Yang Zhao

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

1,166 18 91 30 g-index h-index citations papers 1,683 102 4.42 5.3 L-index avg, IF ext. citations ext. papers

#	Paper	IF	Citations
91	A dynamic model for individualized prognosis prediction in patients with avian influenza A H7N9 <i>Annals of Translational Medicine</i> , 2022 , 10, 149	3.2	
90	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 <i>Hypertension Research</i> , 2022 ,	4.7	1
89	Development of a multi-stage model for intelligent and quantitative appraising of skeletal maturity using cervical vertebras cone-beam CT images of Chinese girls <i>International Journal of Computer Assisted Radiology and Surgery</i> , 2022 , 17, 761	3.9	O
88	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	5.4	
87	Transmission dynamics model and the coronavirus disease 2019 epidemic: applications and challenges 2022 , 2, 89-109		1
86	IPRS: Leveraging Gene-Environment Interaction to Reconstruct Polygenic Risk Score <i>Frontiers in Genetics</i> , 2022 , 13, 801397	4.5	0
85	Prediction of acute kidney injury after cardiac surgery: model development using a Chinese electronic health record dataset <i>Journal of Translational Medicine</i> , 2022 , 20, 166	8.5	0
84	APOLLO: An accurate and independently validated prediction model of lower-grade gliomas overall survival and a comparative study of model performance <i>EBioMedicine</i> , 2022 , 79, 104007	8.8	0
83	Dynamic trajectory of platelet counts after the first cycle of induction chemotherapy in AML patients <i>BMC Cancer</i> , 2022 , 22, 477	4.8	O
82	Common Statistical Methods and Reporting of Results in Medical Research. <i>Cardiovascular Innovations and Applications</i> , 2022 , 6, 117-125	0.1	
81	Better modelling of infectious diseases: lessons from covid-19 in China. <i>BMJ, The</i> , 2021 , 375, n2365	5.9	1
80	Association of Hypokalemia Incidence and Better Treatment Response in NSCLC Patients: A Meta-Analysis and Systematic Review on Anti-EGFR Targeted Therapy Clinical Trials <i>Frontiers in Oncology</i> , 2021 , 11, 757456	5.3	0
79	Comment on the article: "Prediction of the development of acute kidney injury following cardiac surgery by machine learning". <i>Critical Care</i> , 2021 , 25, 374	10.8	
78	Potential of Antithrombin III as a Biomarker of Antidepressive Effect in Major Depressive Disorder. <i>Frontiers in Psychiatry</i> , 2021 , 12, 678384	5	1
77	Potential clinical value of circular RNAs as peripheral biomarkers for the diagnosis and treatment of major depressive disorder. <i>EBioMedicine</i> , 2021 , 66, 103337	8.8	13
76	Comprehensive analyses of m6A regulators and interactive coding and non-coding RNAs across 32 cancer types. <i>Molecular Cancer</i> , 2021 , 20, 67	42.1	13
75	A case report of acute simultaneous cardiocerebral infarction: possible pathophysiology. <i>Annals of Palliative Medicine</i> , 2021 , 10, 5887-5890	1.7	2

(2021-2021)

74	SHARED: Efficacy and safety of sintilimab in combination with concurrent chemoradiotherapy (cCRT) in patients with locally advanced gastric (G) or gastroesophageal junction (GEJ) adenocarcinoma <i>Journal of Clinical Oncology</i> , 2021 , 39, 4040-4040	2.2	О
73	A multi-omics study links TNS3 and SEPT7 to long-term former smoking NSCLC survival. <i>Npj Precision Oncology</i> , 2021 , 5, 39	9.8	1
7 ²	Multiple arterial conduits for multi-vessel coronary artery bypass grafting in patients with mild to moderate left ventricular systolic dysfunction: a multicenter retrospective study. <i>Journal of Cardiothoracic Surgery</i> , 2021 , 16, 123	1.6	1
71	Associations of dietary PUFA with dyslipidaemia among the US adults: the findings from National Health and Nutrition Examination Survey (NHANES) 2009-2016. <i>British Journal of Nutrition</i> , 2021 , 1-9	3.6	1
70	A predictive paradigm for COVID-19 prognosis based on the longitudinal measure of biomarkers. <i>Briefings in Bioinformatics</i> , 2021 , 22,	13.4	2
69	Association of assisted reproductive technology, germline de novo mutations and congenital heart defects in a prospective birth cohort study. <i>Cell Research</i> , 2021 , 31, 919-928	24.7	7
68	Task-related functional magnetic resonance imaging-based neuronavigation for the treatment of depression by individualized repetitive transcranial magnetic stimulation of the visual cortex. <i>Science China Life Sciences</i> , 2021 , 64, 96-106	8.5	10
67	Accuracy of three-dimensional photogrammetry and cone beam computed tomography based on linear measurements in patients with facial deformities. <i>Dentomaxillofacial Radiology</i> , 2021 , 50, 202000	જે1 ⁹	2
66	Association between dietary sodium, potassium intake and lung cancer risk: evidence from the prostate, lung, colorectal and ovarian cancer screening trial and the Women® Health Initiative. <i>Translational Lung Cancer Research</i> , 2021 , 10, 45-56	4.4	1
65	The relationship between treatment-induced hypertension and efficacy of anlotinib in recurrent or metastatic esophageal squamous cell carcinoma. <i>Cancer Biology and Medicine</i> , 2021 ,	5.2	6
64	Mid-term outcomes of coronary artery bypass grafting in patients with mild left ventricular systolic dysfunction: a multicentre retrospective cohort study. <i>Interactive Cardiovascular and Thoracic Surgery</i> , 2021 , 32, 855-863	1.8	
63	Anlotinib for previously treated advanced or metastatic esophageal squamous cell carcinoma: A double-blind randomized phase 2 trial. <i>Cancer Medicine</i> , 2021 , 10, 1681-1689	4.8	8
62	Assisted reproductive technology and birth defects in a Chinese birth cohort study. <i>The Lancet Regional Health - Western Pacific</i> , 2021 , 7, 100090	5	5
61	Genome-wide gene-smoking interaction study identified novel susceptibility loci for non-small cell lung cancer in Chinese populations. <i>Carcinogenesis</i> , 2021 , 42, 1154-1161	4.6	О
60	Safety profile of poly (ADP-ribose) polymerase (PARP) inhibitors in cancer: a network meta-analysis of randomized controlled trials. <i>Annals of Translational Medicine</i> , 2021 , 9, 1229	3.2	2
59	Comprehensive estimation for the length and dispersion of COVID-19 incubation period: a systematic review and meta-analysis. <i>Infection</i> , 2021 , 1	5.8	2
58	Intelligent quantitative assessment of skeletal maturation based on multi-stage model: a retrospective cone-beam CT study of cervical vertebrae. <i>Oral Radiology</i> , 2021 , 1	2.5	1
57	Cord serum elementomics profiling of 56 elements depicts risk of preterm birth: Evidence from a prospective birth cohort in rural Bangladesh. <i>Environment International</i> , 2021 , 156, 106731	12.9	4

56	Association between aspirin use and lung cancer incidence depends on high-frequency use, bodyweight, and age in U.S. adults. <i>Translational Lung Cancer Research</i> , 2021 , 10, 392-401	4.4	O
55	Effects of interaction between single nucleotide polymorphisms and psychosocial factors on the response to antidepressant treatment in patients with major depressive disorder <i>Journal of Genetics and Genomics</i> , 2021 , 49, 587-587	4	
54	Implementation of Clinical Diagnostic Criteria and Universal Symptom Survey Contributed to Lower Magnitude and Faster Resolution of the COVID-19 Epidemic in Wuhan. <i>Engineering</i> , 2020 , 6, 1141-1146	9.7	10
53	Appealing for efficient, well organized clinical trials on COVID-19. <i>Annals of Translational Medicine</i> , 2020 , 8, 632	3.2	2
52	Identification of genetic features associated with fine particulate matter (PM2.5) modulated DNA damage using improved random forest analysis. <i>Gene</i> , 2020 , 740, 144570	3.8	O
51	Cardiac Injury and COVID-19: A Systematic Review and Meta-analysis. <i>CJC Open</i> , 2020 , 2, 386-394	2	23
50	Modeling the transmission dynamics of COVID-19 epidemic: a systematic review. <i>Journal of Biomedical Research</i> , 2020 , 34, 422-430	1.5	14
49	Optimal, minimax and admissible two-stage design for phase II oncology clinical trials. <i>BMC Medical Research Methodology</i> , 2020 , 20, 126	4.7	1
48	Association between early treatment with Qingfei Paidu decoction and favorable clinical outcomes in patients with COVID-19: A retrospective multicenter cohort study. <i>Pharmacological Research</i> , 2020 , 161, 105290	10.2	33
47	Identifying Plasma Biomarkers with high specificity for major depressive disorder: A multi-level proteomics study. <i>Journal of Affective Disorders</i> , 2020 , 277, 620-630	6.6	4
46	Identifying and exploiting gene-pathway interactions from RNA-seq data for binary phenotype. <i>BMC Genetics</i> , 2019 , 20, 36	2.6	2
45	Elevated Platelet Count Appears to Be Causally Associated with Increased Risk of Lung Cancer: A Mendelian Randomization Analysis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2019 , 28, 935-942	4	12
44	Trans-omics biomarker model improves prognostic prediction accuracy for early-stage lung adenocarcinoma. <i>Aging</i> , 2019 , 11, 6312-6335	5.6	8
43	TOBMI: trans-omics block missing data imputation using a k-nearest neighbor weighted approach. <i>Bioinformatics</i> , 2019 , 35, 1278-1283	7.2	18
42	Development and validation of an immune gene-set based Prognostic signature in ovarian cancer. <i>EBioMedicine</i> , 2019 , 40, 318-326	8.8	106
41	Association between oral contraceptives and risk of hemorrhagic stroke: a meta-analysis of observational studies. <i>Archives of Gynecology and Obstetrics</i> , 2018 , 297, 1181-1191	2.5	11
40	Epigenetic modifications in lysine demethylases associate with survival of early-stage NSCLC. <i>Clinical Epigenetics</i> , 2018 , 10, 41	7.7	11
39	Circulating PCSK9 levels and 2-hPG are positively correlated in metabolic diseases in a Chinese Han population. <i>Lipids in Health and Disease</i> , 2018 , 17, 15	4.4	6

(2015-2018)

Effect of Anlotinib as a Third-Line or Further Treatment on Overall Survival of Patients With Advanced Non-Small Cell Lung Cancer: The ALTER 0303 Phase 3 Randomized Clinical Trial. <i>JAMA Oncology</i> , 2018 , 4, 1569-1575	13.4	218
Evaluating the effect of multiple genetic risk score models on colorectal cancer risk prediction. <i>Gene</i> , 2018 , 673, 174-180	3.8	9
Mutant-allele fraction heterogeneity is associated with non-small cell lung cancer patient survival. <i>Oncology Letters</i> , 2018 , 15, 795-802	2.6	9
Prognostic factors of refractory NSCLC patients receiving anlotinib hydrochloride as the third- or further-line treatment. <i>Cancer Biology and Medicine</i> , 2018 , 15, 443-451	5.2	19
Bayesian variable selection for parametric survival model with applications to cancer omics data. <i>Human Genomics</i> , 2018 , 12, 49	6.8	7
The Efficacy and Toxicity of Gefitinib in Treating Non-small Cell Lung Cancer: A Meta-analysis of 19 Randomized Clinical Trials. <i>Journal of Cancer</i> , 2018 , 9, 1455-1465	4.5	11
Profiling and Bioinformatic Analyses Indicate Differential circRNA and miRNA/isomiR Expression and Interactions. <i>BioMed Research International</i> , 2018 , 2018, 8518563	3	5
Exome-Wide Association Study Identified New Risk Loci for Hirschsprung B Disease. <i>Molecular Neurobiology</i> , 2017 , 54, 1777-1785	6.2	7
ESR2 Genetic Variants and Combined Oral Contraceptive Use Associated with the Risk of Stroke. <i>Archives of Medical Research</i> , 2017 , 48, 203-211	6.6	4
A fast algorithm for Bayesian multi-locus model in genome-wide association studies. <i>Molecular Genetics and Genomics</i> , 2017 , 292, 923-934	3.1	4
Seven-CpG-based prognostic signature coupled with gene expression predicts survival of oral squamous cell carcinoma. <i>Clinical Epigenetics</i> , 2017 , 9, 88	7.7	39
Application of cloud computing in biomedicine big data analysis cloud computing in big data 2017,		2
A seven-gene prognostic signature for rapid determination of head and neck squamous cell carcinoma survival. <i>Oncology Reports</i> , 2017 , 38, 3403-3411	3.5	21
Association of three single nucleotide polymorphisms of ESR1with breast cancer susceptibility: a meta-analysis. <i>Journal of Biomedical Research</i> , 2017 , 31, 213-225	1.5	13
Variance component testing for identifying differentially expressed genes in RNA-seq data. <i>PeerJ</i> , 2017 , 5, e3797	3.1	3
Relationship between Branched-Chain Amino Acids, Metabolic Syndrome, and Cardiovascular Risk Profile in a Chinese Population: A Cross-Sectional Study. <i>International Journal of Endocrinology</i> , 2016 , 2016, 8173905	2.7	22
A Comparative Study of Five Association Tests Based on CpG Set for Epigenome-Wide Association Studies. <i>PLoS ONE</i> , 2016 , 11, e0156895	3.7	13
Permutation-based variance component test in generalized linear mixed model with application to multilocus genetic association study. <i>BMC Medical Research Methodology</i> , 2015 , 15, 37	4.7	7
	Advanced Non-Small Cell Lung Cancer: The ALTER 0303 Phase 3 Randomized Clinical Trial. JAMA Oncology, 2018, 4, 1569-1575 Evaluating the effect of multiple genetic risk score models on colorectal cancer risk prediction. Gene, 2018, 673, 174-180 Mutant-allele fraction heterogeneity is associated with non-small cell lung cancer patient survival. Oncology Letters, 2018, 15, 795-802 Prognostic factors of refractory NSCLC patients receiving anlotinib hydrochloride as the third- or further-line treatment. Cancer Biology and Medicine, 2018, 15, 443-451 Bayesian variable selection for parametric survival model with applications to cancer omics data. Human Genomics, 2018, 12, 49 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 Profiling and Bioinformatic Analyses Indicate Differential circRNA and miRNA/isomiR Expression and Interactions. BioMed Research International, 2018, 2018, 8518563 Exome-Wide Association Study Identified New Risk Loci for Hirschsprung® Disease. Molecular Neurobiology, 2017, 54, 1777-1785 ESR2 Genetic Variants and Combined Oral Contraceptive Use Associated with the Risk of Stroke. Archives of Medical Research, 2017, 48, 203-211 A fast algorithm for Bayesian multi-locus model in genome-wide association studies. Molecular Genetics and Genomics, 2017, 292, 923-934 Seven-CpG-based prognostic signature coupled with gene expression predicts survival of oral squamous cell carcinoma. Clinical Epigenetics, 2017, 9, 88 Application of cloud computing in biomedicine big data analysis cloud computing in big data 2017, A seven-gene prognostic signature for rapid determination of head and neck squamous cell carcinoma survival. Oncology Reports, 2017, 38, 3403-3411 Association of three single nucleotide polymorphisms of ESR1with breast cancer susceptibility; a meta-analysis. Journal of Biomedical Research, 2017, 31, 213-225 Variance component testing for identifying differentially expr	Advanced Non-Small Cell Lung Cancer: The ALTER 0303 Phase 3 Randomized Clinical Trial. JAMA Oncology, 2018, 4, 1569-1575 Evaluating the effect of multiple genetic risk score models on colorectal cancer risk prediction. Gene, 2018, 673, 174-180 Mutant-allele fraction heterogeneity is associated with non-small cell lung cancer patient survival. Oncology Letters, 2018, 15, 795-802 Prognostic factors of refractory NSCLC patients receiving anlotinib hydrochloride as the third- or further-line treatment. Cancer Biology and Medicine, 2018, 15, 443-451 Bayesian variable selection for parametric survival model with applications to cancer omics data. Human Genomics, 2018, 12, 49 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 Profiling and Bioinformatic Analyses Indicate Differential circRNA and miRNA/isomiR Expression and Interactions. BioMed Research International, 2018, 2018, 8518563 Exome-Wide Association Study Identified New Risk Loci for HirschsprungB Disease. Molecular Neurobiology, 2017, 54, 1777-1785 ESSR Cenetic Variants and Combined Oral Contraceptive Use Associated with the Risk of Stroke. Archives of Medical Research, 2017, 48, 203-211 A fast algorithm for Bayesian multi-locus model in genome-wide association studies. Malecular Genetics and Genomics, 2017, 292, 923-934 Seven-CpC-based prognostic signature coupled with gene expression predicts survival of oral squamous cell carcinoma. Clinical Epigenetics, 2017, 9, 88 Application of cloud computing in biomedicine big data analysis cloud computing in big data 2017, A seven-gene prognostic signature for rapid determination of head and neck squamous cell carcinoma survival. Oncology Reports, 2017, 38, 3403-3411 Association of three single nucleotide polymorphisms of ESR with breast cancer susceptibility: a meta-analysis. Journal of Biomedical Research, 2017, 31, 213-225 Variance component testing for identifying differentially exp

20	Mitochondrial Variations in Non-Small Cell Lung Cancer (NSCLC) Survival. <i>Cancer Informatics</i> , 2015 , 14, 1-9	2.4	10
19	Computed tomography measurement of pulmonary artery for diagnosis of COPD and its comorbidity pulmonary hypertension. <i>International Journal of COPD</i> , 2015 , 10, 2525-33	3	17
18	A Systematic Evaluation of Feature Selection and Classification Algorithms Using Simulated and Real miRNA Sequencing Data. <i>Computational and Mathematical Methods in Medicine</i> , 2015 , 2015, 17857	2 ^{2.8}	6
17	Evolutionary and expression analysis of miR-#-5p and miR-#-3p at the miRNAs/isomiRs levels. BioMed Research International, 2015 , 2015, 168358	3	16
16	Comparisons of isomiR patterns and classification performance using the rank-based MANOVA and 10-fold cross-validation. <i>Gene</i> , 2015 , 569, 21-6	3.8	21
15	A coding single-nucleotide polymorphism in lysine demethylase KDM4A associates with increased sensitivity to mTOR inhibitors. <i>Cancer Discovery</i> , 2015 , 5, 245-54	24.4	20
14	Statistical analysis for genome-wide association study. <i>Journal of Biomedical Research</i> , 2015 , 29, 285-97	1.5	40
13	Variable selection approach for zero-inflated count data via adaptive lasso. <i>Journal of Applied Statistics</i> , 2014 , 41, 879-894	1	16
12	Polymorphisms in MicroRNAs are associated with survival in non-small cell lung cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014 , 23, 2503-11	4	19
11	Reconstructing the coding and non-coding RNA regulatory networks of miRNAs and mRNAs in breast cancer. <i>Gene</i> , 2014 , 548, 6-13	3.8	17
10	A genome-wide gene-environment interaction analysis for tobacco smoke and lung cancer susceptibility. <i>Carcinogenesis</i> , 2014 , 35, 1528-35	4.6	35
9	Invasive adenocarcinoma of the lung is associated with the upper lung regions. <i>Lung Cancer</i> , 2014 , 84, 145-50	5.9	21
8	Gene copy number alterations in the azoospermia-associated AZFc region and their effect on spermatogenic impairment. <i>Molecular Human Reproduction</i> , 2014 , 20, 836-43	4.4	19
7	An integrated analysis of miRNA, lncRNA, and mRNA expression profiles. <i>BioMed Research International</i> , 2014 , 2014, 345605	3	39
6	Global analysis of miRNA gene clusters and gene families reveals dynamic and coordinated expression. <i>BioMed Research International</i> , 2014 , 2014, 782490	3	12
5	Integrative analysis of miRNA-mRNA and miRNA-miRNA interactions. <i>BioMed Research International</i> , 2014 , 2014, 907420	3	36
4	Adiponectin gene polymorphisms and acute respiratory distress syndrome susceptibility and mortality. <i>PLoS ONE</i> , 2014 , 9, e89170	3.7	17
3	Rare variants detection with kernel machine learning based on likelihood ratio test. <i>PLoS ONE</i> , 2014 , 9, e93355	3.7	7

- 2 Genetic association analysis using sibship data: a multilevel model approach. PLoS ONE, 2012, 7, e31134 3.7 2
- Association between delta-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, 1986-94

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