Xiangyu

List of Publications by Year
in descending order

43 papers	1,624 citations	394286 19 h-index	315616 38 g-index
43 all docs	43 docs citations	43 times ranked	3445 citing authors

#	Article	IF	Citations
1	Short- and long-term prognosis of glycemic control in COVID-19 patients with type 2 diabetes. QJM - Monthly Journal of the Association of Physicians, 2022, 115, 131-139.	0.2	17
2	OUP accepted manuscript. QJM - Monthly Journal of the Association of Physicians, 2022, , .	0.2	1
3	Post-sequelae one year after hospital discharge among older COVID-19 patients: A multi-center prospective cohort study. Journal of Infection, 2022, 84, 179-186.	1.7	31
4	Development and validation of a prognostic nomogram for predicting in-hospital mortality of COVID-19: a multicenter retrospective cohort study of 4086 cases in China. Aging, 2021, 13, 3176-3189.	1.4	14
5	Singleâ€nucleotide polymorphism rs4142441 and MYC coâ€modulated long nonâ€coding RNA OSER1â€AS1 suppresses nonâ€small cell lung cancer by sequestering ELAVL1. Cancer Science, 2021, 112, 2272-2286.	1.7	17
6	The Association Between Short-Term Air Pollution Exposure and Post-Adolescent Acne: The Evidence from a Time Series Analysis in Xiâ€™an, China. Clinical, Cosmetic and Investigational Dermatology, 2021, Volume 14, 723-731.	0.8	3
7	Symptoms and Health Outcomes Among Survivors of COVID-19 Infection 1 Year After Discharge From Hospitals in Wuhan, China. JAMA Network Open, 2021, 4, e2127403.	2.8	146
8	Circulating Long Noncoding RNAs Act as Diagnostic Biomarkers in Non-Small Cell Lung Cancer. Frontiers in Oncology, 2020, 10, 537120.	1.3	17
9	Mendelian randomization analyses of genetically predicted circulating levels of cytokines with risk of breast cancer. Npj Precision Oncology, 2020, 4, 25.	2.3	18
10	Systematic meta-analyses, field synopsis and global assessment of the evidence of genetic association studies in colorectal cancer. Gut, 2020, 69, 1460-1471.	6.1	27
11	Epidemiological, comorbidity factors with severity and prognosis of COVID-19: a systematic review and meta-analysis. Aging, 2020, 12, 12493-12503.	1.4	298
12	Bioinformatics analyses and biological function of lncRNA ZFPM2â€AS1 and ZFPM2 gene in hepatocellular carcinoma. Oncology Letters, 2020, 19, 3677-3686.	0.8	6
13	Consumption of Very Low Mineral Water Is Associated with Lower Bone Mineral Content in Children. Journal of Nutrition, 2019, 149, 1994-2000.	1.3	18
14	Burnout among obstetricians and paediatricians: a cross-sectional study from China. BMJ Open, 2019, 9, e024205.	0.8	14
15	Hypoxiaâ€sensitive <scp>LINC</scp>01436 is regulated by E2F6 and acts as an oncogene by targeting miRâ€30aâ€3p in nonâ€small cell lung cancer. Molecular Oncology, 2019, 13, 840-856.	2.1	47
16	Genetic predisposition to lung cancer: comprehensive literature integration, meta-analysis, and multiple evidence assessment of candidate-gene association studies. Scientific Reports, 2017, 7, 8371.	1.6	87
17	GPC5, a novel epigenetically silenced tumor suppressor, inhibits tumor growth by suppressing Wnt/Î²-catenin signaling in lung adenocarcinoma. Oncogene, 2016, 35, 6120-6131.	2.6	86
18	Whole-exome sequencing identifies OR2W3 mutation as a cause of autosomal dominant retinitis pigmentosa. Scientific Reports, 2015, 5, 9236.	1.6	25

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19	Relationship between *CYP11B2*-344T>C polymorphsim and atrial fibrillation: A meta-analysis. JRAAS - Journal of the Renin-Angiotensin-Aldosterone System, 2015, 16, 185-188.	1.0	4
20	Association of Hemostatic Markers with Atrial Fibrillation: A Meta-Analysis and Meta-Regression. PLoS ONE, 2015, 10, e0124716.	1.1	32
21	Liver enzymes and metabolic syndrome: a large-scale case-control study. Oncotarget, 2015, 6, 26782-26788.	0.8	25
22	A Community-Based Comprehensive Intervention Program for 7200 Patients with Type 2 Diabetes Mellitus in Chongqing (China). International Journal of Environmental Research and Public Health, 2014, 11, 11450-11463.	1.2	10
23	Complete genome sequence of 285P, a novel T7-like polyvalent E. coli bacteriophage. Virus Genes, 2014, 48, 528-533.	0.7	5
24	Genetic variants associated with colorectal cancer risk: comprehensive research synopsis, meta-analysis, and epidemiological evidence. Gut, 2014, 63, 326-336.	6.1	101
25	Effectiveness of evidence-based medicine training for undergraduate students at a Chinese Military Medical University: a self-controlled trial. BMC Medical Education, 2014, 14, 133.	1.0	9
26	No association between genetic variants in angiogenesis and inflammation pathway genes and breast cancer survival among Chinese women. Cancer Epidemiology, 2013, 37, 619-624.	0.8	8
27	Association of inflammatory factors with occurrence and recurrence of atrial fibrillation: A meta-analysis. International Journal of Cardiology, 2013, 169, 62-72.	0.8	187
28	Impact: China needs to review its metrics. Nature, 2013, 503, 198-198.	13.7	1
29	No Association between Ovarian Cancer Susceptibility Variants and Breast Cancer Risk among Chinese Women. Cancer Epidemiology Biomarkers and Prevention, 2013, 22, 467-469.	1.1	1
30	Anthropometric measures and epithelial ovarian cancer risk among Chinese women: results from the Shanghai Women's Health Study. British Journal of Cancer, 2013, 109, 751-755.	2.9	30
31	Pathway Analyses Identify *TGFBR2* as Potential Breast Cancer Susceptibility Gene: Results from a Consortium Study among Asians. Cancer Epidemiology Biomarkers and Prevention, 2012, 21, 1176-1184.	1.1	22
32	Glycemic load, glycemic index and risk of cardiovascular diseases: Meta-analyses of prospective studies. Atherosclerosis, 2012, 223, 491-496.	0.4	95
33	Associations of the ATP-binding cassette transporter A1 R219K polymorphism with HDL-C level and coronary artery disease risk: A meta-analysis. Atherosclerosis, 2011, 215, 428-434.	0.4	41
34	Association between gene polymorphisms of vitamin D receptor and pulmonary tuberculosis susceptibility: a meta-analysis. Journal of Medical Colleges of PLA, 2011, 26, 63-75.	0.1	4
35	Association between CYP19 polymorphisms and breast cancer risk: results from 10,592 cases and 11,720 controls. Breast Cancer Research and Treatment, 2010, 122, 495-501.	1.1	40
36	No association between SOD2 Val16Ala polymorphism and breast cancer susceptibility: a meta-analysis based on 9,710 cases and 11,041 controls. Breast Cancer Research and Treatment, 2010, 122, 509-514.	1.1	17

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37	Methylenetetrahydrofolate reductase polymorphisms and breast cancer risk: a meta-analysis from 41 studies with 16,480 cases and 22,388 controls. Breast Cancer Research and Treatment, 2010, 123, 499-506.	1.1	54
38	Extremely low-frequency electromagnetic fields exposure and female breast cancer risk: a meta-analysis based on 24,338 cases and 60,628 controls. Breast Cancer Research and Treatment, 2010, 123, 569-576.	1.1	30
39	Transforming growth factorÎ²1 L10P variant plays an active role on the breast cancer susceptibility in Caucasian: evidence from 10,392 cases and 11,697 controls. Breast Cancer Research and Treatment, 2010, 124, 453-457.	1.1	20
40	Molecular characterization, structural analysis and determination of host range of a novel bacteriophage LSB-1. Virology Journal, 2010, 7, 255.	1.4	4
41	Molecular Characterization of Tb, a New Approach for an Ancient Brucellaphage. International Journal of Molecular Sciences, 2009, 10, 2999-3011.	1.8	7
42	Isolation of a novel polyvalent virulent bacteriophage of E. coli. Journal of Medical Colleges of PLA, 2007, 22, 261-267.	0.1	4
43	Association between interleukin-4 polymorphisms and asthma. Journal of Medical Colleges of PLA, 2007, 22, 312-319.	0.1	1