptide XBP1SBM encoded by lncRNA promotes angiogenesis and metastasis of TNBC via XBP1s pathway. <i>Oncogene</i> , 2022, 41, 2163-2172.	5.9	15
2	A polygenic risk score for nasopharyngeal carcinoma shows potential for risk stratification and personalized screening. <i>Nature Communications</i> , 2022, 13, 1966.	12.8	19
3	Long non-coding RNA LSAMP-1 is down-regulated in non-small cell lung cancer and predicts a poor prognosis. <i>Cancer Cell International</i> , 2022, 22, 181.	4.1	6
4	A peptide CORO1C-47aa encoded by the circular noncoding RNA circ-0000437 functions as a negative regulator in endometrium tumor angiogenesis. <i>Journal of Biological Chemistry</i> , 2021, 297, 101182.	3.4	23
5	Micropeptide <sc>CIP</sc> 2A&€•<sc>BP</sc> encoded by <sc>LINC</sc> 00665 inhibits triple&€negative breast cancer progression. <i>EMBO Journal</i> , 2020, 39, e102190.	7.8	138
6	LncRNA-encoded polypeptide ASRPS inhibits triple-negative breast cancer angiogenesis. <i>Journal of Experimental Medicine</i> , 2020, 217, .	8.5	136
7	Upregulation of long noncoding RNA RAB11B-AS1 promotes tumor metastasis and predicts poor prognosis in lung cancer. <i>Annals of Translational Medicine</i> , 2020, 8, 582-582.	1.7	9
8	A Novel Micropeptide Encoded by Y-Linked LINC00278 Links Cigarette Smoking and AR Signaling in Male Esophageal Squamous Cell Carcinoma. <i>Cancer Research</i> , 2020, 80, 2790-2803.	0.9	91
9	Long noncoding RNA PANDAR inhibits the development of lung cancer by regulating autophagy and apoptosis pathways. <i>Journal of Cancer</i> , 2020, 11, 4783-4790.	2.5	14
10	Down expression of lnc-BMP1-1 decreases that of Caveolin-1 is associated with the lung cancer susceptibility and cigarette smoking history. <i>Aging</i> , 2020, 12, 462-480.	3.1	6
11	Analysis of Survival-Related lncRNA Landscape Identifies A Role for LINC01537 in Energy Metabolism and Lung Cancer Progression. <i>International Journal of Molecular Sciences</i> , 2019, 20, 3713.	4.1	31
12	X chromosome&€linked long noncoding RNA <i>Lnc&€XLEC1</i> regulates <i>c&€Myc</i>&€dependent cell growth by collaborating with MBP&€1 in endometrial cancer. <i>International Journal of Cancer</i> , 2019, 145, 927-940.	5.1	12
13	Exosomal FMR1-AS1 facilitates maintaining cancer stem-like cell dynamic equilibrium via TLR7/NF&€B/c-Myc signaling in female esophageal carcinoma. <i>Molecular Cancer</i> , 2019, 18, 22.	19.2	93
14	A functional CNVR_3425.1 damping lincRNA FENDRR increases lifetime risk of lung cancer and COPD in Chinese. <i>Carcinogenesis</i> , 2018, 39, 347-359.	2.8	16
15	Upregulation of LncRNA FEZF-AS1 is associated with advanced clinical stages and family history of cancer in patients with NSCLC. <i>Pathology Research and Practice</i> , 2018, 214, 857-861.	2.3	14
16	Long non&€coding RNA AGER&€1 functionally upregulates the innate immunity gene AGER and approximates its anti&€tumor effect in lung cancer. <i>Molecular Carcinogenesis</i> , 2018, 57, 305-318.	2.7	29
17	Long Noncoding RNA PRRG4-4 Promotes Viability, Cell Cycle, Migration, and Invasion in Lung Cancer Cells. <i>DNA and Cell Biology</i> , 2018, 37, 953-966.	1.9	3
18	PIWI-interacting RNA-54265 is oncogenic and a potential therapeutic target in colorectal adenocarcinoma. <i>Theranostics</i> , 2018, 8, 5213-5230.	10.0	115

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19	Upregulation of long non-coding RNA RAB1A-2 induces FGF1 expression worsening lung cancer prognosis. <i>Cancer Letters</i> , 2018, 438, 116-125.	7.2	31
20	Functional Polymorphism in the MSI1 Gene Promoter Confers a Decreased Risk of Lung Cancer in Chinese by Reducing MSI1 Expression. <i>Current Genomics</i> , 2018, 19, 375-383.	1.6	6
21	Association of nsv823469 copy number loss with decreased risk of chronic obstructive pulmonary disease and pulmonary function in Chinese. <i>Scientific Reports</i> , 2017, 7, 40060.	3.3	10
22	Long non-coding RNA LINC00672 contributes to p53 protein-mediated gene suppression and promotes endometrial cancer chemosensitivity. <i>Journal of Biological Chemistry</i> , 2017, 292, 5801-5813.	3.4	54
23	Rare variant of <i>MAP2K7</i> is associated with increased risk of COPD in southern and eastern Chinese. <i>Respirology</i> , 2017, 22, 691-698.	2.3	1
24	The MKK7 p.Glu116Lys Rare Variant Serves as a Predictor for Lung Cancer Risk and Prognosis in Chinese. <i>PLoS Genetics</i> , 2016, 12, e1005955.	3.5	14
25	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
26	Circular RNA ITCH has inhibitory effect on ESCC by suppressing the Wnt/ $\beta$ -catenin pathway. <i>Oncotarget</i> , 2015, 6, 6001-6013.	1.8	626
27	Polymorphisms of NF $\kappa$ B1 and B1 and Their Synergistic Effect on Nasopharyngeal Carcinoma Susceptibility. <i>BioMed Research International</i> , 2015, 2015, 1-9.	1.9	7
28	Polymorphism in mature microRNA-608 sequence is associated with an increased risk of nasopharyngeal carcinoma. <i>Gene</i> , 2015, 565, 180-186.	2.2	34
29	Sequence Variation in Mature MicroRNA-499 Confers Unfavorable Prognosis of Lung Cancer Patients Treated with Platinum-Based Chemotherapy. <i>Clinical Cancer Research</i> , 2015, 21, 1602-1613.	7.0	37
30	Duplicated copy of CHRNA7 increases risk and worsens prognosis of COPD and lung cancer. <i>European Journal of Human Genetics</i> , 2015, 23, 1019-1024.	2.8	20
31	The Functional Copy Number Variation-67048 in <i>WWOX</i> Contributes to Increased Risk of COPD in Southern and Eastern Chinese. <i>COPD: Journal of Chronic Obstructive Pulmonary Disease</i> , 2015, 12, 494-501.	1.6	8
32	Functional polymorphisms in the <i>CD44</i> gene and acute myeloid leukemia cancer risk in a Chinese population. <i>Molecular Carcinogenesis</i> , 2015, 54, 102-110.	2.7	13
33	A Functional Polymorphism in the 3'-UTR of PXR Interacts with Smoking to Increase Lung Cancer Risk in Southern and Eastern Chinese Smoker. <i>International Journal of Molecular Sciences</i> , 2014, 15, 17457-17468.	4.1	11
34	Genetic variant in the 3'-untranslated region of VEGFR1 gene influences chronic obstructive pulmonary disease and lung cancer development in Chinese population. <i>Mutagenesis</i> , 2014, 29, 311-317.	2.6	14
35	The functional polymorphism of NBS1 p.Glu185Gln is associated with an increased risk of lung cancer in Chinese populations: Case-control and a meta-analysis. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2014, 770, 61-68.	1.0	18
36	Identification of chimeric TSNAX-DISC1 resulting from intergenic splicing in endometrial carcinoma through high-throughput RNA sequencing. <i>Carcinogenesis</i> , 2014, 35, 2687-2697.	2.8	36

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37	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
38	Increased Levels of the Long Intergenic Non-coding Protein Coding RNA POU3F3 Promote DNA Methylation in Esophageal Squamous Cell Carcinoma Cells. <i>Gastroenterology</i> , 2014, 146, 1714-1726.e5.	1.3	169
39	A Polymorphism rs12325489C>T in the lincRNA-ENST00000515084 Exon Was Found to Modulate Breast Cancer Risk via GWAS-Based Association Analyses. <i>PLoS ONE</i> , 2014, 9, e98251.	2.5	36
40	A Newfound Association between MDC1 Functional Polymorphism and Lung Cancer Risk in Chinese. <i>PLoS ONE</i> , 2014, 9, e106794.	2.5	7
41	NBS1 rs1805794G>C polymorphism is associated with decreased risk of acute myeloid leukemia in a Chinese population. <i>Molecular Biology Reports</i> , 2013, 40, 3749-3756.	2.3	15
42	Functional polymorphisms in PIN1 promoter and esophageal carcinoma susceptibility in Chinese population. <i>Molecular Biology Reports</i> , 2013, 40, 829-838.	2.3	21
43	A genetic polymorphism in lincRNA-uc003opf.1 is associated with susceptibility to esophageal squamous cell carcinoma in Chinese populations. <i>Carcinogenesis</i> , 2013, 34, 2908-2917.	2.8	61
44	IL-21 gene polymorphism is associated with the prognosis of breast cancer in Chinese populations. <i>Breast Cancer Research and Treatment</i> , 2013, 137, 893-901.	2.5	23
45	A Sequence Polymorphism in miR-608 Predicts Recurrence after Radiotherapy for Nasopharyngeal Carcinoma. <i>Cancer Research</i> , 2013, 73, 5151-5162.	0.9	64
46	Heterozygous Genetic Variations of FOXP3 in Xp11.23 Elevate Breast Cancer Risk in Chinese Population via Skewed X-Chromosome Inactivation. <i>Human Mutation</i> , 2013, 34, n/a-n/a.	2.5	26
47	A functional polymorphism at microRNA-629-binding site in the 3'-untranslated region of NBS1 gene confers an increased risk of lung cancer in Southern and Eastern Chinese population. <i>Carcinogenesis</i> , 2012, 33, 338-347.	2.8	80
48	CD44 rs13347 C>T polymorphism predicts breast cancer risk and prognosis in Chinese populations. <i>Breast Cancer Research</i> , 2012, 14, R105.	5.0	52
49	Functional genetic variations in the IL-23 receptor gene are associated with risk of breast, lung and nasopharyngeal cancer in Chinese populations. <i>Carcinogenesis</i> , 2012, 33, 2409-2416.	2.8	55
50	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
51	Functional polymorphism in the EpCAM gene is associated with occurrence and advanced disease status of cervical cancer in Chinese population. <i>Molecular Biology Reports</i> , 2012, 39, 7303-7309.	2.3	8
52	The protective role of polymorphism MKK4 rs1304 T>G in nasopharyngeal carcinoma is modulated by Epstein-Barr virus' infection status. <i>International Journal of Cancer</i> , 2012, 130, 1981-1990.	5.1	32
53	Genetic Variations in CD14 Promoter and Acute Lymphoblastic Leukemia Susceptibility in a Chinese Population. <i>DNA and Cell Biology</i> , 2011, 30, 777-782.	1.9	10
54	Genome-wide association study identifies three new susceptibility loci for esophageal squamous-cell carcinoma in Chinese populations. <i>Nature Genetics</i> , 2011, 43, 679-684.	21.4	260

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55	A genome-wide association study identifies new susceptibility loci for non-cardia gastric cancer at 3q13.31 and 5p13.1. <i>Nature Genetics</i> , 2011, 43, 1215-1218.	21.4	250
56	Functional variant (â <sup>1304T</sup> &gt;G) in the <i>MKK4</i> promoter is associated with decreased risk of acute myeloid leukemia in a southern Chinese population. <i>Cancer Science</i> , 2011, 102, 1462-1468.	3.9	13
57	Functional polymorphisms in the NBS1 gene and acute lymphoblastic leukemia susceptibility in a Chinese population. <i>European Journal of Haematology</i> , 2011, 86, 199-205.	2.2	23
58	Functional genetic variations of cyclooxygenase-2 and susceptibility to acute myeloid leukemia in a Chinese population. <i>European Journal of Haematology</i> , 2011, 87, 486-493.	2.2	18
59	A non-synonymous polymorphism Thr115Met in the EpCAM gene is associated with an increased risk of breast cancer in Chinese population. <i>Breast Cancer Research and Treatment</i> , 2011, 126, 487-495.	2.5	45
60	Functional <i>NBS1</i> polymorphism is associated with occurrence and advanced disease status of nasopharyngeal carcinoma. <i>Molecular Carcinogenesis</i> , 2011, 50, 689-696.	2.7	48
61	The polymorphism and haplotypes of PIN1 gene are associated with the risk of lung cancer in southern and eastern chinese populations. <i>Human Mutation</i> , 2011, 32, 1299-1308.	2.5	59
62	A protein-based electrochemical method for label-free characterization of sequence-specific proteinâ€DNA interactions. <i>Electrochimica Acta</i> , 2011, 56, 5759-5765.	5.2	6
63	Association between the Cytotoxic T-Lymphocyte Antigen 4 +49G &gt; A polymorphism and cancer risk: a meta-analysis. <i>BMC Cancer</i> , 2010, 10, 522.	2.6	30