## Yang Zhao

## List of Publications by Year in descending order

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279778 289230 2,145 96 23 40 citations h-index g-index papers 102 102 102 3463 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Effect of Anlotinib as a Third-Line or Further Treatment on Overall Survival of Patients With Advanced Non–Small Cell Lung Cancer. JAMA Oncology, 2018, 4, 1569.	7.1	388
2	Development and validation of an immune gene-set based Prognostic signature in ovarian cancer. EBioMedicine, 2019, 40, 318-326.	6.1	209
3	Statistical analysis for genome-wide association study. Journal of Biomedical Research, 2015, 29, 285.	1.6	73
4	Association between early treatment with Qingfei Paidu decoction and favorable clinical outcomes in patients with COVID-19: A retrospective multicenter cohort study. Pharmacological Research, 2020, 161, 105290.	7.1	67
5	Seven-CpG-based prognostic signature coupled with gene expression predicts survival of oral squamous cell carcinoma. Clinical Epigenetics, 2017, 9, 88.	4.1	65
6	Comprehensive analyses of m6A regulators and interactive coding and non-coding RNAs across 32 cancer types. Molecular Cancer, 2021, 20, 67.	19.2	65
7	A genome-wide gene-environment interaction analysis for tobacco smoke and lung cancer susceptibility. Carcinogenesis, 2014, 35, 1528-1535.	2.8	47
8	An Integrated Analysis of miRNA, IncRNA, and mRNA Expression Profiles. BioMed Research International, 2014, 2014, 1-12.	1.9	45
9	Integrative Analysis of miRNA-mRNA and miRNA-miRNA Interactions. BioMed Research International, 2014, 2014, 1-8.	1.9	45
10	Association Between δ-Aminolevulinic Acid Dehydratase (ALAD) Polymorphism and Blood Lead Levels: A Meta-regression Analysis. Journal of Toxicology and Environmental Health - Part A: Current Issues, 2007, 70, 1986-1994.	2.3	44
11	Relationship between Branched-Chain Amino Acids, Metabolic Syndrome, and Cardiovascular Risk Profile in a Chinese Population: A Cross-Sectional Study. International Journal of Endocrinology, 2016, 2016, 1-10.	1.5	39
12	Anlotinib for previously treated advanced or metastatic esophageal squamous cell carcinoma: A doubleâ€blind randomized phase 2 trial. Cancer Medicine, 2021, 10, 1681-1689.	2.8	39
13	Modeling the transmission dynamics of COVID-19 epidemic: a systematic review. Journal of Biomedical Research, 2020, 34, 422.	1.6	35
14	Cardiac Injury and COVID-19: A Systematic Review and Meta-analysis. CJC Open, 2020, 2, 386-394.	1.5	33
15	Task-related functional magnetic resonance imaging-based neuronavigation for the treatment of depression by individualized repetitive transcranial magnetic stimulation of the visual cortex. Science China Life Sciences, 2021, 64, 96-106.	4.9	33
16	Potential clinical value of circular RNAs as peripheral biomarkers for the diagnosis and treatment of major depressive disorder. EBioMedicine, 2021, 66, 103337.	6.1	33
17	TOBMI: trans-omics block missing data imputation using a k-nearest neighbor weighted approach. Bioinformatics, 2019, 35, 1278-1283.	4.1	32
18	Invasive adenocarcinoma of the lung is associated with the upper lung regions. Lung Cancer, 2014, 84, 145-150.	2.0	31

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19	Comparisons of isomiR patterns and classification performance using the rank-based MANOVA and 10-fold cross-validation. Gene, 2015, 569, 21-26.	2.2	29
20	A seven-gene prognostic signature for rapid determination of head and neck squamous cell carcinoma survival. Oncology Reports, 2017, 38, 3403-3411.	2.6	29
21	Prognostic factors of refractory NSCLC patients receiving anlotinib hydrochloride as the third- or further-line treatment. Cancer Biology and Medicine, 2018, 15, 443.	3.0	29
22	Gene copy number alterations in the azoospermia-associated AZFc region and their effect on spermatogenic impairment. Molecular Human Reproduction, 2014, 20, 836-843.	2.8	27
23	Association of assisted reproductive technology, germline de novo mutations and congenital heart defects in a prospective birth cohort study. Cell Research, 2021, 31, 919-928.	12.0	26
24	A Coding Single-Nucleotide Polymorphism in Lysine Demethylase <i>KDM4A</i> Associates with Increased Sensitivity to mTOR Inhibitors. Cancer Discovery, 2015, 5, 245-254.	9.4	25
25	Computed tomography measurement of pulmonary artery for diagnosis of COPD and its comorbidity pulmonary hypertension. International Journal of COPD, 2015, 10, 2525.	2.3	24
26	Evolutionary and Expression Analysis of miR-#-5p and miR-#-3p at the miRNAs/isomiRs Levels. BioMed Research International, 2015, 2015, 1-14.	1.9	24
27	Assisted reproductive technology and birth defects in a Chinese birth cohort study. The Lancet Regional Health - Western Pacific, 2021, 7, 100090.	2.9	24
28	Polymorphisms in MicroRNAs Are Associated with Survival in Non–Small Cell Lung Cancer. Cancer Epidemiology Biomarkers and Prevention, 2014, 23, 2503-2511.	2.5	22
29	Elevated Platelet Count Appears to Be Causally Associated with Increased Risk of Lung Cancer: A Mendelian Randomization Analysis. Cancer Epidemiology Biomarkers and Prevention, 2019, 28, 935-942.	2.5	21
30	Association of three single nucleotide polymorphisms of ESR1 with breast cancer susceptibility: a meta-analysis. Journal of Biomedical Research, 2017, 31, 213.	1.6	20
31	Variable selection approach for zero-inflated count data via adaptive lasso. Journal of Applied Statistics, 2014, 41, 879-894.	1.3	19
32	Reconstructing the coding and non-coding RNA regulatory networks of miRNAs and mRNAs in breast cancer. Gene, 2014, 548, 6-13.	2.2	18
33	Association between oral contraceptives and risk of hemorrhagic stroke: a meta-analysis of observational studies. Archives of Gynecology and Obstetrics, 2018, 297, 1181-1191.	1.7	18
34	Adiponectin Gene Polymorphisms and Acute Respiratory Distress Syndrome Susceptibility and Mortality. PLoS ONE, 2014, 9, e89170.	2.5	18
35	A Large-Scale Genome-Wide Gene-Gene Interaction Study of Lung Cancer Susceptibility in Europeans With a Trans-Ethnic Validation in Asians. Journal of Thoracic Oncology, 2022, 17, 974-990.	1.1	18
36	Cord serum elementomics profiling of 56 elements depicts risk of preterm birth: Evidence from a prospective birth cohort in rural Bangladesh. Environment International, 2021, 156, 106731.	10.0	17

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37	Identifying Plasma Biomarkers with high specificity for major depressive disorder: A multi-level proteomics study. Journal of Affective Disorders, 2020, 277, 620-630.	4.1	16
38	Association between dietary sodium, potassium intake and lung cancer risk: evidence from the prostate, lung, colorectal and ovarian cancer screening trial and the Women's Health Initiative. Translational Lung Cancer Research, 2021, 10, 45-56.	2.8	16
39	Mitochondrial Variations in Non-Small Cell Lung Cancer (NSCLC) Survival. Cancer Informatics, 2015, 14s1, CIN.S13976.	1.9	15
40	A Comparative Study of Five Association Tests Based on CpG Set for Epigenome-Wide Association Studies. PLoS ONE, 2016, 11, e0156895.	2.5	15
41	The Efficacy and Toxicity of Gefitinib in Treating Non-small Cell Lung Cancer: A Meta-analysis of 19 Randomized Clinical Trials. Journal of Cancer, 2018, 9, 1455-1465.	2.5	15
42	Implementation of Clinical Diagnostic Criteria and Universal Symptom Survey Contributed to Lower Magnitude and Faster Resolution of the COVID-19 Epidemic in Wuhan. Engineering, 2020, 6, 1141-1146.	6.7	14
43	Global Analysis of miRNA Gene Clusters and Gene Families Reveals Dynamic and Coordinated Expression. BioMed Research International, 2014, 2014, 1-7.	1.9	13
44	Trans-omics biomarker model improves prognostic prediction accuracy for early-stage lung adenocarcinoma. Aging, 2019, 11, 6312-6335.	3.1	13
45	Better modelling of infectious diseases: lessons from covid-19 in China. BMJ, The, 2021, 375, n2365.	6.0	13
46	Prediction of acute kidney injury after cardiac surgery: model development using a Chinese electronic health record dataset. Journal of Translational Medicine, 2022, 20, 166.	4.4	13
47	Permutation-based variance component test in generalized linear mixed model with application to multilocus genetic association study. BMC Medical Research Methodology, 2015, 15, 37.	3.1	12
48	Exome-Wide Association Study Identified New Risk Loci for Hirschsprung's Disease. Molecular Neurobiology, 2017, 54, 1777-1785.	4.0	12
49	Epigenetic modifications in KDM lysine demethylases associate with survival of early-stage NSCLC. Clinical Epigenetics, 2018, 10, 41.	4.1	12
50	Evaluating the effect of multiple genetic risk score models on colorectal cancer risk prediction. Gene, 2018, 673, 174-180.	2.2	12
51	Comprehensive estimation for the length and dispersion of COVID-19 incubation period: a systematic review and meta-analysis. Infection, 2022, 50, 803-813.	4.7	12
52	Mutantâ€'allele fraction heterogeneity is associated with nonâ€'small cell lung cancer patient survival. Oncology Letters, 2017, 15, 795-802.	1.8	11
53	Bayesian variable selection for parametric survival model with applications to cancer omics data. Human Genomics, 2018, 12, 49.	2.9	11
54	Safety profile of poly (ADP-ribose) polymerase (PARP) inhibitors in cancer: a network meta-analysis of randomized controlled trials. Annals of Translational Medicine, 2021, 9, 1229-1229.	1.7	10

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55	Global COVID-19 Pandemic Waves: Limited Lessons Learned Worldwide over the Past Year. Engineering, 2022, 13, 91-98.	6.7	10
56	APOLLO: An accurate and independently validated prediction model of lower-grade gliomas overall survival and a comparative study of model performance. EBioMedicine, 2022, 79, 104007.	6.1	10
57	SCALE-1: Safety and efficacy of short course neoadjuvant chemo-radiotherapy plus toripalimab for locally advanced resectable squamous cell carcinoma of esophagus Journal of Clinical Oncology, 2022, 40, 4063-4063.	1.6	10
58	A multi-omics study links TNS3 and SEPT7 to long-term former smoking NSCLC survival. Npj Precision Oncology, 2021, 5, 39.	5.4	9
59	A predictive paradigm for COVID-19 prognosis based on the longitudinal measure of biomarkers. Briefings in Bioinformatics, 2021, 22, .	<b>6.</b> 5	9
60	Rare Variants Detection with Kernel Machine Learning Based on Likelihood Ratio Test. PLoS ONE, 2014, 9, e93355.	2.5	9
61	A Systematic Evaluation of Feature Selection and Classification Algorithms Using Simulated and Real miRNA Sequencing Data. Computational and Mathematical Methods in Medicine, 2015, 2015, 1-11.	1.3	8
62	ESR2 Genetic Variants and Combined Oral Contraceptive Use Associated with the Risk of Stroke. Archives of Medical Research, 2017, 48, 203-211.	3.3	8
63	SHARED: Efficacy and safety of sintilimab in combination with concurrent chemoradiotherapy (cCRT) in patients with locally advanced gastric (G) or gastroesophageal junction (GEJ) adenocarcinoma Journal of Clinical Oncology, 2021, 39, 4040-4040.	1.6	8
64	Profiling and Bioinformatic Analyses Indicate Differential circRNA and miRNA/isomiR Expression and Interactions. BioMed Research International, 2018, 2018, 1-9.	1.9	7
65	Circulating PCSK9 levels and 2-hPG are positively correlated in metabolic diseases in a Chinese Han population. Lipids in Health and Disease, 2018, 17, 15.	3.0	7
66	The relationship between treatment-induced hypertension and efficacy of anlotinib in recurrent or metastatic esophageal squamous cell carcinoma. Cancer Biology and Medicine, 2021, 18, 562-568.	3.0	7
67	Optimal, minimax and admissible two-stage design for phase II oncology clinical trials. BMC Medical Research Methodology, 2020, 20, 126.	3.1	7
68	Association between dietary intake of polyunsaturated fatty acid and prevalence of hypertension in U.S. adults: A cross-sectional study using data from NHANES 2009–2016. Hypertension Research, 2022, 45, 516-526.	2.7	7
69	Accuracy of three-dimensional photogrammetry and cone beam computed tomography based on linear measurements in patients with facial deformities. Dentomaxillofacial Radiology, 2021, 50, 20200001.	2.7	6
70	Associations of dietary PUFA with dyslipidaemia among the US adults: the findings from National Health and Nutrition Examination Survey (NHANES) 2009–2016. British Journal of Nutrition, 2022, 127, 1386-1394.	2.3	6
71	A fast algorithm for Bayesian multi-locus model in genome-wide association studies. Molecular Genetics and Genomics, 2017, 292, 923-934.	2.1	5
72	A case report of acute simultaneous cardiocerebral infarction: possible pathophysiology. Annals of Palliative Medicine, 2021, 10, 5887-5890.	1.2	5

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73	Appealing for efficient, well organized clinical trials on COVID-19. Annals of Translational Medicine, 2020, 8, 632-632.	1.7	4
74	Variance component testing for identifying differentially expressed genes in RNA-seq data. PeerJ, 2017, 5, e3797.	2.0	4
75	IPRS: Leveraging Gene-Environment Interaction to Reconstruct Polygenic Risk Score. Frontiers in Genetics, 2022, 13, 801397.	2.3	4
76	Application of cloud computing in biomedicine big data analysis cloud computing in big data., 2017,,.		3
77	Intelligent quantitative assessment of skeletal maturation based on multi-stage model: a retrospective cone-beam CT study of cervical vertebrae. Oral Radiology, 2022, 38, 378-388.	1.9	3
78	Transmission dynamics model and the coronavirus disease 2019 epidemic: applications and challenges. Medical Review, 2022, 2, 89-109.	1.2	3
79	Identifying and exploiting gene-pathway interactions from RNA-seq data for binary phenotype. BMC Genetics, 2019, 20, 36.	2.7	2
80	Identification of genetic features associated with fine particulate matter (PM2.5) modulated DNA damage using improved random forest analysis. Gene, 2020, 740, 144570.	2.2	2
81	Multiple arterial conduits for multi-vessel coronary artery bypass grafting in patients with mild to moderate left ventricular systolic dysfunction: a multicenter retrospective study. Journal of Cardiothoracic Surgery, 2021, 16, 123.	1.1	2
82	Association between aspirin use and lung cancer incidence depends on high-frequency use, bodyweight, and age in U.S. adults. Translational Lung Cancer Research, 2021, 10, 392-401.	2.8	2
83	Genetic Association Analysis Using Sibship Data: A Multilevel Model Approach. PLoS ONE, 2012, 7, e31134.	2.5	2
84	Dynamic trajectory of platelet counts after the first cycle of induction chemotherapy in AML patients. BMC Cancer, 2022, 22, 477.	2.6	2
85	Common Statistical Methods and Reporting of Results in Medical Research. Cardiovascular Innovations and Applications, 2022, 6, .	0.3	2
86	Forecasting of COVID-19: transmission models and beyond. Journal of Thoracic Disease, 2020, 12, 1762-1765.	1.4	1
87	Genome-wide gene–smoking interaction study identified novel susceptibility loci for non-small cell lung cancer in Chinese populations. Carcinogenesis, 2021, 42, 1154-1161.	2.8	1
88	Potential of Antithrombin III as a Biomarker of Antidepressive Effect in Major Depressive Disorder. Frontiers in Psychiatry, 2021, 12, 678384.	2.6	1
89	Association of Hypokalemia Incidence and Better Treatment Response in NSCLC Patients: A Meta-Analysis and Systematic Review on Anti-EGFR Targeted Therapy Clinical Trials. Frontiers in Oncology, 2021, 11, 757456.	2.8	1
90	Development of a multi-stage model for intelligent and quantitative appraising of skeletal maturity using cervical vertebras cone-beam CT images of Chinese girls. International Journal of Computer Assisted Radiology and Surgery, 2022, 17, 761.	2.8	1

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91	Heterogeneity of Treatment Effects for Intensive Blood Pressure Therapy by Individual Components of FRS: An Unsupervised Data-Driven Subgroup Analysis in SPRINT and ACCORD. Frontiers in Cardiovascular Medicine, 2022, 9, 778756.	2.4	1
92	Mid-term outcomes of coronary artery bypass grafting in patients with mild left ventricular systolic dysfunction: a multicentre retrospective cohort study. Interactive Cardiovascular and Thoracic Surgery, 2021, 32, 855-863.	1.1	0
93	Comment on the article: "Prediction of the development of acute kidney injury following cardiac surgery by machine learning― Critical Care, 2021, 25, 374.	5.8	O
94	A dynamic model for individualized prognosis prediction in patients with avian influenza A H7N9. Annals of Translational Medicine, 2022, 10, 149-149.	1.7	0
95	Effects of interaction between single nucleotide polymorphisms and psychosocial factors on the response to antidepressant treatment in patients with major depressive disorder. Journal of Genetics and Genomics, 2021, 49, 587-587.	3.9	0
96	Risk difference, relative risk, and odds ratio for non-inferiority clinical trials with risk rate endpoint. Journal of Biopharmaceutical Statistics, $0$ , , $1-16$ .	0.8	0