List of Publications by Year in descending order

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YUE LIANC

#	Article	IF	CITATIONS
1	Plasma mi <scp>RNA</scp> s as early biomarkers for detecting hepatocellular carcinoma. International Journal of Cancer, 2015, 137, 1679-1690.	5.1	188
2	Identification of risk loci and a polygenic risk score for lung cancer: a large-scale prospective cohort study in Chinese populations. Lancet Respiratory Medicine,the, 2019, 7, 881-891.	10.7	167
3	Genomic Landscape Survey Identifies SRSF1 as a Key Oncodriver in Small Cell Lung Cancer. PLoS Genetics, 2016, 12, e1005895.	3.5	144
4	Systematic identification of genes with a cancer-testis expression pattern in 19 cancer types. Nature Communications, 2016, 7, 10499.	12.8	124
5	Genetic risk, incident gastric cancer, and healthy lifestyle: a meta-analysis of genome-wide association studies and prospective cohort study. Lancet Oncology, The, 2020, 21, 1378-1386.	10.7	123
6	Associations Between Hepatitis B Virus Infection and Risk of All Cancer Types. JAMA Network Open, 2019, 2, e195718.	5.9	114
7	Identification of new susceptibility loci for gastric non-cardia adenocarcinoma: pooled results from two Chinese genome-wide association studies. Gut, 2017, 66, 581-587.	12.1	68
8	Whole-genome sequencing reveals genomic signatures associated with the inflammatory microenvironments in Chinese NSCLC patients. Nature Communications, 2018, 9, 2054.	12.8	68
9	Estimation of heritability for nine common cancers using data from genomeâ€wide association studies in Chinese population. International Journal of Cancer, 2017, 140, 329-336.	5.1	66
10	Comprehensive analyses of m6A regulators and interactive coding and non-coding RNAs across 32 cancer types. Molecular Cancer, 2021, 20, 67.	19.2	65
11	Association analysis identifies new risk loci for non-obstructive azoospermia in Chinese men. Nature Communications, 2014, 5, 3857.	12.8	64
12	Plasma metabolomics identified novel metabolites associated with risk of type 2 diabetes in two prospective cohorts of Chinese adults. International Journal of Epidemiology, 2016, 45, 1507-1516.	1.9	64
13	A cancer-testis non-coding RNA LIN28B-AS1 activates driver gene LIN28B by interacting with IGF2BP1 in lung adenocarcinoma. Oncogene, 2019, 38, 1611-1624.	5.9	61
14	Identification of Novel T1D Risk Loci and Their Association With Age and Islet Function at Diagnosis in Autoantibody-Positive T1D Individuals: Based on a Two-Stage Genome-Wide Association Study. Diabetes Care, 2019, 42, 1414-1421.	8.6	60
15	Exome Array Analysis Identifies Variants in SPOCD1 and BTN3A2 That Affect Risk for Gastric Cancer. Gastroenterology, 2017, 152, 2011-2021.	1.3	58
16	Evaluation of genetic variants in microRNA biosynthesis genes and risk of breast cancer in Chinese women. International Journal of Cancer, 2013, 133, 2216-2224.	5.1	50
17	Genetic variation in a hsa-let-7 binding site in RAD52 is associated with breast cancer susceptibility. Carcinogenesis, 2013, 34, 689-693.	2.8	43
18	Meta-analysis of genome-wide association studies and functional assays decipher susceptibility genes for gastric cancer in Chinese populations. Gut, 2020, 69, 641-651.	12.1	36

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19	Combinational therapeutic targeting of BRD4 and CDK7 synergistically induces anticancer effects in head and neck squamous cell carcinoma. Cancer Letters, 2020, 469, 510-523.	7.2	36
20	Hepatitis B virus genotype, mutations, human leukocyte antigen polymorphisms and their interactions in hepatocellular carcinoma: a multi-centre case-control study. Scientific Reports, 2015, 5, 16489.	3.3	35
21	A Screen for Genomic Disorders of Infertility Identifies MAST2 Duplications Associated with Nonobstructive Azoospermia in Humans1. Biology of Reproduction, 2015, 93, 61.	2.7	30
22	Genetic Variants at 1p11.2 and Breast Cancer Risk: A Two-Stage Study in Chinese Women. PLoS ONE, 2011, 6, e21563.	2.5	24
23	Evaluation of genetic variants in autophagy pathway genes as prognostic biomarkers for breast cancer. Gene, 2017, 627, 549-555.	2.2	21
24	Comprehensive characterization of functional eRNAs in lung adenocarcinoma reveals novel regulators and a prognosis-related molecular subtype. Theranostics, 2020, 10, 11264-11277.	10.0	20
25	Genetic Variations in the Flanking Regions of miR-101-2 Are Associated with Increased Risk of Breast Cancer. PLoS ONE, 2014, 9, e86319.	2.5	19
26	Immune landscape and subtypes in primary resectable oral squamous cell carcinoma: prognostic significance and predictive of therapeutic response. , 2021, 9, e002434.		19
27	Association of KCTD10, MVK, and MMAB polymorphisms with dyslipidemia and coronary heart disease in Han Chinese population. Lipids in Health and Disease, 2016, 15, 171.	3.0	18
28	Circulating Tumor Cells Enriched by the Depletion of Leukocytes with Bi-Antibodies in Non-Small Cell Lung Cancer: Potential Clinical Application. PLoS ONE, 2015, 10, e0137076.	2.5	18
29	Predictive value of prognostic nutritional index in patients with oral squamous cell carcinoma. Oral Diseases, 2020, 26, 903-911.	3.0	17
30	Potentially functional polymorphisms in aminoacyl-tRNA synthetases genes are associated with breast cancer risk in a Chinese population. Molecular Carcinogenesis, 2015, 54, 577-583.	2.7	15
31	Evaluation of CpG-SNPs in miRNA promoters and risk of breast cancer. Gene, 2018, 651, 1-8.	2.2	15
32	Genetic variants at 9p21.3 are associated with risk of esophageal squamous cell carcinoma in a Chinese population. Cancer Science, 2017, 108, 250-255.	3.9	14
33	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
34	Epigenetic–smoking interaction reveals histologically heterogeneous effects of TRIM27 DNA methylation on overall survival among earlyâ€stage NSCLC patients. Molecular Oncology, 2020, 14, 2759-2774.	4.6	13
35	Functional genetic variants in centrosome-related genes CEP72 and YWHAG confer susceptibility to gastric cancer. Archives of Toxicology, 2020, 94, 2861-2872.	4.2	13
36	Associations Between Sleep Quality and Health Span: A Prospective Cohort Study Based on 328,850 UK Biobank Participants. Frontiers in Genetics, 2021, 12, 663449.	2.3	13

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37	Low-frequency germline variants across 6p22.2–6p21.33 are associated with non-obstructive azoospermia in Han Chinese men. Human Molecular Genetics, 2015, 24, 5628-5636.	2.9	12
38	Polymorphisms in CARS are associated with gastric cancer risk: a two-stage case–control study in the Chinese population. Gastric Cancer, 2017, 20, 940-947.	5.3	12
39	Lowâ€frequency nonsynonymous variants in <i>FKBPL</i> and <i>ARPC1B</i> genes are associated with breast cancer risk in Chinese women. Molecular Carcinogenesis, 2017, 56, 774-780.	2.7	12
40	Claudin-7 Inhibits Proliferation and Metastasis in Salivary Adenoid Cystic Carcinoma Through Wnt/β-Catenin Signaling. Cell Transplantation, 2020, 29, 096368972094358.	2.5	12
41	Potentially functional variants in IncRNAs are associated with breast cancer risk in a Chinese population. Molecular Carcinogenesis, 2017, 56, 2048-2057.	2.7	11
42	Integrating of genomic and transcriptomic profiles for the prognostic assessment of breast cancer. Breast Cancer Research and Treatment, 2019, 175, 691-699.	2.5	11
43	Genetic characterisation of sarcomatoid carcinomas reveals multiple novel actionable mutations and identifies <i>KRAS</i> mutation as a biomarker of poor prognosis. Journal of Medical Genetics, 2022, 59, 10-17.	3.2	11
44	Epidemiological, clinical, and 3-dimentional CBCT radiographic characterizations of supernumerary teeth in a non-syndromic adult population: a single-institutional study from 60,104 Chinese subjects. Clinical Oral Investigations, 2020, 24, 4271-4281.	3.0	11
45	Epidemiological, clinical, radiographic characterization of nonâ€syndromic supernumerary teeth in Chinese children and adolescents. Oral Diseases, 2021, 27, 981-992.	3.0	11
46	U-shaped association between telomere length and esophageal squamous cell carcinoma risk: a case-control study in Chinese population. Frontiers of Medicine, 2015, 9, 478-486.	3.4	10
47	Leukocyte telomere length, lipid parameters and gestational diabetes risk: a case-control study in a Chinese population. Scientific Reports, 2019, 9, 8483.	3.3	10
48	ldentification of an autophagyâ€related prognostic signature in head and neck squamous cell carcinoma. Journal of Oral Pathology and Medicine, 2021, 50, 1040-1049.	2.7	10
49	Early Second-Trimester Serum MicroRNAs as Potential Biomarker for Nondiabetic Macrosomia. BioMed Research International, 2014, 2014, 1-6.	1.9	8
50	A cis-eQTL genetic variant of the cancer–testis gene CCDC116 is associated with risk of multiple cancers. Human Genetics, 2017, 136, 987-997.	3.8	7
51	Genetic variants affecting telomere length are associated with the prognosis of esophageal squamous cell carcinoma in a Chinese population. Molecular Carcinogenesis, 2017, 56, 1021-1029.	2.7	7
52	Genetic Variants in the Promoter Region of <i> miR-10b</i> and the Risk of Breast Cancer. BioMed Research International, 2017, 2017, 1-7.	1.9	7
53	Genetic Variation in the 3'-Untranslated Region of NBN Gene Is Associated with Gastric Cancer Risk in a Chinese Population. PLoS ONE, 2015, 10, e0139059.	2.5	7
54	Peptidomic analysis of zebrafish embryos exposed to polychlorinated biphenyls and their impact on eye development. Ecotoxicology and Environmental Safety, 2019, 175, 164-172.	6.0	6

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55	The sonic hedgehog signaling pathway is suppressed following PCB ₁₂₅₄ exposure during retinal development. Environmental Toxicology, 2019, 34, 340-347.	4.0	6
56	Transcriptomeâ \in wide association study for persistent hepatitis B virus infection and related hepatocellular carcinoma. Liver International, 2020, 40, 2117-2127.	3.9	6
57	Functional Dissection of CD26 and Its Pharmacological Inhibition by Sitagliptin During Skin Wound Healing. Medical Science Monitor, 2021, 27, e928933.	1.1	5
58	Genetic variants at 10q23.33 are associated with plasma lipid levels in a Chinese population. Journal of Biomedical Research, 2014, 28, 53-8.	1.6	5
59	Weight Change across Adulthood in Relation to Non-Alcoholic Fatty Liver Disease among Non-Obese Individuals. Nutrients, 2022, 14, 2140.	4.1	5
60	Identification of Recurrent Variants in BRCA1 and BRCA2 across Multiple Cancers in the Chinese Population. BioMed Research International, 2020, 2020, 1-11.	1.9	4
61	Sex-Specific Associations of Testosterone and Genetic Factors With Health Span. Frontiers in Endocrinology, 2021, 12, 773464.	3.5	4
62	Pharmacological inhibition of CDK7 by THZ1 impairs tumor growth in p53â€mutated HNSCC. Oral Diseases, 2022, 28, 611-620.	3.0	3
63	Investigation of Causal Effect of Type 2 Diabetes Mellitus on Lung Cancer: A Mendelian Randomization Study. Frontiers in Genetics, 2021, 12, 673687.	2.3	3
64	The rs6771157 C/G polymorphism in SCN10A is associated with the risk of atrial fibrillation in a Chinese Han population. Scientific Reports, 2016, 6, 35212.	3.3	2
65	Prognostic role of CD11b+ myeloid-derived suppressor cells in oral squamous cell carcinoma. Archives of Medical Science, 2021, , .	0.9	2
66	Functional characterization and clinical significance of superâ€enhancers in lung adenocarcinoma. Molecular Carcinogenesis, 2022, 61, 776-786.	2.7	2
67	Genetic Variations in miR-30 Family Member Regulatory Regions Are Associated with Breast Cancer Risk in a Chinese Population. BioMed Research International, 2020, 2020, 1-8.	1.9	1
68	The lab-confirmed interval of COVID-19 clusters and its application in the strength evaluation of prevention and control measures. BMC Infectious Diseases, 2021, 21, 226.	2.9	1
69	Potential functional variants of KIAA genes are associated with breast cancer risk in a case control study. Annals of Translational Medicine, 2021, 9, 549-549.	1.7	1
70	Coding variants in the PCNT and CEP295 genes contribute to breast cancer risk in Chinese women. Pathology Research and Practice, 2021, 225, 153581.	2.3	1