

# Yuanqing Ye

## List of Publications by Year in descending order

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Version: 2024-02-01

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	Intratumor heterogeneity in localized lung adenocarcinomas delineated by multiregion sequencing. <i>Science</i> , 2014, 346, 256-259.	12.6	834
2	Large-scale association analysis identifies new lung cancer susceptibility loci and heterogeneity in genetic susceptibility across histological subtypes. <i>Nature Genetics</i> , 2017, 49, 1126-1132.	21.4	472
3	Rare variants of large effect in BRCA2 and CHEK2 affect risk of lung cancer. <i>Nature Genetics</i> , 2014, 46, 736-741.	21.4	360
4	Genome-wide association study of renal cell carcinoma identifies two susceptibility loci on 2p21 and 11q13.3. <i>Nature Genetics</i> , 2011, 43, 60-65.	21.4	220
5	Genetic Variations in MicroRNA-Related Genes Are Novel Susceptibility Loci for Esophageal Cancer Risk. <i>Cancer Prevention Research</i> , 2008, 1, 460-469.	1.5	206
6	Genome-wide association study identifies multiple susceptibility loci for diffuse large B cell lymphoma. <i>Nature Genetics</i> , 2014, 46, 1233-1238.	21.4	147
7	Comprehensive T cell repertoire characterization of non-small cell lung cancer. <i>Nature Communications</i> , 2020, 11, 603.	12.8	140
8	Mosaic loss of chromosome Y is associated with common variation near TCL1A. <i>Nature Genetics</i> , 2016, 48, 563-568.	21.4	134
9	Prognostic significance of pretreatment serum levels of albumin, LDH and total bilirubin in patients with non-metastatic breast cancer. <i>Carcinogenesis</i> , 2015, 36, 243-248.	2.8	124
10	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
11	Genome-wide association study identifies multiple risk loci for renal cell carcinoma. <i>Nature Communications</i> , 2017, 8, 15724.	12.8	106
12	Cancer risk associated with chronic diseases and disease markers: prospective cohort study. <i>BMJ: British Medical Journal</i> , 2018, 360, k134.	2.3	97
13	Genome-wide Association Study Identifies Five Susceptibility Loci for Follicular Lymphoma outside the HLA Region. <i>American Journal of Human Genetics</i> , 2014, 95, 462-471.	6.2	96
14	Meta-analysis of genome-wide association studies discovers multiple loci for chronic lymphocytic leukemia. <i>Nature Communications</i> , 2016, 7, 10933.	12.8	94
15	A genome-wide association study identifies a novel susceptibility locus for renal cell carcinoma on 12p11.23. <i>Human Molecular Genetics</i> , 2012, 21, 456-462.	2.9	81
16	Identification of susceptibility pathways for the role of chromosome 15q25.1 in modifying lung cancer risk. <i>Nature Communications</i> , 2018, 9, 3221.	12.8	60
17	The influence of obesity-related factors in the etiology of renal cell carcinoma—A mendelian randomization study. <i>PLoS Medicine</i> , 2019, 16, e1002724.	8.4	59
18	A genome-wide association study of marginal zone lymphoma shows association to the HLA region. <i>Nature Communications</i> , 2015, 6, 5751.	12.8	58

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19	Angiogenin/Ribonuclease 5 Is an EGFR Ligand and a Serum Biomarker for Erlotinib Sensitivity in Pancreatic Cancer. <i>Cancer Cell</i> , 2018, 33, 752-769.e8.	16.8	58
20	Common variation at 2q22.3 (ZEB2) influences the risk of renal cancer. <i>Human Molecular Genetics</i> , 2013, 22, 825-831.	2.9	54
21	Genetic variants in cell cycle control pathway confer susceptibility to bladder cancer. <i>Cancer</i> , 2008, 112, 2467-2474.	4.1	52
22	Genetically predicted longer telomere length is associated with increased risk of B-cell lymphoma subtypes. <i>Human Molecular Genetics</i> , 2016, 25, 1663-1676.	2.9	52
23	The somatic mutation landscape of premalignant colorectal adenoma. <i>Gut</i> , 2018, 67, 1299-1305.	12.1	52
24	A 5-microRNA signature identified from serum microRNA profiling predicts survival in patients with advanced stage non-small cell lung cancer. <i>Carcinogenesis</i> , 2019, 40, 643-650.	2.8	52
25	The Ability of Bilirubin in Identifying Smokers with Higher Risk of Lung Cancer: A Large Cohort Study in Conjunction with Global Metabolomic Profiling. <i>Clinical Cancer Research</i> , 2015, 21, 193-200.	7.0	51
26	Global and targeted serum metabolic profiling of colorectal cancer progression. <i>Cancer</i> , 2017, 123, 4066-4074.	4.1	51
27	Personalized Prognostic Prediction Models for Breast Cancer Recurrence and Survival Incorporating Multidimensional Data. <i>Journal of the National Cancer Institute</i> , 2017, 109, .	6.3	42
28	CYP2A6 reduced activity gene variants confer reduction in lung cancer risk in African American smokersâ€™ findings from two independent populations. <i>Carcinogenesis</i> , 2015, 36, 99-103.	2.8	41
29	Cancer Incidence in First- and Second-Degree Relatives of <i>BRCA1</i> and <i>BRCA2</i> Mutation Carriers. <i>Oncologist</i> , 2016, 21, 869-874.	3.7	41
30	Genetic Variants Related to Longer Telomere Length are Associated with Increased Risk of Renal Cell Carcinoma. <i>European Urology</i> , 2017, 72, 747-754.	1.9	39
31	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. <i>Human Molecular Genetics</i> , 2016, 25, 1203-1214.	2.9	38
32	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. <i>American Journal of Pathology</i> , 2018, 188, 2487-2496.	3.8	34
33	Depressive Symptoms and Short Telomere Length Are Associated with Increased Mortality in Bladder Cancer Patients. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 336-343.	2.5	33
34	Glycemic Index, Glycemic Load, and Lung Cancer Risk in Non-Hispanic Whites. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 532-539.	2.5	33
35	Genetic Variants in the Wnt/ $\beta$ -Catenin Signaling Pathway as Indicators of Bladder Cancer Risk. <i>Journal of Urology</i> , 2015, 194, 1771-1776.	0.4	32
36	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. <i>Annals of Surgical Oncology</i> , 2015, 22, 4104-4110.	1.5	31

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37	Serum MicroRNA-150 Predicts Prognosis for Early-Stage Non-Small Cell Lung Cancer and Promotes Tumor Cell Proliferation by Targeting Tumor Suppressor Gene <i>SRGIN1</i> . <i>Clinical Pharmacology and Therapeutics</i> , 2018, 103, 1061-1073.	4.7	31
38	Identification of Serum Markers of Esophageal Adenocarcinoma by Global and Targeted Metabolic Profiling. <i>Clinical Gastroenterology and Hepatology</i> , 2015, 13, 1730-1737.e9.	4.4	29
39	Personalized Risk Assessment in Never, Light, and Heavy Smokers in a prospective cohort in Taiwan. <i>Scientific Reports</i> , 2016, 6, 36482.	3.3	29
40	Cohort Profile: The Mexican American Mano a Mano Cohort. <i>International Journal of Epidemiology</i> , 2017, 46, e3-e3.	1.9	28
41	Sex specific associations in genome wide association analysis of renal cell carcinoma. <i>European Journal of Human Genetics</i> , 2019, 27, 1589-1598.	2.8	27
42	Serum miR-331-3p predicts tumor recurrence in esophageal adenocarcinoma. <i>Scientific Reports</i> , 2018, 8, 14006.	3.3	26
43	Gene-environment interaction of genome-wide association study-identified susceptibility loci and meat-cooking mutagens in the etiology of renal cell carcinoma. <i>Cancer</i> , 2016, 122, 108-115.	4.1	24
44	Hypoxia-targeted gold nanorods for cancer photothermal therapy. <i>Oncotarget</i> , 2018, 9, 26556-26571.	1.8	24
45	Dietary patterns and risk of recurrence and progression in non-muscle-invasive bladder cancer. <i>International Journal of Cancer</i> , 2018, 142, 1797-1804.	5.1	23
46	Social-demographics, health behaviors, and telomere length in the Mexican American Mano a Mano Cohort. <i>Oncotarget</i> , 2017, 8, 96553-96567.	1.8	23
47	Genetic variations in cell-cycle pathway and the risk of oral premalignant lesions. <i>Cancer</i> , 2008, 113, 2488-2495.	4.1	21
48	Application of Multi-SNP Approaches Bayesian LASSO and AUC-RF to Detect Main Effects of Inflammatory-Gene Variants Associated with Bladder Cancer Risk. <i>PLoS ONE</i> , 2013, 8, e83745.	2.5	21
49	Predictors of health-related quality of life and association with survival may identify colorectal cancer patients at high risk of poor prognosis. <i>Quality of Life Research</i> , 2017, 26, 319-330.	3.1	21
50	Determinants and prognostic value of quality of life in patients with pancreatic ductal adenocarcinoma. <i>European Journal of Cancer</i> , 2018, 92, 20-32.	2.8	21
51	Elevated Platelet Count Appears to Be Causally Associated with Increased Risk of Lung Cancer: A Mendelian Randomization Analysis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2019, 28, 935-942.	2.5	21
52	Allostatic score and its associations with demographics, healthy behaviors, tumor characteristics, and mitochondrial DNA among breast cancer patients. <i>Breast Cancer Research and Treatment</i> , 2021, 187, 587-596.	2.5	21
53	D-mannose: a Novel Prognostic Biomarker for Patients with Esophageal Adenocarcinoma. <i>Carcinogenesis</i> , 2017, 38, bgw207.	2.8	19
54	Is folic acid safe for non-muscle-invasive bladder cancer patients? An evidence-based cohort study. <i>American Journal of Clinical Nutrition</i> , 2018, 107, 208-216.	4.7	19

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55	Prevalence of Aflatoxin-Associated <i>TP53R249S</i> Mutation in Hepatocellular Carcinoma in Hispanics in South Texas. <i>Cancer Prevention Research</i> , 2018, 11, 103-112.	1.5	19
56	Mitochondrial DNA Content as Risk Factor for Bladder Cancer and Its Association with Mitochondrial DNA Polymorphisms. <i>Cancer Prevention Research</i> , 2015, 8, 607-613.	1.5	18
57	Different dietary patterns and reduction of lung cancer risk: A large case-control study in the U.S.. <i>Scientific Reports</i> , 2016, 6, 26760.	3.3	18
58	High baseline levels of interleukin-8 in leukocytes and urine predict tumor recurrence in non-muscle invasive bladder cancer patients receiving bacillus Calmette-Guérin therapy: A long-term survival analysis. <i>Oncotarget</i> , 2017, 6, e1265719.	4.6	18
59	Polymorphisms in genes related to epithelial-mesenchymal transition and risk of non-small cell lung cancer. <i>Carcinogenesis</i> , 2017, 38, 1029-1035.	2.8	18
60	Germline genetic variants in somatically significantly mutated genes in tumors are associated with renal cell carcinoma risk and outcome. <i>Carcinogenesis</i> , 2018, 39, 752-757.	2.8	18
61	Unruptured intracranial aneurysm growth trajectory: occurrence and rate of enlargement in 520 longitudinally followed cases. <i>Journal of Neurosurgery</i> , 2020, 132, 1077-1087.	1.6	17
62	Inflammation-Related Genetic Variations and Survival in Patients With Advanced Non-Small Cell Lung Cancer Receiving First-Line Chemotherapy. <i>Clinical Pharmacology and Therapeutics</i> , 2014, 96, 360-369.	4.7	16
63	Functional variants in DCAF4 associated with lung cancer risk in European populations. <i>Carcinogenesis</i> , 2017, 38, 541-551.	2.8	16
64	Socio-demographic, Clinical, and Genetic Determinants of Quality of Life in Lung Cancer Patients. <i>Scientific Reports</i> , 2018, 8, 10640.	3.3	16
65	Circulating metabolite profiles to predict overall survival in advanced non-small cell lung cancer patients receiving first-line chemotherapy. <i>Lung Cancer</i> , 2017, 114, 70-78.	2.0	15
66	Common, germline genetic variations in the novel tumor suppressor <i>BAP1</i> and risk of developing different types of cancer. <i>Oncotarget</i> , 2017, 8, 74936-74946.	1.8	15
67	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. <i>Cancers</i> , 2016, 8, 23.	3.7	13
68	Effect of physiological factors on the biochemical properties of colon tissue – an <i>in vivo</i> Raman spectroscopy study. <i>Journal of Raman Spectroscopy</i> , 2017, 48, 902-909.	2.5	13
69	Breast cancer risk in relation to plasma metabolites among Hispanic and African American women. <i>Breast Cancer Research and Treatment</i> , 2019, 176, 687-696.	2.5	13
70	A <i>LIN28B</i> polymorphism predicts for colon cancer survival. <i>Cancer Biology and Therapy</i> , 2012, 13, 1390-1395.	3.4	12
71	Pathway analysis of bladder cancer genome-wide association study identifies novel pathways involved in bladder cancer development. <i>Genes and Cancer</i> , 2016, 7, 229-239.	1.9	12
72	Genetic variation in the TNF/TRAF2/ASK1/p38 kinase signaling pathway as markers for postoperative pulmonary complications in lung cancer patients. <i>Scientific Reports</i> , 2015, 5, 12068.	3.3	11

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73	HIF3A DNA methylation, obesity and weight gain, and breast cancer risk among Mexican American women. <i>Obesity Research and Clinical Practice</i> , 2020, 14, 548-553.	1.8	11
74	Human ribonuclease 1 serves as a secretory ligand of ephrin A4 receptor and induces breast tumor initiation. <i>Nature Communications</i> , 2021, 12, 2788.	12.8	11
75	Serum microRNAs as predictors of risk for non-muscle invasive bladder cancer. <i>Oncotarget</i> , 2018, 9, 14895-14908.	1.8	11
76	Genetic Variants in Epigenetic Pathways and Risks of Multiple Cancers in the GAME-ON Consortium. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 816-825.	2.5	10
77	Genetic variants of PTPN2 are associated with lung cancer risk: a re-analysis of eight GWASs in the TRICL-ILCCO consortium. <i>Scientific Reports</i> , 2017, 7, 825.	3.3	10
78	Associations between genetic variants in mRNA splicing-related genes and risk of lung cancer: a pathway-based analysis from published GWASs. <i>Scientific Reports</i> , 2017, 7, 44634.	3.3	10
79	Susceptibility loci of <i>CNOT6</i> in the general mRNA degradation pathway and lung cancer risk—A re-analysis of eight GWASs. <i>Molecular Carcinogenesis</i> , 2017, 56, 1227-1238.	2.7	10
80	Metabolic hormones and breast cancer risk among Mexican American Women in the Mano a Mano Cohort Study. <i>Scientific Reports</i> , 2019, 9, 9989.	3.3	10
81	Patterns of racial/ethnic disparities in baseline health-related quality of life and relationship with overall survival in patients with colorectal cancer. <i>Quality of Life Research</i> , 2020, 29, 2977-2986.	3.1	10
82	Joint Modeling of Time Series Measures and Recurrent Events and Analysis of the Effects of Air Quality on Respiratory Symptoms. <i>Journal of the American Statistical Association</i> , 2008, 103, 48-60.	3.1	9
83	Novel genetic variants in the P38MAPK pathway gene <i>ZAK</i> and susceptibility to lung cancer. <i>Molecular Carcinogenesis</i> , 2018, 57, 216-224.	2.7	9
84	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
85	MiRNA-Related Genetic Variations Associated with Radiotherapy-Induced Toxicities in Patients with Locally Advanced Non-Small Cell Lung Cancer. <i>PLoS ONE</i> , 2016, 11, e0150467.	2.5	7
86	Polymorphisms of the centrosomal gene ( <i>FGFR1OP</i> ) and lung cancer risk: a meta-analysis of 14 463 cases and 44 188 controls. <i>Carcinogenesis</i> , 2016, 37, 280-289.	2.8	7
87	Phase I study of nab-paclitaxel, gemcitabine, and bevacizumab in patients with advanced cancers. <i>British Journal of Cancer</i> , 2018, 118, 1419-1424.	6.4	7
88	Genetic associations of T cell cancer immune response with tumor aggressiveness in localized prostate cancer patients and disease reclassification in an active surveillance cohort. <i>Oncology</i> , 2019, 8, e1483303.	4.6	7
89	Reply to “Mosaic loss of chromosome Y in leukocytes matters”. <i>Nature Genetics</i> , 2019, 51, 7-9.	21.4	7
90	Associations of blood mitochondrial DNA copy number with social-demographics and cancer risk: results from the Mano-A-Mano Mexican American Cohort. <i>Oncotarget</i> , 2018, 9, 25491-25502.	1.8	6

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91	Genetic variants in cytokine signaling pathways and clinical outcomes in early-stage lung cancer patients. <i>Journal of Thoracic and Cardiovascular Surgery</i> , 2018, 155, 2635-2645.e15.	0.8	5
92	Nanotrap-enabled quantification of KRAS-induced peptide hydroxylation in blood for cancer early detection. <i>Nano Research</i> , 2019, 12, 1445-1452.	10.4	5
93	Genetic variants of the Wnt signaling pathway as predictors of aggressive disease and reclassification in men with early stage prostate cancer on active surveillance. <i>Carcinogenesis</i> , 2016, 37, 965-971.	2.8	4
94	Potential Susceptibility Loci Identified for Renal Cell Carcinoma by Targeting Obesity-Related Genes. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 1436-1442.	2.5	2
95	Multilevel-analysis identify a cis-expression quantitative trait locus associated with risk of renal cell carcinoma. <i>Oncotarget</i> , 2015, 6, 4097-4109.	1.8	1
96	Validation of plasma metabolites associated with breast cancer risk among Mexican Americans. <i>Cancer Epidemiology</i> , 2020, 69, 101826.	1.9	1
97	Land use mix and leukocyte telomere length in Mexican Americans. <i>Scientific Reports</i> , 2021, 11, 19742.	3.3	1
98	Response. <i>Journal of the National Cancer Institute</i> , 2017, 109, .	6.3	0
99	Leukocyte telomere length associated with glioma risk and survival. <i>Ageing and Cancer</i> , 2020, 1, 71-78.	1.6	0
100	Genetic variants in epithelial-mesenchymal transition genes as predictors of clinical outcomes in localized prostate cancer. <i>Carcinogenesis</i> , 2020, 41, 1057-1064.	2.8	0