Na Shen

List of Publications by Year
in descending order

80	1,179	430754	501076
papers	citations	18 h-index	28 g-index

82	82	82	2274
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Exome-wide analyses identify low-frequency variant in CYP26B1 and additional coding variants associated with esophageal squamous cell carcinoma. Nature Genetics, 2018, 50, 338-343.	9.4	75
2	Clinical characteristics of 80 hospitalized frontline medical workers infected with COVID-19 in Wuhan, China. Journal of Hospital Infection, 2020, 105, 399-403.	1.4	64
3	Non-linear doseâ€“response relationship between cigarette smoking and pancreatic cancer risk: Evidence from a meta-analysis of 42 observational studies. European Journal of Cancer, 2014, 50, 193-203.	1.3	63
4	A Rare Missense Variant in TCF7L2 Associates with Colorectal Cancer Risk by Interacting with a GWAS-Identified Regulatory Variant in the MYC Enhancer. Cancer Research, 2018, 78, 5164-5172.	0.4	54
5	Dietary Mushroom Intake May Reduce the Risk of Breast Cancer: Evidence from a Meta-Analysis of Observational Studies. PLoS ONE, 2014, 9, e93437.	1.1	40
6	TPX2 promotes migration and invasion of human breast cancer cells. Asian Pacific Journal of Tropical Medicine, 2015, 8, 1064-1070.	0.4	35
7	Genetic variants in the SWI/SNF complex and smoking collaborate to modify the risk of pancreatic cancer in a Chinese population. Molecular Carcinogenesis, 2015, 54, 761-768.	1.3	35
8	A Genetic Variant rs1801274 in FCGR2A as a Potential Risk Marker for Kawasaki Disease: A Case-Control Study and Meta-Analysis. PLoS ONE, 2014, 9, e103329.	1.1	32
9	Exome-wide analysis identifies three low-frequency missense variants associated with pancreatic cancer risk in Chinese populations. Nature Communications, 2018, 9, 3688.	5.8	32
10	Systematic Confirmation Study of GWAS-Identified Genetic Variants for Kawasaki Disease in A Chinese Population. Scientific Reports, 2015, 5, 8194.	1.6	31
11	Meta-analysis Reveals the Prognostic Value of Circulating Tumour Cells Detected in the Peripheral Blood in Patients with Non-Metastatic Colorectal Cancer. Scientific Reports, 2017, 7, 905.	1.6	30
12	The significant prognostic value of circulating tumor cells in triple-negative breast cancer: a meta-analysis. Oncotarget, 2016, 7, 37361-37369.	0.8	27
13	Urinary bisphenol A and its interaction with ESR1 genetic polymorphism associated with non-small cell lung cancer: findings from a case-control study in Chinese population. Chemosphere, 2020, 254, 126835.	4.2	26
14	Genetic Variant in MTRR, but Not MTR, Is Associated with Risk of Congenital Heart Disease: An Integrated Meta-Analysis. PLoS ONE, 2014, 9, e89609.	1.1	24
15	Leukemia cell-derived microvesicles induce T cell exhaustion via miRNA delivery. OncoImmunology, 2018, 7, e1448330.	2.1	24
16	Association between rs2853669 in TERT gene and the risk and prognosis of human cancer: a systematic review and meta-analysis. Oncotarget, 2017, 8, 50864-50872.	0.8	24
17	Genetic variant in SWI/SNF complexes influences hepatocellular carcinoma risk: a new clue for the contribution of chromatin remodeling in carcinogenesis. Scientific Reports, 2014, 4, 4147.	1.6	23
18	Association between the p.V37I variant of GJB2 and hearing loss: a pedigree and meta-analysis. Oncotarget, 2017, 8, 46681-46690.	0.8	21

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19	MAD1L1 Arg558His and MAD2L1 Leu84Met interaction with smoking increase the risk of colorectal cancer. Scientific Reports, 2015, 5, 12202.	1.6	20
20	Parity and pancreatic cancer risk: evidence from a meta-analysis of twenty epidemiologic studies. Scientific Reports, 2014, 4, 5313.	1.6	20
21	Evaluating the diagnostic accuracy of the Xpert MTB/RIF assay on bronchoalveolar lavage fluid: A retrospective study. International Journal of Infectious Diseases, 2018, 71, 14-19.	1.5	20
22	SF3A1 and pancreatic cancer: new evidence for the association of the spliceosome and cancer. Oncotarget, 2015, 6, 37750-37757.	0.8	20
23	Clinicopathological features of breast cancer with different molecular subtypes in chinese women. Journal of Huazhong University of Science and Technology [Medical Sciences], 2013, 33, 117-121.	1.0	19
24	Long nonâ€'coding RNA nuclear paraspeckle assembly transcript 1 interacts with microRNAâ€'107 to modulate breast cancer growth and metastasis by targeting carnitine palmitoyltransferaseâ€'1. International Journal of Oncology, 2019, 55, 1125-1136.	1.4	18
25	Association of lncRNA H19 rs217727 polymorphism and cancer risk in the Chinese population: a meta-analysis. Oncotarget, 2016, 7, 59580-59588.	0.8	17
26	Construction of a Risk Prediction Model for Subsequent Bloodstream Infection in Intestinal Carriers of Carbapenem-Resistant Enterobacteriaceae: A Retrospective Study in Hematology Department and Intensive Care Unit. Infection and Drug Resistance, 2021, Volume 14, 815-824.	1.1	16
27	The Roles of Ca2+/NFAT Signaling Genes in Kawasaki Disease: Single- and Multiple-Risk Genetic Variants. Scientific Reports, 2014, 4, 5208.	1.6	15
28	CpG-methylation-based risk score predicts progression in colorectal cancer. Epigenomics, 2020, 12, 605-615.	1.0	15
29	Vitamin D status in Mainland of China: A systematic review and meta-analysis. EClinicalMedicine, 2021, 38, 101017.	3.2	15
30	Sleep Duration and the Risk of Fatty Liver Disease: A Systematic Review and Meta-analysis. Scientific Reports, 2016, 6, 31956.	1.6	14
31	Vitamin D receptor polymorphisms and the susceptibility of Parkinsonâ€™s disease. Neuroscience Letters, 2019, 699, 206-211.	1.0	13
32	LINC01149 variant modulates MICA expression that facilitates hepatitis B virus spontaneous recovery but increases hepatocellular carcinoma risk. Oncogene, 2020, 39, 1944-1956.	2.6	13
33	Alpha and beta-Thalassemia mutations in Hubei area of China. BMC Medical Genetics, 2020, 21, 6.	2.1	12
34	8p22â€“23-rs2254546 as a Susceptibility Locus for Kawasaki Disease: a Case-control Study and a Meta-analysis. Scientific Reports, 2014, 4, 4247.	1.6	11
35	Influence of tumor extent on central lymph node metastasis in solitary papillary thyroid microcarcinomas: a retrospective study of 1092 patients. World Journal of Surgical Oncology, 2017, 15, 133.	0.8	11
36	Bisphenol A exposure, interaction with genetic variants and colorectal cancer via mediating oxidative stress biomarkers. Environmental Pollution, 2021, 287, 117630.	3.7	11

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37	Genome-wide gene-bisphenol A, F and triclosan interaction analyses on urinary oxidative stress markers. Science of the Total Environment, 2022, 807, 150753.	3.9	11
38	Hypermethylation of the SEPT9 Gene Suggests Significantly Poor Prognosis in Cancer Patients: A Systematic Review and Meta-Analysis. Frontiers in Genetics, 2019, 10, 887.	1.1	10
39	Hepatocellular carcinoma risk variant modulates lncRNA HLA-DQB1-AS1 expression via a long-range enhancerâ€“promoter interaction. Carcinogenesis, 2021, 42, 1347-1356.	1.3	10
40	Integrative Genomic Analysis Identifies That SERPINA6-rs1998056 Regulated by FOXA/ERÎ± Is Associated with Female Hepatocellular Carcinoma. PLoS ONE, 2014, 9, e107246.	1.1	9
41	Identification of a novel de novo ANK1 R1426* nonsense mutation in a Chinese family with hereditary spherocytosis by NGS. Oncotarget, 2017, 8, 96791-96797.	0.8	9
42	Novel hereditary spherocytosis-associated splice site mutation in the ANK1 gene caused by parental gonosomal mosaicism. Haematologica, 2018, 103, e219-e222.	1.7	9
43	T cells expressing CD26-specific chimeric antigen receptors exhibit extensive self-antigen-driven fratricide. Immunopharmacology and Immunotoxicology, 2019, 41, 490-496.	1.1	9
44	Functional polymorphisms on chromosome 5p15.33 disturb telomere biology and confer the risk of nonâ€small cell lung cancer in Chinese population. Molecular Carcinogenesis, 2019, 58, 913-921.	1.3	9
45	A missense variant in PTPN12 associated with the risk of colorectal cancer by modifying Ras/MEK/ERK signaling. Cancer Epidemiology, 2019, 59, 109-114.	0.8	9
46	A functional variant rs1537373 in 9p21.3 region is associated with pancreatic cancer risk. Molecular Carcinogenesis, 2019, 58, 760-766.	1.3	9
47	Mutation analysis of a Chinese family with oculocutaneous albinism. Oncotarget, 2016, 7, 84981-84988.	0.8	8
48	SLC10A1 S267F variant influences susceptibility to HBV infection and reduces cholesterol level by impairing bile acid uptake. Journal of Viral Hepatitis, 2019, 26, 1178-1185.	1.0	8
49	Tumor necrosis factor α knockout impaired tumorigenesis in chronic myeloid leukemia cells partly by metabolism modification and miRNA regulation . OncoTargets and Therapy, 2019, Volume 12, 2355-2364.	1.0	8
50	Whole-exome sequencing identifies a novel missense variant within LOXHD1 causing rare hearing loss in a Chinese family. BMC Medical Genetics, 2019, 20, 30.	2.1	8
51	The interaction of tumor cells and myeloid-derived suppressor cells in chronic myelogenous leukemia. Leukemia and Lymphoma, 2020, 61, 128-137.	0.6	8
52	Urinary bisphenol A and its interaction with CYP17A1 rs743572 are associated with breast cancer risk. Chemosphere, 2022, 286, 131880.	4.2	8
53	Ruxolitinib add-on in corticosteroid-ref ractory graft-vs-host disease after allogeneic stem cell transplantation: Results from a retrospective study on 38 Chinese patients. World Journal of Clinical Cases, 2020, 8, 1065-1073.	0.3	8
54	A Phosphorylation-Related Variant ADD1-rs4963 Modifies the Risk of Colorectal Cancer. PLoS ONE, 2015, 10, e0121485.	1.1	8

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55	Successful treatment of a kidney transplant patient with COVID-19 and late-onset Pneumocystis jirovecii pneumonia. Annals of Clinical Microbiology and Antimicrobials, 2021, 20, 83.	1.7	8
56	Exacerbation of ichthyosis vulgaris phenotype by co-inheritance of STS and FLG mutations in a Chinese family with ichthyosis: a case report. BMC Medical Genetics, 2018, 19, 120.	2.1	7
57	An ANK1 IVS3-2A>C mutation causes exon 4 skipping in two patients from a Chinese family with hereditary spherocytosis. Oncotarget, 2017, 8, 113282-113286.	0.8	6
58	A folate receptor 3 SNP promotes mitochondriaâ€induced clonogenicity of CML leukemia cells: Implications for treatment free remission. Clinical and Translational Medicine, 2021, 11, e317.	1.7	6
59	Harnessing Big Data to Optimize an Algorithm for Rapid Diagnosis of Pulmonary Tuberculosis in a Real-World Setting. Frontiers in Cellular and Infection Microbiology, 2021, 11, 650163.	1.8	6
60	Association of RAGE rs1800625 Polymorphism and Cancer Risk: A Meta-Analysis of 18 Case-Control Studies. Medical Science Monitor, 2019, 25, 7026-7034.	0.5	6
61	Possible association of CCDC62 rs12817488 polymorphism and Parkinsonâ€™s disease risk in Chinese population: a meta-analysis. Scientific Reports, 2016, 6, 23991.	1.6	5
62	A de novo deletion mutation in SOX10 in a Chinese family with Waardenburg syndrome type 4. Scientific Reports, 2017, 7, 41513.	1.6	5
63	MBOAT7-TMC4 rs641738 Is Not Associated With the Risk of Hepatocellular Carcinoma or Persistent Hepatitis B Infection. Frontiers in Oncology, 2021, 11, 639438.	1.3	5
64	Hereditary elliptocytosis with variable expression and incomplete penetrance in a Chinese family. British Journal of Haematology, 2019, 186, e159-e162.	1.2	3
65	CDC25B is associated with the risk of hepatocellular carcinoma, but not related to persistent infection of hepatitis B virus in a Chinese population. Molecular Biology Reports, 2020, 47, 3361-3368.	1.0	3
66	Mutational analysis of a Chinese family with oculocutaneous albinism type 2. Oncotarget, 2017, 8, 70345-70355.	0.8	3
67	The epigenetic effect of microRNA in BCR-ABL1-positive microvesicles during the transformation of normal hematopoietic transplants. Oncology Reports, 2017, 38, 3278-3284.	1.2	2
68	Early genetic testing of STK11 is important for management and genetic counseling for Peutzâ€“Jeghers syndrome. Digestive and Liver Disease, 2019, 51, 1353-1355.	0.4	2
69	Mutation of Factor IX Cys178 is intolerant and may cause severe hemophilia B. Thrombosis Research, 2019, 183, 108-110.	0.8	2
70	TERT-rs33963617 and CLPTM1L-rs77518573 reduce the risk of non-small cell lung cancer in Chinese population. Gene, 2020, 731, 144357.	1.0	2
71	PNPLA3 rs738409 is not associated with the risk of hepatocellular carcinoma and persistent infection of hepatitis B virus (HBV) in HBV-related subjects: A case-control study and meta-analysis on Asians. Gene, 2020, 742, 144585.	1.0	2
72	Congenital fibrinogen disorder caused by digenic mutations of the FGA and FGB genes. Hematology, 2020, 25, 145-148.	0.7	2

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73	The experimental research on the inhibiting impacts of RNAi on Cyclin E in breast cancer cell line. Chinese-German Journal of Clinical Oncology, 2011, 10, 502-505.	0.1	1
74	Identification of a Homozygous Missense Mutation in the TYR Gene in a Chinese Family with OCA1. Current Medical Science, 2018, 38, 932-936.	0.7	1
75	ATM rs189037 significantly increases the risk of cancer in non-smokers rather than smokers: an updated meta-analysis. Bioscience Reports, 2019, 39, .	1.1	1
76	Monitoring and Analysis of Chinese Chronic Myeloid Leukemia Patients Who Have Stopped Tyrosine Kinase Inhibitor Therapy. Current Medical Science, 2019, 39, 211-216.	0.7	1
77	F8 IVS9+5G>A mutation causes moderate haemophilia A. Haemophilia, 2019, 25, e132-e135.	1.0	1
78	Nonreceptor protein tyrosine phosphatases (NRPTPs) gene family associates with the risk of hepatocellular carcinoma in a Chinese hepatitis B virusâ€related subjects. Molecular Carcinogenesis, 2020, 59, 980-988.	1.3	1
79	Normal activated partial thromboplastin time in Chinese patients with mild hemophilia B. Hematology, 2020, 25, 484-488.	0.7	0
80	The Interaction of Tumor Cells and Myeloid-Derived Suppressor Cells in Chronic Myelogenous Leukemia. Blood, 2019, 134, 1636-1636.	0.6	0