Yuanqing Ye

List of Publications by Year
in descending order

100 papers	5,069 citations	136950 32 h-index	102487 66 g-index
103 all docs	103 docs citations	103 times ranked	11788 citing authors

#	Article	IF	Citations
1	Human ribonuclease 1 serves as a secretory ligand of ephrin A4 receptor and induces breast tumor initiation. Nature Communications, 2021, 12, 2788.	12.8	11
2	Allostatic score and its associations with demographics, healthy behaviors, tumor characteristics, and mitochondrial DNA among breast cancer patients. Breast Cancer Research and Treatment, 2021, 187, 587-596.	2.5	21
3	Land use mix and leukocyte telomere length in Mexican Americans. Scientific Reports, 2021, 11, 19742.	3.3	1
4	HIF3A DNA methylation, obesity and weight gain, and breast cancer risk among Mexican American women. Obesity Research and Clinical Practice, 2020, 14, 548-553.	1.8	11
5	Genetic associations of T cell cancer immune response-related genes with T cell phenotypes and clinical outcomes of early-stage lung cancer. , 2020, 8, e000336.		9
6	Validation of plasma metabolites associated with breast cancer risk among Mexican Americans. Cancer Epidemiology, 2020, 69, 101826.	1.9	1
7	Leukocyte telomere length associated with glioma risk and survival. Aging and Cancer, 2020, 1, 71-78.	1.6	0
8	Genetic variants in epithelialâ€“mesenchymal transition genes as predictors of clinical outcomes in localized prostate cancer. Carcinogenesis, 2020, 41, 1057-1064.	2.8	0
9	Patterns of racial/ethnic disparities in baseline health-related quality of life and relationship with overall survival in patients with colorectal cancer. Quality of Life Research, 2020, 29, 2977-2986.	3.1	10
10	Comprehensive T cell repertoire characterization of non-small cell lung cancer. Nature Communications, 2020, 11, 603.	12.8	140
11	Unruptured intracranial aneurysm growth trajectory: occurrence and rate of enlargement in 520 longitudinally followed cases. Journal of Neurosurgery, 2020, 132, 1077-1087.	1.6	17
12	Metabolic hormones and breast cancer risk among Mexican American Women in the Mano a Mano Cohort Study. Scientific Reports, 2019, 9, 9989.	3.3	10
13	Sex specific associations in genome wide association analysis of renal cell carcinoma. European Journal of Human Genetics, 2019, 27, 1589-1598.	2.8	27
14	Elevated Platelet Count Appears to Be Causally Associated with Increased Risk of Lung Cancer: A Mendelian Randomization Analysis. Cancer Epidemiology Biomarkers and Prevention, 2019, 28, 935-942.	2.5	21
15	Nanotrap-enabled quantification of KRAS-induced peptide hydroxylation in blood for cancer early detection. Nano Research, 2019, 12, 1445-1452.	10.4	5
16	Breast cancer risk in relation to plasma metabolites among Hispanic and African American women. Breast Cancer Research and Treatment, 2019, 176, 687-696.	2.5	13
17	Soluble immune checkpoint-related proteins as predictors of tumor recurrence, survival, and T cell phenotypes in clear cell renal cell carcinoma patients. , 2019, 7, 334.		107
18	The influence of obesity-related factors in the etiology of renal cell carcinomaâ€”A mendelian randomization study. PLoS Medicine, 2019, 16, e1002724.	8.4	59

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19	A 5-microRNA signature identified from serum microRNA profiling predicts survival in patients with advanced stage non-small cell lung cancer. Carcinogenesis, 2019, 40, 643-650.	2.8	52
20	Genetic associations of T cell cancer immune response with tumor aggressiveness in localized prostate cancer patients and disease reclassification in an active surveillance cohort. OncoImmunology, 2019, 8, e1483303.	4.6	7
21	Reply to â€˜Mosaic loss of chromosome Y in leukocytes mattersâ€™. Nature Genetics, 2019, 51, 7-9.	21.4	7
22	Is folic acid safe for nonâ€“muscle-invasive bladder cancer patients? An evidence-based cohort study. American Journal of Clinical Nutrition, 2018, 107, 208-216.	4.7	19
23	Genetic variants in cytokine signaling pathways and clinical outcomes in early-stage lung cancer patients. Journal of Thoracic and Cardiovascular Surgery, 2018, 155, 2635-2645.e15.	0.8	5
24	Germline genetic variants in somatically significantly mutated genes in tumors are associated with renal cell carcinoma risk and outcome. Carcinogenesis, 2018, 39, 752-757.	2.8	18
25	Determinants and prognostic value of quality of life in patients with pancreatic ductal adenocarcinoma. European Journal of Cancer, 2018, 92, 20-32.	2.8	21
26	Cancer risk associated with chronic diseases and disease markers: prospective cohort study. BMJ: British Medical Journal, 2018, 360, k134.	2.3	97
27	Phase I study of nab-paclitaxel, gemcitabine, and bevacizumab in patients with advanced cancers. British Journal of Cancer, 2018, 118, 1419-1424.	6.4	7
28	Angiogenin/Ribonuclease 5 Is an EGFR Ligand and a Serum Biomarker for Erlotinib Sensitivity in Pancreatic Cancer. Cancer Cell, 2018, 33, 752-769.e8.	16.8	58
29	The somatic mutation landscape of premalignant colorectal adenoma. Gut, 2018, 67, 1299-1305.	12.1	52
30	Novel genetic variants in the P38MAPK pathway gene ZAK and susceptibility to lung cancer. Molecular Carcinogenesis, 2018, 57, 216-224.	2.7	9
31	Serum MicroRNAâ€150 Predicts Prognosis for Earlyâ€Stage Nonâ€Small Cell Lung Cancer and Promotes Tumor Cell Proliferation by Targeting Tumor Suppressor Gene SRCIN1. Clinical Pharmacology and Therapeutics, 2018, 103, 1061-1073.	4.7	31
32	Prevalence of Aflatoxin-Associated TP53R249S Mutation in Hepatocellular Carcinoma in Hispanics in South Texas. Cancer Prevention Research, 2018, 11, 103-112.	1.5	19
33	Dietary patterns and risk of recurrence and progression in nonâ€muscleâ€invasive bladder cancer. International Journal of Cancer, 2018, 142, 1797-1804.	5.1	23
34	Serum miR-331-3p predicts tumor recurrence in esophageal adenocarcinoma. Scientific Reports, 2018, 8, 14006.	3.3	26
35	Global and Targeted miRNA Expression Profiling in Clear Cell Renal Cell Carcinoma Tissues Potentially Links miR-155-5p and miR-210-3p to both Tumorigenesis and Recurrence. American Journal of Pathology, 2018, 188, 2487-2496.	3.8	34
36	Socio-demographic, Clinical, and Genetic Determinants of Quality of Life in Lung Cancer Patients. Scientific Reports, 2018, 8, 10640.	3.3	16

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37	Identification of susceptibility pathways for the role of chromosome 15q25.1 in modifying lung cancer risk. Nature Communications, 2018, 9, 3221.	12.8	60
38	Serum microRNAs as predictors of risk for non-muscle invasive bladder cancer. Oncotarget, 2018, 9, 14895-14908.	1.8	11
39	Associations of blood mitochondrial DNA copy number with social-demographics and cancer risk: results from the Mano-A-Mano Mexican American Cohort. Oncotarget, 2018, 9, 25491-25502.	1.8	6
40	Hypoxia-targeted gold nanorods for cancer photothermal therapy. Oncotarget, 2018, 9, 26556-26571.	1.8	24
41	Cohort Profile: The Mexican American Mano a Mano Cohort. International Journal of Epidemiology, 2017, 46, e3-e3.	1.9	28
42	D-mannose: a Novel Prognostic Biomarker for Patients with Esophageal Adenocarcinoma. Carcinogenesis, 2017, 38, bgw207.	2.8	19
43	Genetic Variants in Epigenetic Pathways and Risks of Multiple Cancers in the GAME-ON Consortium. Cancer Epidemiology Biomarkers and Prevention, 2017, 26, 816-825.	2.5	10
44	Genetic variants of PTPN2 are associated with lung cancer risk: a re-analysis of eight GWASs in the TRICL-ILCCO consortium. Scientific Reports, 2017, 7, 825.	3.3	10
45	Personalized Prognostic Prediction Models for Breast Cancer Recurrence and Survival Incorporating Multidimensional Data. Journal of the National Cancer Institute, 2017, 109, .	6.3	42
46	Effect of physiological factors on the biochemical properties of colon tissue â€“ an in vivo Raman spectroscopy study. Journal of Raman Spectroscopy, 2017, 48, 902-909.	2.5	13
47	Potential Susceptibility Loci Identified for Renal Cell Carcinoma by Targeting Obesity-Related Genes. Cancer Epidemiology Biomarkers and Prevention, 2017, 26, 1436-1442.	2.5	2
48	Genome-wide association study identifies multiple risk loci for renal cell carcinoma. Nature Communications, 2017, 8, 15724.	12.8	106
49	Large-scale association analysis identifies new lung cancer susceptibility loci and heterogeneity in genetic susceptibility across histological subtypes. Nature Genetics, 2017, 49, 1126-1132.	21.4	472
50	Functional variants in DCAF4 associated with lung cancer risk in European populations. Carcinogenesis, 2017, 38, 541-551.	2.8	16
51	High baseline levels of interleukin-8 in leukocytes and urine predict tumor recurrence in non-muscle invasive bladder cancer patients receiving bacillus Calmetteâ€“Guerin therapy: A long-term survival analysis. OncoImmunology, 2017, 6, e1265719.	4.6	18
52	Associations between genetic variants in mRNA splicing-related genes and risk of lung cancer: a pathway-based analysis from published GWASs. Scientific Reports, 2017, 7, 44634.	3.3	10
53	Polymorphisms in genes related to epithelialâ€“mesenchymal transition and risk of non-small cell lung cancer. Carcinogenesis, 2017, 38, 1029-1035.	2.8	18
54	Genetic Variants Related to Longer Telomere Length are Associated with Increased Risk of Renal Cell Carcinoma. European Urology, 2017, 72, 747-754.	1.9	39

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55	Circulating metabolite profiles to predict overall survival in advanced non-small cell lung cancer patients receiving first-line chemotherapy. Lung Cancer, 2017, 114, 70-78.	2.0	15
56	Global and targeted serum metabolic profiling of colorectal cancer progression. Cancer, 2017, 123, 4066-4074.	4.1	51
57	Susceptibility loci of CNOT6 in the general mRNA degradation pathway and lung cancer riskâ€"A reâ€analysis of eight GWASs. Molecular Carcinogenesis, 2017, 56, 1227-1238.	2.7	10
58	Predictors of health-related quality of life and association with survival may identify colorectal cancer patients at high risk of poor prognosis. Quality of Life Research, 2017, 26, 319-330.	3.1	21
59	Response. Journal of the National Cancer Institute, 2017, 109, .	6.3	0
60	Common, germline genetic variations in the novel tumor suppressor BAP1 and risk of developing different types of cancer. Oncotarget, 2017, 8, 74936-74946.	1.8	15
61	Social-demographics, health behaviors, and telomere length in the Mexican American Mano a Mano Cohort. Oncotarget, 2017, 8, 96553-96567.	1.8	23
62	Association between Genetic Variants in DNA Double-Strand Break Repair Pathways and Risk of Radiation Therapy-Induced Pneumonitis and Esophagitis in Non-Small Cell Lung Cancer. Cancers, 2016, 8, 23.	3.7	13
63	MiRNA-Related Genetic Variations Associated with Radiotherapy-Induced Toxicities in Patients with Locally Advanced Nonâ€"Small Cell Lung Cancer. PLoS ONE, 2016, 11, e0150467.	2.5	7
64	Glycemic Index, Glycemic Load, and Lung Cancer Risk in Non-Hispanic Whites. Cancer Epidemiology Biomarkers and Prevention, 2016, 25, 532-539.	2.5	33
65	Mosaic loss of chromosome Y is associated with common variation near TCL1A. Nature Genetics, 2016, 48, 563-568.	21.4	134
66	Cancer Incidence in First- and Second-Degree Relatives of BRCA1 and BRCA2 Mutation Carriers. Oncologist, 2016, 21, 869-874.	3.7	41
67	Geneâ€environment interaction of genomeâ€wide association studyâ€identified susceptibility loci and meatâ€cooking mutagens in the etiology of renal cell carcinoma. Cancer, 2016, 122, 108-115.	4.1	24
68	Genetic variants of the Wnt signaling pathway as predictors of aggressive disease and reclassification in men with early stage prostate cancer on active surveillance. Carcinogenesis, 2016, 37, 965-971.	2.8	4
69	Meta-analysis of genome-wide association studies discovers multiple loci for chronic lymphocytic leukemia. Nature Communications, 2016, 7, 10933.	12.8	94
70	Personalized Risk Assessment in Never, Light, and Heavy Smokers in a prospective cohort in Taiwan. Scientific Reports, 2016, 6, 36482.	3.3	29
71	Different dietary patterns and reduction of lung cancer risk: A large case-control study in the U.S.. Scientific Reports, 2016, 6, 26760.	3.3	18
72	Genetically predicted longer telomere length is associated with increased risk of B-cell lymphoma subtypes. Human Molecular Genetics, 2016, 25, 1663-1676.	2.9	52

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73	Identification of a novel susceptibility locus at 13q34 and refinement of the 20p12.2 region as a multi-signal locus associated with bladder cancer risk in individuals of European ancestry. Human Molecular Genetics, 2016, 25, 1203-1214.	2.9	38
74	Polymorphisms of the centrosomal gene (FGFR1OP) and lung cancer risk: a meta-analysis of 14 463 cases and 44 188 controls. Carcinogenesis, 2016, 37, 280-289.	2.8	7
75	Pathway analysis of bladder cancer genome-wide association study identifies novel pathways involved in bladder cancer development. Genes and Cancer, 2016, 7, 229-239.	1.9	12
76	Genetic variation in the TNF/TRAF2/ASK1/p38 kinase signaling pathway as markers for postoperative pulmonary complications in lung cancer patients. Scientific Reports, 2015, 5, 12068.	3.3	11
77	Multilevel-analysis identify a cis-expression quantitative trait locus associated with risk of renal cell carcinoma. Oncotarget, 2015, 6, 4097-4109.	1.8	1
78	A genome-wide association study of marginal zone lymphoma shows association to the HLA region. Nature Communications, 2015, 6, 5751.	12.8	58
79	Identification of Serum Markers of Esophageal Adenocarcinoma by Global and Targeted Metabolic Profiling. Clinical Gastroenterology and Hepatology, 2015, 13, 1730-1737.e9.	4.4	29
80	Genetic Variants in the Wnt/Î²-Catenin Signaling Pathway as Indicators of Bladder Cancer Risk. Journal of Urology, 2015, 194, 1771-1776.	0.4	32
81	Depressive Symptoms and Short Telomere Length Are Associated with Increased Mortality in Bladder Cancer Patients. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 336-343.	2.5	33
82	Mitochondrial DNA Content as Risk Factor for Bladder Cancer and Its Association with Mitochondrial DNA Polymorphisms. Cancer Prevention Research, 2015, 8, 607-613.	1.5	18
83	The Ability of Bilirubin in Identifying Smokers with Higher Risk of Lung Cancer: A Large Cohort Study in Conjunction with Global Metabolomic Profiling. Clinical Cancer Research, 2015, 21, 193-200.	7.0	51
84	Genetic Variations in Glutathione Pathway Genes Predict Cancer Recurrence in Patients Treated with Transurethral Resection and Bacillus Calmetteâ€“Guerin Instillation for Non-muscle Invasive Bladder Cancer. Annals of Surgical Oncology, 2015, 22, 4104-4110.	1.5	31
85	CYP2A6 reduced activity gene variants confer reduction in lung cancer risk in African American smokersâ€”findings from two independent populations. Carcinogenesis, 2015, 36, 99-103.	2.8	41
86	Prognostic significance of pretreatment serum levels of albumin, LDH and total bilirubin in patients with non-metastatic breast cancer. Carcinogenesis, 2015, 36, 243-248.	2.8	124
87	Genome-wide association study identifies multiple susceptibility loci for diffuse large B cell lymphoma. Nature Genetics, 2014, 46, 1233-1238.	21.4	147
88	Genome-wide Association Study Identifies Five Susceptibility Loci for Follicular Lymphoma outside the HLA Region. American Journal of Human Genetics, 2014, 95, 462-471.	6.2	96
89	Intratumor heterogeneity in localized lung adenocarcinomas delineated by multiregion sequencing. Science, 2014, 346, 256-259.	12.6	834
90	Rare variants of large effect in BRCA2 and CHEK2 affect risk of lung cancer. Nature Genetics, 2014, 46, 736-741.	21.4	360

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91	Inflammation-Related Genetic Variations and Survival in Patients With Advanced Nonâ€“Small Cell Lung Cancer Receiving First-Line Chemotherapy. Clinical Pharmacology and Therapeutics, 2014, 96, 360-369.	4.7	16
92	Common variation at 2q22.3 (ZEB2) influences the risk of renal cancer. Human Molecular Genetics, 2013, 22, 825-831.	2.9	54
93	Application of Multi-SNP Approaches Bayesian LASSO and AUC-RF to Detect Main Effects of Inflammatory-Gene Variants Associated with Bladder Cancer Risk. PLoS ONE, 2013, 8, e83745.	2.5	21
94	A LIN28B polymorphism predicts for colon cancer survival. Cancer Biology and Therapy, 2012, 13, 1390-1395.	3.4	12
95	A genome-wide association study identifies a novel susceptibility locus for renal cell carcinoma on 12p11.23. Human Molecular Genetics, 2012, 21, 456-462.	2.9	81
96	Genome-wide association study of renal cell carcinoma identifies two susceptibility loci on 2p21 and 11q13.3. Nature Genetics, 2011, 43, 60-65.	21.4	220
97	Genetic variants in cell cycle control pathway confer susceptibility to bladder cancer. Cancer, 2008, 112, 2467-2474.	4.1	52
98	Genetic variations in cellâ€cycle pathway and the risk of oral premalignant lesions. Cancer, 2008, 113, 2488-2495.	4.1	21
99	Joint Modeling of Time Series Measures and Recurrent Events and Analysis of the Effects of Air Quality on Respiratory Symptoms. Journal of the American Statistical Association, 2008, 103, 48-60.	3.1	9
100	Genetic Variations in MicroRNA-Related Genes Are Novel Susceptibility Loci for Esophageal Cancer Risk. Cancer Prevention Research, 2008, 1, 460-469.	1.5	206