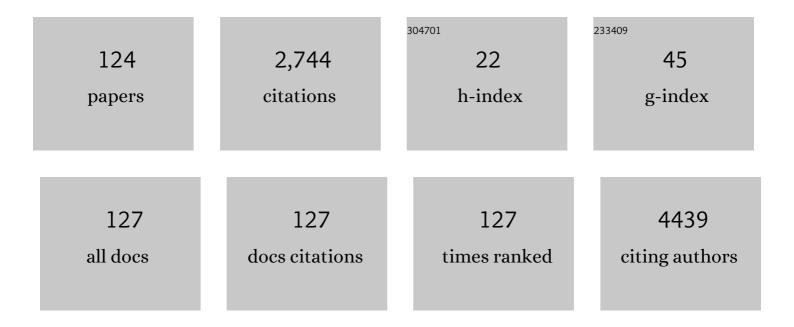
List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Circulating tumour DNA methylation markers for diagnosis and prognosis of hepatocellular carcinoma. Nature Materials, 2017, 16, 1155-1161.	27.5	641
2	Safety and feasibility of CRISPR-edited T cells in patients with refractory non-small-cell lung cancer. Nature Medicine, 2020, 26, 732-740.	30.7	322
3	Genomic monitoring of SARS-CoV-2 uncovers an Nsp1 deletion variant that modulates type I interferon response. Cell Host and Microbe, 2021, 29, 489-502.e8.	11.0	95
4	Homogeneous Visual and Fluorescence Detection of Circulating Tumor Cells in Clinical Samples <i>via</i> Selective Recognition Reaction and Enzyme-Free Amplification. ACS Nano, 2021, 15, 11634-11643.	14.6	81
5	Quercetin inhibits IL-1 beta-induced ICAM-1 expression in pulmonary epithelial cell line A549 through the MAPK pathways. Molecular Biology Reports, 2009, 36, 1825-1832.	2.3	73
6	Association of polymorphisms in the human IL-10 and IL-18 genes with rheumatoid arthritis. Molecular Biology Reports, 2011, 38, 379-385.	2.3	70
7	Integrating exosomal microRNAs and electronic health data improved tuberculosis diagnosis. EBioMedicine, 2019, 40, 564-573.	6.1	53
8	A proline deletion in IFNAR1 impairs IFN-signaling and underlies increased resistance to tuberculosis in humans. Nature Communications, 2018, 9, 85.	12.8	49
9	Detection of Circulating Tumor Cells in Breast Cancer Patients by Nanopore Sensing with Aptamer-Mediated Amplification. ACS Sensors, 2020, 5, 2359-2366.	7.8	43
10	International Expansion of a Novel SARS-CoV-2 Mutant. Journal of Virology, 2020, 94, .	3.4	42
11	Multimode MicroRNA Sensing via Multiple Enzyme-Free Signal Amplification and Cation-Exchange Reaction. ACS Applied Materials & Interfaces, 2019, 11, 36476-36484.	8.0	41
12	Homogeneous Binary Visual and Fluorescence Detection of Tetanus Toxoid in Clinical Samples Based on Enzyme-Free Parallel Hybrid Chain Reaction. Nano Letters, 2022, 22, 1710-1717.	9.1	37
13	Ghrelin inhibit cell apoptosis in pancreatic β cell line HIT-T15 via mitogen-activated protein kinase/phosphoinositide 3-kinase pathways. Toxicology, 2007, 237, 194-202.	4.2	35
14	MiR-124-3p inhibits the migration and invasion of Gastric cancer by targeting ITGB3. Pathology Research and Practice, 2020, 216, 152762.	2.3	35
15	The diagnostic and prognostic role of myocardial injury biomarkers in hospitalized patients with COVID-19. Clinica Chimica Acta, 2020, 510, 186-190.	1.1	35
16	A precision medicine approach to managing 2019 novel coronavirus pneumonia. Precision Clinical Medicine, 2020, 3, 14-21.	3.3	34
17	Association between SNPs in microRNA machinery genes and gastric cancer susceptibility, invasion, and metastasis in Chinese Han population. Oncotarget, 2017, 8, 86435-86446.	1.8	30
18	A finger-driven disposable micro-platform based on isothermal amplification for the application of multiplexed and point-of-care diagnosis of tuberculosis. Biosensors and Bioelectronics, 2022, 195, 113663.	10.1	29

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19	Translation of aptamers toward clinical diagnosis and commercialization. Biosensors and Bioelectronics, 2022, 208, 114168.	10.1	29
20	Epidemiology study of HBV genotypes and antiviral drug resistance in multi-ethnic regions from Western China. Scientific Reports, 2015, 5, 17413.	3.3	27
21	Significance of genetic polymorphisms in long non-coding RNA AC079767.4 in tuberculosis susceptibility and clinical phenotype in Western Chinese Han population. Scientific Reports, 2017, 7, 965.	3.3	27
22	Active DNA unwinding and transport by a membrane-adapted helicase nanopore. Nature Communications, 2019, 10, 5083.	12.8	25
23	Epidemiological surveillance of common respiratory viruses in patients with suspected COVID-19 in Southwest China. BMC Infectious Diseases, 2020, 20, 688.	2.9	25
24	Elevated circulating PAI-1 levels are related to lung function decline, systemic inflammation, and small airway obstruction in chronic obstructive pulmonary disease. International Journal of COPD, 2016, Volume 11, 2369-2376.	2.3	24
25	CRISPR-Cas13a cascade-based viral RNA assay for detecting SARS-CoV-2 and its mutations in clinical samples. Sensors and Actuators B: Chemical, 2022, 362, 131765.	7.8	23
26	Antituberculosis Drugâ€Induced Adverse Events in the Liver, Kidneys, and Blood: Clinical Profiles and Pharmacogenetic Predictors. Clinical Pharmacology and Therapeutics, 2018, 104, 326-334.	4.7	22
27	Systematic Evolution of Ligands by Exponential Enrichment Technologies and Aptamer-Based Applications: Recent Progress and Challenges in Precision Medicine of Infectious Diseases. Frontiers in Bioengineering and Biotechnology, 2021, 9, 704077.	4.1	22
28	Association of the gene polymorphisms in sodium taurocholate cotransporting polypeptide with the outcomes of hepatitis B infection in Chinese Han population. Infection, Genetics and Evolution, 2014, 27, 77-82.	2.3	20
29	Ultrasensitive Nanopore Sensing of Mucin 1 and Circulating Tumor Cells in Whole Blood of Breast Cancer Patients by Analyte-Triggered Triplex-DNA Release. ACS Applied Materials & Interfaces, 2021, 13, 21030-21039.	8.0	20
30	Visual and dual-fluorescence homogeneous sensor for the detection of pyrophosphatase in clinical hyperthyroidism samples based on selective recognition of CdTe QDs and coordination polymerization of Ce ³⁺ . Journal of Materials Chemistry C, 2021, 9, 4141-4149.	5.5	19
31	Long Noncoding RNA and Predictive Model To Improve Diagnosis of Clinically Diagnosed Pulmonary Tuberculosis. Journal of Clinical Microbiology, 2020, 58, .	3.9	18
32	Clinical significance of Inc-AC145676.2.1-6 and Inc-TGS1-1 and their variants in western Chinese tuberculosis patients. International Journal of Infectious Diseases, 2019, 84, 8-14.	3.3	17
33	Applications of laboratory findings in the prevention, diagnosis, treatment, and monitoring of COVID-19. Signal Transduction and Targeted Therapy, 2021, 6, 316.	17.1	17
34	Nearâ€Infraredâ€Enpowered Nanomotorâ€Mediated Targeted Chemotherapy and Mitochondrial Phototherapy to Boost Systematic Antitumor Immunity. Advanced Healthcare Materials, 2022, 11, e2200255.	7.6	17
35	Ghrelin Inhibits Interleukin-6 Production Induced by Cigarette Smoke Extract in the Bronchial Epithelial Cell Via NF-κB Pathway. Inflammation, 2016, 39, 190-198.	3.8	16
36	Exonuclease III-assisted strand displacement reaction-driven cyclic generation of G-quadruplex strategy for homogeneous fluorescent detection of melamine. Talanta, 2019, 203, 255-260.	5.5	16

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37	<p>The diagnostic value of circulating tumor cells and ctDNA for gene mutations in lung cancer</p> . OncoTargets and Therapy, 2019, Volume 12, 2539-2552.	2.0	16
38	Importance of common TLR2 genetic variants on clinical phenotypes and risk in tuberculosis disease in a Western Chinese population. Infection, Genetics and Evolution, 2018, 60, 173-180.	2.3	15
39	Genetic polymorphisms of long noncoding RNA <i>RP11â€37B2.1</i> associate with susceptibility of tuberculosis and adverse events of antituberculosis drugs in west China. Journal of Clinical Laboratory Analysis, 2019, 33, e22880.	2.1	15
40	Homogeneous two-dimensional visual and fluorescence analysis of circulating tumor cells in clinical samples via steric hindrance regulated enzymes recognition cleavage and elongation. Biosensors and Bioelectronics, 2022, 202, 114009.	10.1	15
41	Simultaneous Homogeneous Fluorescence Detection of AFP and GPC3 in Hepatocellular Carcinoma Clinical Samples Assisted by Enzyme-Free Catalytic Hairpin Assembly. ACS Applied Materials & Interfaces, 2022, 14, 28697-28705.	8.0	15
42	Clinical features, Outcomes and Molecular Profiles of Drug Resistance in Tuberculous Meningitis in non-HIV Patients. Scientific Reports, 2016, 6, 19072.	3.3	14
43	Status of drug-resistant tuberculosis in China: A systematic review and meta-analysis. American Journal of Infection Control, 2016, 44, 671-676.	2.3	14
44	Genetic polymorphisms in PXR and NF-κB1 influence susceptibility to anti-tuberculosis drug-induced liver injury. PLoS ONE, 2019, 14, e0222033.	2.5	14
45	Polymorphisms in sodium taurocholate cotransporting polypeptide are not associated with hepatitis B virus clearance in Chinese Tibetans and Uygurs. Infection, Genetics and Evolution, 2016, 41, 128-134.	2.3	13
46	A Novel Genetic Variation in <i>NCF2</i> , the Core Component of NADPH Oxidase, Contributes to the Susceptibility of Tuberculosis in Western Chinese Han Population. DNA and Cell Biology, 2020, 39, 57-62.	1.9	13
47	Autoimmune hemolytic anemia in hospitalized patients. Medicine (United States), 2020, 99, e18739.	1.0	13
48	Low-Cost and Scalable Platform with Multiplexed Microwell Array Biochip for Rapid Diagnosis of COVID-19. Research, 2021, 2021, 2813643.	5.7	13
49	Association between the European GWAS-Identified Susceptibility Locus at Chromosome 4p16 and the Risk of Atrial Septal Defect: A Case-Control Study in Southwest China and a Meta-Analysis. PLoS ONE, 2015, 10, e0123959.	2.5	13
50	SARS-CoV-2 nucleic acid testing is China's key pillar of COVID-19 containment. Lancet, The, 2022, 399, 1690-1691.	13.7	13
51	Pathway Analyses Identify Novel Variants in the WNT Signaling Pathway Associated with Tuberculosis in Chinese Population. Scientific Reports, 2016, 6, 28530.	3.3	12
52	Allele frequency distribution of 21 forensic autosomal STR loci of Goldeneyeâ,,¢ DNA ID 22NC Kit in Chinese Tibetan group. Forensic Science International: Genetics, 2016, 22, e21-e24.	3.1	12
53	Sequence variations of mitochondrial DNA D‑loop region in patients with acute myeloid leukemia. Oncology Letters, 2017, 14, 6269-6276.	1.8	12
54	A general strategy for label-free homogeneous bioassays based on selective recognition and silver ion-mediated conformational switch. Talanta, 2019, 201, 9-15.	5.5	12

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55	Mechanical strain-induced c-fos expression in pulmonary epithelial cell line A549. Biochemical and Biophysical Research Communications, 2006, 347, 369-372.	2.1	11
56	Association between SNPs in miRNA-machinery genes and chronic hepatitis B in the Chinese Han population. Infection, Genetics and Evolution, 2014, 28, 113-117.	2.3	11
57	The prevalence and clinical profiles of FLT3-ITD, FLT3-TKD, NPM1, C-KIT, DNMT3A, and CEBPA mutations in a cohort of patients with de novo acute myeloid leukemia from southwest China. Tumor Biology, 2016, 37, 7357-7370.	1.8	11
58	SFRP1 variations influence susceptibility and immune response to Mycobacterium tuberculosis in a Chinese Han population. Infection, Genetics and Evolution, 2016, 37, 259-265.	2.3	11
59	Plasma levels of ILâ€l Ra are associated with schizophrenia. Psychiatry and Clinical Neurosciences, 2018, 73, 109-115.	1.8	11
60	Development of aÂfluorescent DNA nanomachine for ultrasensitive detection of Salmonella enteritidis without labeling and enzymes. Mikrochimica Acta, 2020, 187, 376.	5.0	11
61	Development and Validation of a Predictive Model for Severe COVID-19: A Case-Control Study in China. Frontiers in Medicine, 2021, 8, 663145.	2.6	11
62	The long non-coding RNA HOXA11-AS activates ITGB3 expression to promote the migration and invasion of gastric cancer by sponging miR-124-3p. Cancer Cell International, 2021, 21, 576.	4.1	11
63	Genetic Variation in miR-146a Is Not Associated with Susceptibility to IgA Nephropathy in Adults from a Chinese Han Population. PLoS ONE, 2015, 10, e0139554.	2.5	10
64	Clinical relevance of the Incâ€HNF1Bâ€3:1 genetic polymorphisms in Western Chinese tuberculosis patients. Journal of Clinical Laboratory Analysis, 2020, 34, e23076.	2.1	10
65	Development and Validation of a LASSO Prediction Model for Better Identification of Ischemic Stroke: A Case-Control Study in China. Frontiers in Aging Neuroscience, 2021, 13, 630437.	3.4	10
66	Multi-responsive nanotheranostics with enhanced tumor penetration and oxygen self-producing capacities for multimodal synergistic cancer therapy. Acta Pharmaceutica Sinica B, 2022, 12, 406-423.	12.0	10
67	Blockade of macrophage-associated programmed death 1 inhibits the pyroptosis signalling pathway in sepsis. Inflammation Research, 2021, 70, 993-1004.	4.0	10
68	TLR1 polymorphisms are significantly associated with the occurrence, presentation and drug-adverse reactions of tuberculosis in Western Chinese adults. Oncotarget, 2018, 9, 1691-1704.	1.8	10
69	A Notch4 missense mutation is associated with susceptibility to tuberculosis in Chinese population. Infection, Genetics and Evolution, 2020, 78, 104145.	2.3	9
70	A novel risk factor for predicting anti-tuberculosis drug resistance in patients with tuberculosis complicated with type 2 diabetes mellitus. International Journal of Infectious Diseases, 2020, 97, 69-77.	3.3	9
71	Systematic evaluation, verification and comparison of tuberculosisâ€related nonâ€coding RNA diagnostic panels. Journal of Cellular and Molecular Medicine, 2021, 25, 184-202.	3.6	9
72	Association of Genetic Variants in Wnt Signaling Pathway with Tuberculosis in Chinese Han Population. PLoS ONE, 2014, 9, e93841.	2.5	8

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73	Clinical relevance of <i>LINC00152</i> and its variants in western Chinese tuberculosis patients. Oncotarget, 2017, 8, 115456-115468.	1.8	8
74	Ultrasensitive DNA Methylation Ratio Detection Based on the Target-Induced Nanoparticle-Coupling and Site-Specific Base Oxidation Damage for Colorectal Cancer. Analytical Chemistry, 2022, 94, 6261-6270.	6.5	8
75	RIPK2 polymorphisms and susceptibility to tuberculosis in a Western Chinese Han population. Infection, Genetics and Evolution, 2019, 75, 103950.	2.3	7
76	Association of ABCC Gene Polymorphism With Susceptibility to Antituberculosis Drug–Induced Hepatotoxicity in Western Han Patients With Tuberculosis. Journal of Clinical Pharmacology, 2020, 60, 361-368.	2.0	7
77	Homogeneous assay based on the pre-reduction and selective cation exchange for detection of multiple targets by atomic spectrometry. Talanta, 2020, 219, 121387.	5.5	7
78	A Promising Preoperative Prediction Model for Microvascular Invasion in Hepatocellular Carcinoma Based on an Extreme Gradient Boosting Algorithm. Frontiers in Oncology, 2022, 12, 852736.	2.8	7
79	Identification of Abnormal 51 CTA/CTG Expansion as Probably the Shortest Pathogenic Allele for Spinocerebellar Ataxia-8 in China. Neuroscience Bulletin, 2018, 34, 859-862.	2.9	6
80	Multimode detection of β-glycosidase and pathogenic bacteria via cation exchange assisted signal amplification. Mikrochimica Acta, 2020, 187, 453.	5.0	6
81	Significance of LncRNA CASC8 genetic polymorphisms on the tuberculosis susceptibility in Chinese population. Journal of Clinical Laboratory Analysis, 2020, 34, e23234.	2.1	6
82	Which sample type is better for Xpert MTB/RIF to diagnose adult and pediatric pulmonary tuberculosis?. Bioscience Reports, 2020, 40, .	2.4	6
83	Deciphering a TB-related DNA methylation biomarker and constructing a TB diagnostic classifier. Molecular Therapy - Nucleic Acids, 2022, 27, 37-49.	5.1	6
84	Novel directions of precision oncology: circulating microbial DNA emerging in cancer-microbiome areas. Precision Clinical Medicine, 2022, 5, .	3.3	6
85	Association of LEPR polymorphisms with predisposition and inflammatory response in anti-tuberculosis drug-induced liver injury: A pilot prospective investigation in Western Chinese Han population. Infection, Genetics and Evolution, 2019, 75, 103970.	2.3	5
86	C3-Epimer of 25-Hydroxyvitamin D3 as a Superior Marker for Predicting the Severity of Chronic Kidney Disease in Rheumatoid Arthritis Patients. Oxidative Medicine and Cellular Longevity, 2022, 2022, 1-8.	4.0	5
87	Detection of nucleic acids via G-quadruplex-controlled l-cysteine oxidation and catalyzed hairpin assembly-assisted signal amplification. RSC Advances, 2018, 8, 40564-40569.	3.6	4
88	The Variant at TGFBRAP1 but Not TGFBR2 Is Associated with Antituberculosis Drug-Induced Liver Injury. Evidence-based Complementary and Alternative Medicine, 2019, 2019, 1-9.	1.2	4
89	<i>ARID5B</i> Genetic Polymorphisms Contribute to the Susceptibility and Prognosis of Male Acute Promyelocytic Leukemia. DNA and Cell Biology, 2019, 38, 1374-1386.	1.9	4
90	Interleukinâ€⊋1 receptor gene polymorphism is associated with hepatitis B virusâ€related hepatocellular carcinoma in Chinese patients. Journal of Clinical Laboratory Analysis, 2019, 33, e22860.	2.1	4

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91	The genetic variants in calcium signaling related genes influence anti-tuberculosis drug induced liver injury. Medicine (United States), 2019, 98, e17821.	1.0	4
92	Correlation between IncRNA AC079767.4 variants and liver injury from antituberculosis treatment in West China. Journal of Infection and Chemotherapy, 2020, 26, 63-68.	1.7	4
93	Association of POR and PPARα polymorphisms with risk of anti-tuberculosis drug-induced liver injury in Western Chinese Han population. Infection, Genetics and Evolution, 2020, 79, 104147.	2.3	4
94	Validations of Top and Novel Susceptibility Variants in All-Age Chinese Patients With Acute Lymphoblastic Leukemia. Frontiers in Genetics, 2020, 11, 1004.	2.3	4
95	No Significant Effects of IL-6 and IL-13 Gene Variants on Tuberculosis Susceptibility in the Chinese Population. DNA and Cell Biology, 2020, 39, 1356-1367.	1.9	4
96	Up-regulated long noncoding RNA AC007128.1 and its genetic polymorphisms associated with Tuberculosis susceptibility. Annals of Translational Medicine, 2021, 9, 1018-1018.	1.7	4
97	Metal–Organic Framework-Derived Hollow and Hierarchical Porous Multivariate Metal-Oxide Microspheres for Efficient Phosphoproteomics Analysis. ACS Applied Materials & Interfaces, 2021, 13, 34762-34772.	8.0	4
98	Fluorescence Aptasensor of Tuberculosis Interferon-Î ³ in Clinical Samples Regulated by Steric Hindrance and Selective Identification. Analytical Chemistry, 2022, 94, 9122-9129.	6.5	4
99	A prospective study on associations between superoxide dismutase gene polymorphisms and antituberculosis drugâ€induced liver injury in a Chinese Han population. Journal of Gene Medicine, 2019, 21, e3121.	2.8	3
100	The dominant model analysis of Sirt3 genetic variants is associated with susceptibility to tuberculosis in a Chinese Han population. Molecular Genetics and Genomics, 2020, 295, 1155-1162.	2.1	3
101	Association of IL27 and STAT3 genetic polymorphism on the susceptibility of tuberculosis in Western Chinese Han population. Infection, Genetics and Evolution, 2020, 83, 104324.	2.3	3
102	Novel Long Non-coding RNA and LASSO Prediction Model to Better Identify Pulmonary Tuberculosis: A Case-Control Study in China. Frontiers in Molecular Biosciences, 2021, 8, 632185.	3.5	3
103	Cancer Genomic Alterations Can Be Potential Biomarkers Predicting Microvascular Invasion and Early Recurrence of Hepatocellular Carcinoma. Frontiers in Oncology, 2022, 12, 783109.	2.8	3
104	Fluorescence and visual immunoassay of HIV-1 p24 antigen in clinical samples via multiple selective recognitions of CdTe QDs. Mikrochimica Acta, 2021, 188, 422.	5.0	3
105	Genetic Profile of Two Tibetan Populations from China by Analysis of 15 STR Loci. Human Biology, 2006, 78, 757-770.	0.2	2
106	Evaluation of three rapid assays for Mycobacterium tuberculosis complex detection in a comprehensive hospital from West China. Clinical Biochemistry, 2015, 48, 79-84.	1.9	2
107	Do genetic polymorphisms of B-cell CLL/lymphoma 2 confer susceptibility to anti-tuberculous therapy-associated drug-induced liver injury?. International Journal of Infectious Diseases, 2020, 91, 223-231.	3.3	2
108	Analysis of Factors Influencing Kidney Function of Recipients After Renal Transplantation in Southwestern China: A Retrospective Study. Frontiers in Medicine, 2020, 7, 519582.	2.6	2

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109	Collagen type XVIII alpha 1 chain (COL18A1) variants affect the risk of antiâ€ŧuberculosis drugâ€induced hepatotoxicity: A prospective study. Journal of Clinical Laboratory Analysis, 2021, 35, e23630.	2.1	2
110	Age-related differences of genetic susceptibility to patients with acute lymphoblastic leukemia. Aging, 2021, 13, 12456-12465.	3.1	2
111	A novel nucleotide deletion in <i>RHAG</i> allele identified in a Chinese Rh _{null} individual. Transfusion, 2018, 58, 826-827.	1.6	1
112	Genetic polymorphisms of 21 autosomal STR loci in Chinese Uygur ethnic population. International Journal of Legal Medicine, 2018, 132, 1637-1639.	2.2	1
113	Weak A variant caused by <i>c.940A>G</i> missense mutation of the ABO gene. Transfusion, 2019, 59, E13-E14.	1.6	1
114	The risk factors for tuberculosis patients with diabetes mellitus living in Western China: a retrospective study conducted from 2014 to 2018. International Journal of Diabetes in Developing Countries, 2020, 40, 538-546.	0.8	1
115	Absence of significant association between <i>UGT2B4</i> genetic variants and the susceptibility to antiâ€tuberculosis drugâ€induced liver injury in a Western Chinese population. Journal of Clinical Pharmacy and Therapeutics, 2021, 46, 66-73.	1.5	1
116	Assessing the role of SH3RF1 and SH3RF2 polymorphisms in susceptibility to tuberculosis: A case-control study in the Han Chinese population. Microbial Pathogenesis, 2021, 152, 104567.	2.9	1
117	Potential Fluid Biomarkers and a Prediction Model for Better Recognition Between Multiple System Atrophy-Cerebellar Type and Spinocerebellar Ataxia. Frontiers in Aging Neuroscience, 2021, 13, 644699.	3.4	1
118	Genetic and Functional Evaluation of the Role of FOXO1 in Antituberculosis Drug-Induced Hepatotoxicity. Evidence-based Complementary and Alternative Medicine, 2021, 2021, 1-13.	1.2	1
119	A case-control study on correlation between the single nucleotide polymorphism of CLEC4E and the susceptibility to tuberculosis among Han people in Western China. BMC Infectious Diseases, 2021, 21, 788.	2.9	1
120	Association analysis of pulmonary tuberculosis and vitamin D Receptor Gene Polymorphisms of Han population in Western China. Microbial Pathogenesis, 2021, 161, 105190.	2.9	1
121	Chinese abnormal compound heterozygote spinocerebellar ataxia type 8: a case report. Neurological Sciences, 2022, 43, 1435-1439.	1.9	1
122	Commentary: Epidemiology of Antibody-Positive Autoimmune Encephalitis in Southwest China: A Multicenter Study. Frontiers in Immunology, 2020, 11, 1976.	4.8	0
123	Genetic Polymorphisms of Delta-Like 1 Homolog Influence the Susceptibility to Antituberculosis Drug-Induced Hepatotoxicity. DNA and Cell Biology, 2021, 40, 231-238.	1.9	Ο
124	Allele frequency and proportion defined by circulating tumor DNA profiling predict tyrosine kinase inhibitors' therapeutic outcomes for non-small cell lung cancer. Journal of Cancer Research and Clinical Oncology, 2022, , .	2.5	0