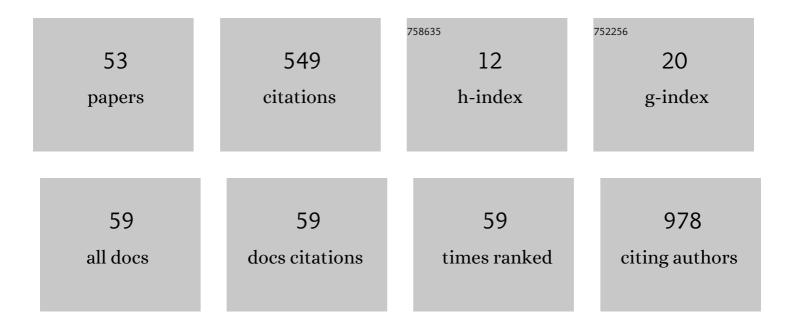
Lin Hua

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	An exploratory analysis of forme fruste keratoconus sensitivity diagnostic parameters. International Ophthalmology, 2022, 42, 2473-2481.	0.6	5
2	Low Serum LH Levels During Ovarian Stimulation With GnRH Antagonist Protocol Decrease the Live Birth Rate After Fresh Embryo Transfers but Have No Impact in Freeze-All Cycles. Frontiers in Endocrinology, 2021, 12, 640047.	1.5	16
3	Diagnostic value of galactomannan test in non-immunocompromised critically ill patients with influenza-associated aspergillosis: data from three consecutive influenza seasons. European Journal of Clinical Microbiology and Infectious Diseases, 2021, 40, 1899-1907.	1.3	4
4	Randomized controlled clinical trial of a highly filled flowable composite in non-carious cervical lesions: 3-year results. Clinical Oral Investigations, 2021, 25, 5955-5965.	1.4	7
5	Association of Venous Thromboembolism and Early Mortality in Patients with Newly Diagnosed Metastatic Non-Small Cell Lung Cancer. Cancer Management and Research, 2021, Volume 13, 4031-4040.	0.9	7
6	Impact of Angiotensin-Converting Enzyme Inhibitors and Angiotensin Receptor Blockers on the Inflammatory Response and Viral Clearance in COVID-19 Patients. Frontiers in Cardiovascular Medicine, 2021, 8, 710946.	1.1	8
7	Incidence of Type 1 Diabetes May Be Underestimated in the Chinese Population: Evidence From 21.7 Million People Between 2007 and 2017. Diabetes Care, 2021, 44, 2503-2509.	4.3	23
8	Development and validation of a predictive score for venous thromboembolism in newly diagnosed non-small cell lung cancer. Thrombosis Research, 2021, 208, 45-51.	0.8	8
9	B Lymphocyte Subsets in Children With Steroid-Sensitive Nephrotic Syndrome: A Longitudinal Study. Frontiers in Pediatrics, 2021, 9, 736341.	0.9	4
10	Decreased Circulating Transitional B-Cell to Memory B-Cell Ratio Is a Risk Factor for Relapse in Children with Steroid-Sensitive Nephrotic Syndrome. Nephron, 2021, 145, 107-112.	0.9	8
11	A Retrospective Comparative Study of 2-μm Thulium Laser During Thoracoscopic Resection of Pulmonary Nodules. Indian Journal of Surgery, 2020, 82, 1021-1025.	0.2	1
12	Early identification of patients with severe influenza-associated aspergillosis (IAA) in the intensive care unit——an IAA prediction score system (Asper-PreSS). Journal of Infection, 2020, 81, 639-646.	1.7	6
13	Integrative analysis of mRNA, miRNA and IncRNA profiles reveals the commonness between bladder cancer and breast cancer. Translational Cancer Research, 2020, 9, 1070-1090.	0.4	2
14	Frequent occurrence of CD19-negative relapse after CD19 CAR T and consolidation therapy in 14 TP53-mutated r/r B-ALL children. Leukemia, 2020, 34, 3382-3387.	3.3	35
15	The Association between Immune Infiltration and Clinical Phenotypes and Prognosis of Prostate Cancer. Iranian Journal of Biotechnology, 2020, 18, e2538.	0.3	1
16	An Exploration of Possible Association between m6A Methylation Regulators and Chronic Obstructive Pulmonary Disease by a Bioinformatics Analysis. , 2020, , .		0
17	Preliminary evidence of the association between DNAm and orbital volumetry in GO. Endocrine Connections, 2020, 9, 617-626.	0.8	2
18	Altered B-Lymphocyte Homeostasis in Idiopathic Nephrotic Syndrome. Frontiers in Pediatrics, 2019, 7, 377.	0.9	19

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19	A clinicopathological comparison between IgA nephropathy and Henoch–Schönlein purpura nephritis in children: use of the Oxford classification. Clinical and Experimental Nephrology, 2019, 23, 1382-1390.	0.7	18
20	Vitamin D deficiency is associated with risk of developing peripheral arterial disease in type 2 diabetic patients. BMC Cardiovascular Disorders, 2019, 19, 145.	0.7	14
21	Identifying and Validating Genes with DNA Methylation Data in the Context of Biological Network for Chinese Patients with Graves' Orbitopathy. International Journal of Endocrinology, 2019, 2019, 1-9.	0.6	8
22	A Pathway Analysis Based on Genome-Wide DNA Methylation of Chinese Patients with Graves' Orbitopathy. BioMed Research International, 2019, 2019, 1-10.	0.9	10
23	Helicobacter pylori infection as a risk factor for serum bilirubin change and less favourable lipid profiles: a hospital-based health examination survey. BMC Infectious Diseases, 2019, 19, 157.	1.3	14
24	The Potential Link between Gut Microbiota and Serum TRAb in Chinese Patients with Severe and Active Graves' Orbitopathy. International Journal of Endocrinology, 2019, 2019, 1-12.	0.6	16
25	The long-term efficacy of one-shot neoadjuvant intra-arterial chemotherapy combined with radical cystectomy versus radical cystectomy alone for bladder cancer: a propensity-score matching study. BMC Urology, 2019, 19, 117.	0.6	3
26	Derivation and Validation of the Potential Core Genes in Pancreatic Cancer for Tumor-Stroma Crosstalk. BioMed Research International, 2018, 2018, 1-11.	0.9	14
27	Exploration of genetics commonness between bladder cancer and breast cancer based on a silcio analysis on disease subtypes. Technology and Health Care, 2018, 26, 361-377.	0.5	3
28	New method of lung cancer detection by saliva test using surfaceâ€enhanced Raman spectroscopy. Thoracic Cancer, 2018, 9, 1556-1561.	0.8	46
29	Prediction of microRNA and gene target from an integrated network in chronic obstructive pulmonary disease based on canonical correlation analysis. Technology and Health Care, 2018, 26, 121-134.	0.5	2
30	Prevalence of diminished ovarian reserve in Chinese women with polycystic ovary syndrome and sensitive diagnostic parameters. Gynecological Endocrinology, 2017, 33, 694-697.	0.7	6
31	Novel Computational Approaches and Applications in Cancer Research. BioMed Research International, 2017, 2017, 1-2.	0.9	1
32	Reanalysis and External Validation of a Decision Tree Model for Detecting Unrecognized Diabetes in Rural Chinese Individuals. International Journal of Endocrinology, 2017, 2017, 1-6.	0.6	5
33	Gene Regulation Network Based Analysis Associated with TGF-beta Stimulation in Lung Adenocarcinoma Cells. Iranian Journal of Biotechnology, 2017, 15, 1-9.	0.3	3
34	Exploring the interaction among EPHX1, GSTP1, SERPINE2, and TGFB1 contributing to the quantitative traits of chronic obstructive pulmonary disease in Chinese Han population. Human Genomics, 2016, 10, 13.	1.4	7
35	Integration of multi-microarray datasets to identify chronic obstructive pulmonary disease-related miRNAs. Bio-Medical Materials and Engineering, 2015, 26, S1903-S1915.	0.4	3
36	ldentification of Novel Breast Cancer Subtype-Specific Biomarkers by Integrating Genomics Analysis of DNA Copy Number Aberrations and miRNA-mRNA Dual Expression Profiling. BioMed Research International, 2015, 2015, 1-17.	0.9	19

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37	Psychometric evaluation of the Simplified Chinese Version of the Posttraumatic Growth Inventory for assessing breast cancer survivors. European Journal of Oncology Nursing, 2015, 19, 391-396.	0.9	12
38	A controlled trial of transcutaneous vagus nerve stimulation for the treatment of pharmacoresistant epilepsy. Epilepsy and Behavior, 2014, 39, 105-110.	0.9	76
39	Explore the associated MiRNAs with emphysema severity of chronic obstructive pulmonary disease. , 2014, , .		0
40	Apply support vector regression to extract the potential susceptibility genes of chronic obstructive pulmonary disease. , 2014, , .		0
41	Combination of microRNA expression profiling with genome-wide SNP genotyping to construct a coronary artery disease-related miRNA-miRNA synergistic network. BioScience Trends, 2014, 8, 297-307.	1.1	23
42	The prevalence of glucose metabolism disturbances in Chinese Muslims and possible risk factors: a study from northwest China. Arquivos Brasileiros De Endocrinologia E Metabologia, 2014, 58, 715-723.	1.3	3
43	Clinical observation of local resection or enucleation for uveal melanoma. Chinese Medical Journal, 2014, 127, 3459-63.	0.9	1
44	Wnt Signaling Pathway-Based Analysis of Variance on Different Subtypes of Breast. , 2013, , .		0
45	Applying Bioinformatics Methods to Detect the Relationship among Complex Diseases. , 2013, , .		0
46	Prioritizing breast cancer subtype related miRNAs using miRNA–mRNA dysregulated relationships extracted from their dual expression profiling. Journal of Theoretical Biology, 2013, 331, 1-11.	0.8	15
47	Identifying Breast Cancer Subtype Related miRNAs from Two Constructed miRNAs Interaction Networks in Silico Method. BioMed Research International, 2013, 2013, 1-13.	0.9	8
48	A comparison of contributions to disease phenotype between damaging and benign non-synonymous SNPs. , 2012, , .		0
49	Explore coronary artery disease related microRNA clusters by combing single nucleotide polymorphisms with microRNA microarray. , 2012, , .		0
50	Mining Functional Gene Modules Linked with Rheumatoid Arthritis Using a SNP-SNP Network. Genomics, Proteomics and Bioinformatics, 2012, 10, 23-34.	3.0	13
51	Identifying obesity indicators which best correlate with type 2 diabetes in a Chinese population. BMC Public Health, 2012, 12, 732.	1.2	24
52	Genome-wide gene-gene interaction analysis of rheumatoid arthritis. , 2010, , .		0
53	A simple tool detected diabetes and prediabetes in rural Chinese. Journal of Clinical Epidemiology, 2010, 63, 1030-1035.	2.4	22