

# Xifeng Wu

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/83402/publications.pdf>

Version: 2024-02-01

400  
papers

25,774  
citations

6613

79  
h-index

10445

139  
g-index

410  
all docs

410  
docs citations

410  
times ranked

33166  
citing authors

#	ARTICLE	IF	CITATIONS
1	Minimum amount of physical activity for reduced mortality and extended life expectancy: a prospective cohort study. <i>Lancet, The</i> , 2011, 378, 1244-1253.	13.7	1,495
2	Genome-wide association scan of tag SNPs identifies a susceptibility locus for lung cancer at 15q25.1. <i>Nature Genetics</i> , 2008, 40, 616-622.	21.4	1,189
3	Intratumor heterogeneity in localized lung adenocarcinomas delineated by multiregion sequencing. <i>Science</i> , 2014, 346, 256-259.	12.6	834
4	Detectable clonal mosaicism and its relationship to aging and cancer. <i>Nature Genetics</i> , 2012, 44, 651-658.	21.4	519
5	Common 5p15.33 and 6p21.33 variants influence lung cancer risk. <i>Nature Genetics</i> , 2008, 40, 1407-1409.	21.4	510
6	A multi-stage genome-wide association study of bladder cancer identifies multiple susceptibility loci. <i>Nature Genetics</i> , 2010, 42, 978-984.	21.4	493
7	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
8	Perinatal depressive and anxiety symptoms of pregnant women during the coronavirus disease 2019 outbreak in China. <i>American Journal of Obstetrics and Gynecology</i> , 2020, 223, 240.e1-240.e9.	1.3	438
9	Telomere Dysfunction: A Potential Cancer Predisposition Factor. <i>Journal of the National Cancer Institute</i> , 2003, 95, 1211-1218.	6.3	436
10	A Risk Model for Prediction of Lung Cancer. <i>Journal of the National Cancer Institute</i> , 2007, 99, 715-726.	6.3	362
11	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
12	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. <i>Nature Genetics</i> , 2017, 49, 680-691.	21.4	356
13	Genetic variation in the prostate stem cell antigen gene PSCA confers susceptibility to urinary bladder cancer. <i>Nature Genetics</i> , 2009, 41, 991-995.	21.4	321
14	Preleukaemic clonal haemopoiesis and risk of therapy-related myeloid neoplasms: a case-control study. <i>Lancet Oncology, The</i> , 2017, 18, 100-111.	10.7	296
15	Focus on bladder cancer. <i>Cancer Cell</i> , 2004, 6, 111-116.	16.8	252
16	Bladder Cancer Predisposition: A Multigenic Approach to DNA-Repair and Cell-Cycle Control Genes. <i>American Journal of Human Genetics</i> , 2006, 78, 464-479.	6.2	249
17	Evaluation of Genetic Variants in MicroRNA-Related Genes and Risk of Bladder Cancer. <i>Cancer Research</i> , 2008, 68, 2530-2537.	0.9	245
18	Mitochondrial DNA Content: Its Genetic Heritability and Association With Renal Cell Carcinoma. <i>Journal of the National Cancer Institute</i> , 2008, 100, 1104-1112.	6.3	237

#	ARTICLE	IF	CITATIONS
19	Identification of six new susceptibility loci for invasive epithelial ovarian cancer. <i>Nature Genetics</i> , 2015, 47, 164-171.	21.4	221
20	Case-Control Study of the D2 Dopamine Receptor Gene and Smoking Status in Lung Cancer Patients. <i>Journal of the National Cancer Institute</i> , 1998, 90, 358-363.	6.3	220
21	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
22	Genetic variants and risk of lung cancer in never smokers: a genome-wide association study. <i>Lancet Oncology</i> , The, 2010, 11, 321-330.	10.7	218
23	p53 Genotypes and Haplotypes Associated With Lung Cancer Susceptibility and Ethnicity. <i>Journal of the National Cancer Institute</i> , 2002, 94, 681-690.	6.3	214
24	The CHRNA5-A3 Region on Chromosome 15q24-25.1 Is a Risk Factor Both for Nicotine Dependence and for Lung Cancer. <i>Journal of the National Cancer Institute</i> , 2008, 100, 1552-1556.	6.3	206
25	Influence of common genetic variation on lung cancer risk: meta-analysis of 14 900 cases and 29 485 controls. <i>Human Molecular Genetics</i> , 2012, 21, 4980-4995.	2.9	196
26	From genotype to phenotype: correlating XRCC1 polymorphisms with mutagen sensitivity. <i>DNA Repair</i> , 2003, 2, 901-908.	2.8	184
27	Replication of Lung Cancer Susceptibility Loci at Chromosomes 15q25, 5p15, and 6p21: A Pooled Analysis From the International Lung Cancer Consortium. <i>Journal of the National Cancer Institute</i> , 2010, 102, 959-971.	6.3	174
28	<i>PALB2</i> , <i>CHEK2</i> and <i>ATM</i> rare variants and cancer risk: data from COGS. <i>Journal of Medical Genetics</i> , 2016, 53, 800-811.	3.2	174
29	Genetic Variations in Radiation and Chemotherapy Drug Action Pathways Predict Clinical Outcomes in Esophageal Cancer. <i>Journal of Clinical Oncology</i> , 2006, 24, 3789-3798.	1.6	165
30	Energy stress-induced lncRNA FILNC1 represses c-Myc-mediated energy metabolism and inhibits renal tumor development. <i>Nature Communications</i> , 2017, 8, 783.	12.8	157
31	Genetic Variations in the PI3K/PTEN/AKT/mTOR Pathway Are Associated With Clinical Outcomes in Esophageal Cancer Patients Treated With Chemoradiotherapy. <i>Journal of Clinical Oncology</i> , 2009, 27, 857-871.	1.6	154
32	Meta-analysis identifies four new loci associated with testicular germ cell tumor. <i>Nature Genetics</i> , 2013, 45, 680-685.	21.4	154
33	XPA polymorphism associated with reduced lung cancer risk and a modulating effect on nucleotide excision repair capacity. <i>Carcinogenesis</i> , 2003, 24, 505-509.	2.8	149
34	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
35	Genetic susceptibility to tobacco-related cancer. <i>Oncogene</i> , 2004, 23, 6500-6523.	5.9	146
36	An Expanded Risk Prediction Model for Lung Cancer. <i>Cancer Prevention Research</i> , 2008, 1, 250-254.	1.5	143

#	ARTICLE	IF	CITATIONS
37	Familial Aggregation of Common Sequence Variants on 15q24-25.1 in Lung Cancer. <i>Journal of the National Cancer Institute</i> , 2008, 100, 1326-1330.	6.3	141
38	Comprehensive T cell repertoire characterization of non-small cell lung cancer. <i>Nature Communications</i> , 2020, 11, 603.	12.8	140
39	Polymorphisms in Inflammation Genes and Bladder Cancer: From Initiation to Recurrence, Progression, and Survival. <i>Journal of Clinical Oncology</i> , 2005, 23, 5746-5756.	1.6	138
40	Genome-wide association study identifies multiple loci associated with bladder cancer risk. <i>Human Molecular Genetics</i> , 2014, 23, 1387-1398.	2.9	137
41	Higher lung cancer risk for younger African-Americans with the Pro/Pro p53 genotype. <i>Carcinogenesis</i> , 1995, 16, 2205-2208.	2.8	134
42	Mosaic loss of chromosome Y is associated with common variation near TCL1A. <i>Nature Genetics</i> , 2016, 48, 563-568.	21.4	134
43	Genetic Variants on 15q25.1, Smoking, and Lung Cancer: An Assessment of Mediation and Interaction. <i>American Journal of Epidemiology</i> , 2012, 175, 1013-1020.	3.4	128
44	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
45	Genetic susceptibility to lung cancer: the role of DNA damage and repair. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2003, 12, 689-98.	2.5	123
46	MicroRNA Expression Signatures in Barrett's Esophagus and Esophageal Adenocarcinoma. <i>Clinical Cancer Research</i> , 2009, 15, 5744-5752.	7.0	120
47	Genetic variations in microRNA-related genes are associated with survival and recurrence in patients with renal cell carcinoma. <i>Carcinogenesis</i> , 2010, 31, 1805-1812.	2.8	120
48	Genetic Variants in MicroRNA Biosynthesis Pathways and Binding Sites Modify Ovarian Cancer Risk, Survival, and Treatment Response. <i>Cancer Research</i> , 2010, 70, 9765-9776.	0.9	118
49	Household transmission of COVID-19-a systematic review and meta-analysis. <i>Journal of Infection</i> , 2020, 81, 979-997.	3.3	117
50	Association of vitamin D levels and risk of ovarian cancer: a Mendelian randomization study. <i>International Journal of Epidemiology</i> , 2016, 45, 1619-1630.	1.9	111
51	Diabetes with early kidney involvement may shorten life expectancy by 16 years. <i>Kidney International</i> , 2017, 92, 388-396.	5.2	109
52	Polymorphisms in DNA Repair Genes, Smoking, and Bladder Cancer Risk: Findings from the International Consortium of Bladder Cancer. <i>Cancer Research</i> , 2009, 69, 6857-6864.	0.9	107
53	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
54	Genome-wide association study identifies multiple risk loci for renal cell carcinoma. <i>Nature Communications</i> , 2017, 8, 15724.	12.8	106

#	ARTICLE	IF	CITATIONS
55	Analysis of Genomes and Transcriptomes of Hepatocellular Carcinomas Identifies Mutations and Gene Expression Changes in the Transforming Growth Factor- $\beta$ Pathway. <i>Gastroenterology</i> , 2018, 154, 195-210.	1.3	105
56	Characterization of Large Structural Genetic Mosaicism in Human Autosomes. <i>American Journal of Human Genetics</i> , 2015, 96, 487-497.	6.2	101
57	An Evolutionary Perspective on Single-Nucleotide Polymorphism Screening in Molecular Cancer Epidemiology. <i>Cancer Research</i> , 2004, 64, 2251-2257.	0.9	100
58	A genome-wide association study of bladder cancer identifies a new susceptibility locus within SLC14A1, a urea transporter gene on chromosome 18q12.3. <i>Human Molecular Genetics</i> , 2011, 20, 4282-4289.	2.9	100
59	Cancer risk associated with chronic diseases and disease markers: prospective cohort study. <i>BMJ: British Medical Journal</i> , 2018, 360, k134.	2.3	97
60	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
61	Hepatocellular Carcinoma Risk Prediction Model for the General Population: The Predictive Power of Transaminases. <i>Journal of the National Cancer Institute</i> , 2012, 104, 1599-1611.	6.3	95
62	Sulfotransferase (SULT) 1A1 polymorphism as a predisposition factor for lung cancer: a case-control analysis. <i>Lung Cancer</i> , 2002, 35, 137-142.	2.0	94
63	Meta-analysis of genome-wide association studies discovers multiple loci for chronic lymphocytic leukemia. <i>Nature Communications</i> , 2016, 7, 10933.	12.8	94
64	Effects of N-acetyl transferase 1 and 2 polymorphisms on bladder cancer risk in Caucasians. <i>Mutation Research - Genetic Toxicology and Environmental Mutagenesis</i> , 2005, 581, 97-104.	1.7	92
65	Molecular epidemiology, cancer-related symptoms, and cytokines pathway. <i>Lancet Oncology</i> , The, 2008, 9, 777-785.	10.7	92
66	Family History and Risk of Renal Cell Carcinoma: Results from a Case-Control Study and Systematic Meta-Analysis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 801-807.	2.5	91
67	Genetic Variants in Inflammation-Related Genes Are Associated with Radiation-Induced Toxicity Following Treatment for Non-Small Cell Lung Cancer. <i>PLoS ONE</i> , 2010, 5, e12402.	2.5	91
68	MicroRNA Expression Signatures during Malignant Progression from Barrett's Esophagus to Esophageal Adenocarcinoma. <i>Cancer Prevention Research</i> , 2013, 6, 196-205.	1.5	91
69	Genetic variation in MicroRNA genes and risk of oral premalignant lesions. <i>Molecular Carcinogenesis</i> , 2010, 49, 183-189.	2.7	90
70	Imputation and subset-based association analysis across different cancer types identifies multiple independent risk loci in the TERT-CLPTM1L region on chromosome 5p15.33. <i>Human Molecular Genetics</i> , 2014, 23, 6616-6633.	2.9	90
71	Genetic Instability in Bladder Cancer Assessed by the Comet Assay. <i>Journal of the National Cancer Institute</i> , 2003, 95, 540-547.	6.3	89
72	MicroRNA in the Pathogenesis and Prognosis of Esophageal Cancer. <i>Current Pharmaceutical Design</i> , 2012, 19, 1292-1300.	1.9	89

#	ARTICLE	IF	CITATIONS
73	Nucleotide Excision Repair Gene Polymorphisms and Recurrence after Treatment for Superficial Bladder Cancer. <i>Clinical Cancer Research</i> , 2005, 11, 1408-1415.	7.0	88
74	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019, 10, 431.	12.8	88
75	High-Order Interactions among Genetic Variants in DNA Base Excision Repair Pathway Genes and Smoking in Bladder Cancer Susceptibility. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007, 16, 84-91.	2.5	87
76	Female chromosome X mosaicism is age-related and preferentially affects the inactivated X chromosome. <i>Nature Communications</i> , 2016, 7, 11843.	12.8	86
77	Genetic variants of myeloperoxidase and lung cancer risk. <i>Carcinogenesis</i> , 2000, 21, 1163-1166.	2.8	85
78	Genetic variations in PI3K-AKT-mTOR pathway and bladder cancer risk. <i>Carcinogenesis</i> , 2009, 30, 2047-2052.	2.8	85
79	A myeloperoxidase polymorphism associated with reduced risk of lung cancer. <i>Lung Cancer</i> , 2002, 37, 35-40.	2.0	84
80	Dietary isothiocyanates, GSTM1, GSTT1, NAT2 polymorphisms and bladder cancer risk. <i>International Journal of Cancer</i> , 2007, 120, 2208-2213.	5.1	82
81	Genome-Wide Association Study of Survival in Non-Small Cell Lung Cancer Patients Receiving Platinum-Based Chemotherapy. <i>Journal of the National Cancer Institute</i> , 2011, 103, 817-825.	6.3	81
82	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
83	A Serological Biopsy Using Five Stomach-Specific Circulating Biomarkers for Gastric Cancer Risk Assessment: A Multi-Phase Study. <i>American Journal of Gastroenterology</i> , 2017, 112, 704-715.	0.4	81
84	Genome-wide profiling of chromosomal alterations in renal cell carcinoma using high-density single nucleotide polymorphism arrays. <i>International Journal of Cancer</i> , 2009, 125, 2342-2348.	5.1	80
85	Pathway-Based Serum microRNA Profiling and Survival in Patients with Advanced Stage Non-Small Cell Lung Cancer. <i>Cancer Research</i> , 2013, 73, 4801-4809.	0.9	80
86	Telomere Length in Peripheral Blood Leukocytes and Lung Cancer Risk: A Large Case-Control Study in Caucasians. <i>Cancer Research</i> , 2014, 74, 2476-2486.	0.9	80
87	Obesity, metabolic factors and risk of different histological types of lung cancer: A Mendelian randomization study. <i>PLoS ONE</i> , 2017, 12, e0177875.	2.5	79
88	An association between NQO1 genetic polymorphism and risk of bladder cancer. <i>Mutation Research - Genetic Toxicology and Environmental Mutagenesis</i> , 2003, 536, 131-137.	1.7	78
89	Mutagen Sensitivity Has High Heritability: Evidence from a Twin Study. <i>Cancer Research</i> , 2006, 66, 5993-5996.	0.9	78
90	High Serum Iron Is Associated with Increased Cancer Risk. <i>Cancer Research</i> , 2014, 74, 6589-6597.	0.9	77

#	ARTICLE	IF	CITATIONS
91	Modulation of DNA damage/DNA repair capacity by XPC polymorphisms. <i>DNA Repair</i> , 2008, 7, 141-148.	2.8	76
92	Mutagen Sensitivity: A Genetic Predisposition Factor for Cancer: Table 1.. <i>Cancer Research</i> , 2007, 67, 3493-3495.	0.9	75
93	Genetic susceptibility to esophageal cancer: the role of the nucleotide excision repair pathway. <i>Carcinogenesis</i> , 2009, 30, 785-792.	2.8	75
94	Telomere Dysfunction in Peripheral Lymphocytes as a Potential Predisposition Factor for Renal Cancer. <i>Journal of Urology</i> , 2007, 178, 1492-1496.	0.4	74
95	Opposing Effects of Emphysema, Hay Fever, and Select Genetic Variants on Lung Cancer Risk. <i>American Journal of Epidemiology</i> , 2005, 161, 412-422.	3.4	73
96	Genome-Wide Catalogue of Chromosomal Aberrations in Barrett's Esophagus and Esophageal Adenocarcinoma: A High-Density Single Nucleotide Polymorphism Array Analysis. <i>Cancer Prevention Research</i> , 2010, 3, 1176-1186.	1.5	73
97	A Genome-Wide Association Study Identifies a Locus on Chromosome 14q21 as a Predictor of Leukocyte Telomere Length and as a Marker of Susceptibility for Bladder Cancer. <i>Cancer Prevention Research</i> , 2011, 4, 514-521.	1.5	73
98	Serum Levels of Insulin Growth Factor (IGF-I) and IGF-Binding Protein Predict Risk of Second Primary Tumors in Patients with Head and Neck Cancer. <i>Clinical Cancer Research</i> , 2004, 10, 3988-3995.	7.0	72
99	Matrix Metalloproteinase Polymorphisms and Bladder Cancer Risk. <i>Cancer Research</i> , 2006, 66, 11644-11648.	0.9	71
100	Adult body mass index and risk of ovarian cancer by subtype: a Mendelian randomization study. <i>International Journal of Epidemiology</i> , 2016, 45, 884-895.	1.9	71
101	Cohort Profile: The Taiwan MJ Cohort: half a million Chinese with repeated health surveillance data. <i>International Journal of Epidemiology</i> , 2017, 46, 1744-1744g.	1.9	70
102	Constitutive Short Telomere Length of Chromosome 17p and 12q but not 11q and 2p Is Associated with an Increased Risk for Esophageal Cancer. <i>Cancer Prevention Research</i> , 2009, 2, 459-465.	1.5	69
103	A parallel study of in vitro sensitivity to benzo[a]pyrene diol epoxide and bleomycin in lung carcinoma cases and controls. <i>Cancer</i> , 1998, 83, 1118-1127.	4.1	67
104	Projecting Individualized Probabilities of Developing Bladder Cancer in White Individuals. <i>Journal of Clinical Oncology</i> , 2007, 25, 4974-4981.	1.6	67
105	Analysis of aromatic DNA adducts and 7,8-dihydro-8-oxo-2'-deoxyguanosine in lymphocyte DNA from a case-control study of lung cancer involving minority populations. , 2000, 27, 34-46.		65
106	High-order interactions among genetic polymorphisms in nucleotide excision repair pathway genes and smoking in modulating bladder cancer risk. <i>Carcinogenesis</i> , 2007, 28, 2160-2165.	2.8	64
107	Pharmacogenomics of platinum-based chemotherapy in NSCLC. <i>Expert Opinion on Drug Metabolism and Toxicology</i> , 2009, 5, 745-755.	3.3	63
108	Cis-eQTL analysis and functional validation of candidate susceptibility genes for high-grade serous ovarian cancer. <i>Nature Communications</i> , 2015, 6, 8234.	12.8	63

#	ARTICLE	IF	CITATIONS
109	Benzo[a]pyrene Diol Epoxide and Bleomycin Sensitivity and Susceptibility to Cancer of Upper Aerodigestive Tract. <i>Journal of the National Cancer Institute</i> , 1998, 90, 1393-1399.	6.3	62
110	Nicotinic Acetylcholine Receptor Region on Chromosome 15q25 and Lung Cancer Risk Among African Americans: A Case-Control Study. <i>Journal of the National Cancer Institute</i> , 2010, 102, 1199-1205.	6.3	62
111	Genetic Variants in Cell Cycle Control Pathway Confer Susceptibility to Lung Cancer. <i>Clinical Cancer Research</i> , 2007, 13, 5974-5981.	7.0	61
112	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
113	Epidemiology and genetic susceptibility to bladder cancer. <i>BJU International</i> , 2008, 102, 1207-1215.	2.5	59
114	Novel genetic variants in the chromosome 5p15.33 region associate with lung cancer risk. <i>Carcinogenesis</i> , 2011, 32, 1493-1499.	2.8	59
115	Testicular germ cell tumor susceptibility associated with the UCK2 locus on chromosome 1q23. <i>Human Molecular Genetics</i> , 2013, 22, 2748-2753.	2.9	59
116	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
117	Sulfotransferase 1A1 (SULT1A1) polymorphism and bladder cancer risk: a case-control study. <i>Cancer Letters</i> , 2003, 202, 61-69.	7.2	58
118	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
119	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
120	Genetic Susceptibility to Tobacco Carcinogenesis. <i>Cancer Investigation</i> , 1999, 17, 645-659.	1.3	55
121	Nucleotide Excision Repair Pathway Genes and Oral Premalignant Lesions. <i>Clinical Cancer Research</i> , 2007, 13, 3753-3758.	7.0	55
122	Novel Susceptibility Loci for Second Primary Tumors/Recurrence in Head and Neck Cancer Patients: Large-Scale Evaluation of Genetic Variants. <i>Cancer Prevention Research</i> , 2009, 2, 617-624.	1.5	55
123	Common variation at 2q22.3 (ZEB2) influences the risk of renal cancer. <i>Human Molecular Genetics</i> , 2013, 22, 825-831.	2.9	54
124	Genetic variations of the PI3K-AKT-mTOR pathway and clinical outcome in muscle invasive and metastatic bladder cancer patients. <i>Carcinogenesis</i> , 2010, 31, 1387-1391.	2.8	53
125	Genome-Wide Association Study of Genetic Predictors of Overall Survival for Non-Small Cell Lung Cancer in Never Smokers. <i>Cancer Research</i> , 2013, 73, 4028-4038.	0.9	53
126	Genetic variants in cell cycle control pathway confer susceptibility to bladder cancer. <i>Cancer</i> , 2008, 112, 2467-2474.	4.1	52



#	ARTICLE	IF	CITATIONS
127	MicroRNA-related genetic variations as predictors for risk of second primary tumor and/or recurrence in patients with early-stage head and neck cancer. <i>Carcinogenesis</i> , 2010, 31, 2118-2123.	2.8	52
128	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
129	The somatic mutation landscape of premalignant colorectal adenoma. <i>Gut</i> , 2018, 67, 1299-1305.	12.1	52
130	Global and targeted circulating microRNA profiling of colorectal adenoma and colorectal cancer. <i>Cancer</i> , 2018, 124, 785-796.	4.1	52
131	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
132	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
133	Global and targeted serum metabolic profiling of colorectal cancer progression. <i>Cancer</i> , 2017, 123, 4066-4074.	4.1	51
134	A validated miRNA profile predicts response to therapy in esophageal adenocarcinoma. <i>Cancer</i> , 2014, 120, 3635-3641.	4.1	50
135	Profiling of Genetic Variations in Inflammation Pathway Genes in Relation to Bladder Cancer Predisposition. <i>Clinical Cancer Research</i> , 2008, 14, 2236-2244.	7.0	49
136	Genome-wide association study confirms lung cancer susceptibility loci on chromosomes 5p15 and 15q25 in an African-American population. <i>Lung Cancer</i> , 2016, 98, 33-42.	2.0	49
137	Metabolomics profiling in plasma samples from glioma patients correlates with tumor phenotypes. <i>Oncotarget</i> , 2016, 7, 20486-20495.	1.8	49
138	Association between asbestos exposure, cigarette smoking, myeloperoxidase (MPO) genotypes, and lung cancer risk. <i>American Journal of Industrial Medicine</i> , 2002, 42, 29-37.	2.1	48
139	Myeloperoxidase G-463A polymorphism and lung cancer: A HuGE Genetic Susceptibility to Environmental Carcinogens pooled analysis. <i>Genetics in Medicine</i> , 2007, 9, 67-73.	2.4	47
140	Genetic susceptibility to bladder cancer risk and outcome. <i>Personalized Medicine</i> , 2011, 8, 365-374.	1.5	46
141	GWAS-identified colorectal cancer susceptibility loci associated with clinical outcomes. <i>Carcinogenesis</i> , 2012, 33, 1327-1331.	2.8	46
142	Association of mitochondrial DNA copy number in peripheral blood leukocytes with risk of esophageal adenocarcinoma. <i>Carcinogenesis</i> , 2013, 34, 2521-2524.	2.8	46
143	Prognostic role of elevated mir-24-3p in breast cancer and its association with the metastatic process. <i>Oncotarget</i> , 2018, 9, 12868-12878.	1.8	46
144	Glutathione peroxidase 1 gene polymorphism and risk of recurrence in patients with superficial bladder cancer. <i>Urology</i> , 2005, 66, 769-774.	1.0	45

#	ARTICLE	IF	CITATIONS
145	Genetic Variations in the Sonic Hedgehog Pathway Affect Clinical Outcomes in Non-muscle-Invasive Bladder Cancer. <i>Cancer Prevention Research</i> , 2010, 3, 1235-1245.	1.5	45
146	Cyclin D1 gene polymorphism as a risk factor for oral premalignant lesions. <i>Carcinogenesis</i> , 2006, 27, 2034-2037.	2.8	44
147	Dietary Intake of Vegetables and Fruits and the Modification Effects of <i>GSTM1</i> and <i>NAT2</i> Genotypes on Bladder Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009, 18, 2090-2097.	2.5	44
148	Common Genetic Variation In Cellular Transport Genes and Epithelial Ovarian Cancer (EOC) Risk. <i>PLoS ONE</i> , 2015, 10, e0128106.	2.5	44
149	Mutagen Sensitivity and Genetic Variants in Nucleotide Excision Repair Pathway: Genotype-Phenotype Correlation. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007, 16, 2065-2071.	2.5	43
150	Germline genetic variations in drug action pathways predict clinical outcomes in advanced lung cancer treated with platinum-based chemotherapy. <i>Pharmacogenetics and Genomics</i> , 2008, 18, 955-965.	1.5	43
151	MicroRNA-Related Genetic Variants Associated with Clinical Outcomes in Early-Stage Non-small Cell Lung Cancer Patients. <i>Cancer Research</i> , 2013, 73, 1867-1875.	0.9	43
152	Fine mapping of MHC region in lung cancer highlights independent susceptibility loci by ethnicity. <i>Nature Communications</i> , 2018, 9, 3927.	12.8	43
153	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
154	Genome-wide association study of familial lung cancer. <i>Carcinogenesis</i> , 2018, 39, 1135-1140.	2.8	42
155	CYP2A6 reduced activity gene variants confer reduction in lung cancer risk in African American smokers—findings from two independent populations. <i>Carcinogenesis</i> , 2015, 36, 99-103.	2.8	41
156	Genetic Risk Can Be Decreased: Quitting Smoking Decreases and Delays Lung Cancer for Smokers With High and Low <i>CHRNA5</i> Risk Genotypes – A Meta-Analysis. <i>EBioMedicine</i> , 2016, 11, 219-226.	6.1	40
157	Immune checkpoint-related serum proteins and genetic variants predict outcomes of localized prostate cancer, a cohort study. <i>Cancer Immunology, Immunotherapy</i> , 2021, 70, 701-712.	4.2	40
158	Radiation-induced chromatid breaks as a predictor of breast cancer risk. <i>International Journal of Radiation Oncology Biology Physics</i> , 2001, 49, 533-537.	0.8	39
159	Case-Control Analysis of Dietary Folate and Risk of Bladder Cancer. <i>Nutrition and Cancer</i> , 2005, 53, 144-151.	2.0	39
160	Genetic Susceptibility to Renal Cell Carcinoma: The Role of DNA Double-Strand Break Repair Pathway. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2008, 17, 2366-2373.	2.5	39
161	Smoking-related Genomic Signatures in Non-small Cell Lung Cancer. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2008, 178, 1164-1172.	5.6	39
162	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

#	ARTICLE	IF	CITATIONS
163	Personal Permanent Hair Dye Use Is Not Associated with Bladder Cancer Risk: Evidence from a Case-Control Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006, 15, 1746-1749.	2.5	38
164	Genotypes, haplotypes and diplotypes of XPC and risk of bladder cancer. <i>Carcinogenesis</i> , 2006, 28, 698-703.	2.8	38
165	Genetic and intermediate phenotypic susceptibility markers of gastric cancer in Hispanic Americans: A case-control study. <i>Cancer</i> , 2014, 120, 3040-3048.	4.1	38
166	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
167	Dietary Carotenoids and Genetic Instability Modify Bladder Cancer Risk. <i>Journal of Nutrition</i> , 2004, 134, 3362-3369.	2.9	37
168	Reduced mitochondrial DNA copy number in peripheral blood leukocytes increases the risk of soft tissue sarcoma. <i>Carcinogenesis</i> , 2013, 34, 1039-1043.	2.8	37
169	Energy Balance, the PI3K-AKT-mTOR Pathway Genes, and the Risk of Bladder Cancer. <i>Cancer Prevention Research</i> , 2010, 3, 505-517.	1.5	36
170	Fine-mapping of the 5p15.33, 6p22.1-p21.31, and 15q25.1 Regions Identifies Functional and Histology-Specific Lung Cancer Susceptibility Loci in African-Americans. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013, 22, 251-260.	2.5	36
171	Chromosome instability in lymphocytes: a potential indicator of predisposition to oral premalignant lesions. <i>Cancer Research</i> , 2002, 62, 2813-8.	0.9	36
172	Microsomal epoxide hydrolase polymorphisms and lung cancer risk in non-Hispanic whites. <i>Molecular Carcinogenesis</i> , 2002, 33, 99-104.	2.7	35
173	Sensitivity to NNKOAc is associated with renal cancer risk. <i>Carcinogenesis</i> , 2009, 30, 706-710.	2.8	35
174	Genome-wide association studies of bladder cancer risk: a field synopsis of progress and potential applications. <i>Cancer and Metastasis Reviews</i> , 2009, 28, 269-280.	5.9	35
175	Long telomeres in peripheral blood leukocytes are associated with an increased risk of soft tissue sarcoma. <i>Cancer</i> , 2013, 119, 1885-1891.	4.1	35
176	ABO blood types and cancer risk—A cohort study of 339,432 subjects in Taiwan. <i>Cancer Epidemiology</i> , 2015, 39, 150-156.	1.9	35
177	Novel fluorescence <i>in situ</i> hybridization-based definition of bacille Calmette-Guérin (BCG) failure for use in enhancing recruitment into clinical trials of intravesical therapies. <i>BJU International</i> , 2016, 117, 754-760.	2.5	35
178	Bladder cancer risk as modified by family history and smoking. <i>Cancer</i> , 2006, 107, 705-711.	4.1	34
179	Association of Polymorphisms in Oxidative Stress Genes with Clinical Outcomes for Bladder Cancer Treated with Bacillus Calmette-Guérin. <i>PLoS ONE</i> , 2012, 7, e38533.	2.5	34
180	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

#	ARTICLE	IF	CITATIONS
181	Variants in Inflammation Genes Are Implicated in Risk of Lung Cancer in Never Smokers Exposed to Second-hand Smoke. <i>Cancer Discovery</i> , 2011, 1, 420-429.	9.4	33
182	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
183	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
184	Mutagen Sensitivity and Neoplastic Progression in Patients with Barrett's Esophagus: A Prospective Analysis. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2006, 15, 1935-1940.	2.5	32
185	Plasma vitamins E and A and risk of bladder cancer: a case-control analysis. <i>Cancer Causes and Control</i> , 2008, 19, 981-992.	1.8	32
186	Genetic variations in the transforming growth factor-beta pathway as predictors of survival in advanced non-small cell lung cancer. <i>Carcinogenesis</i> , 2011, 32, 1050-1056.	2.8	32
187	Short telomere lengths in peripheral blood leukocytes are associated with an increased risk of oral premalignant lesion and oral squamous cell carcinoma. <i>Cancer</i> , 2013, 119, 4277-4283.	4.1	32
188	Genetic Variants in the Wnt/ $\beta$ 2-Catenin Signaling Pathway as Indicators of Bladder Cancer Risk. <i>Journal of Urology</i> , 2015, 194, 1771-1776.	0.4	32
189	Mendelian Randomization and mediation analysis of leukocyte telomere length and risk of lung and head and neck cancers. <i>International Journal of Epidemiology</i> , 2019, 48, 751-766.	1.9	32
190	Serum insulin-like growth factor (IGF) and IGF-binding protein levels and risk of lung cancer: a case-control study nested in the beta-Carotene and Retinol Efficacy Trial Cohort. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2002, 11, 1413-8.	2.5	32
191	Energy Balance, Polymorphisms in the mTOR Pathway, and Renal Cell Carcinoma Risk. <i>Journal of the National Cancer Institute</i> , 2013, 105, 424-432.	6.3	31
192	The Prostate Cancer Susceptibility Variant rs2735839 Near <i>KLK3</i> Gene Is Associated with Aggressive Prostate Cancer and Can Stratify Gleason Score 7 Patients. <i>Clinical Cancer Research</i> , 2014, 20, 5133-5139.	7.0	31
193	Genetic Variations in Glutathione Pathway Genes Predict Cancer Recurrence in Patients Treated with Transurethral Resection and Bacillus Calmette-Guerin Instillation for Non-muscle Invasive Bladder Cancer. <i>Annals of Surgical Oncology</i> , 2015, 22, 4104-4110.	1.5	31
194	Serum MicroRNA-150 Predicts Prognosis for Early-Stage Non-Small Cell Lung Cancer and Promotes Tumor Cell Proliferation by Targeting Tumor Suppressor Gene <i>SRCIN1</i> . <i>Clinical Pharmacology and Therapeutics</i> , 2018, 103, 1061-1073.	4.7	31
195	Consensus report of the 8 and 9th Weinman Symposia on Gene x Environment Interaction in carcinogenesis: novel opportunities for precision medicine. <i>Cell Death and Differentiation</i> , 2018, 25, 1885-1904.	11.2	31
196	Cell cycle checkpoints, DNA damage/repair, and lung cancer risk. <i>Cancer Research</i> , 2005, 65, 349-57.	0.9	31
197	Copy number alterations detected as clonal hematopoiesis of indeterminate potential. <i>Blood Advances</i> , 2017, 1, 1031-1036.	5.2	30
198	Serum levels of insulin-like growth factor I and risk of squamous intraepithelial lesions of the cervix. <i>Clinical Cancer Research</i> , 2003, 9, 3356-61.	7.0	30

#	ARTICLE	IF	CITATIONS
199	ATM sequence variants associate with susceptibility to non-small cell lung cancer. <i>International Journal of Cancer</i> , 2007, 121, 2254-2259.	5.1	29
200	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
201	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
202	Genome-wide interaction study of smoking behavior and non-small cell lung cancer risk in Caucasian population. <i>Carcinogenesis</i> , 2018, 39, 336-346.	2.8	29
203	Circulating adipokine concentrations and risk of five obesity-related cancers: A Mendelian randomization study. <i>International Journal of Cancer</i> , 2021, 148, 1625-1636.	5.1	29
204	Genetic polymorphism in bladder cancer. <i>Frontiers in Bioscience - Landmark</i> , 2007, 12, 192.	3.0	29
205	Urinary tract diseases and bladder cancer risk: a case-control study. <i>Cancer Causes and Control</i> , 2007, 18, 839-845.	1.8	28
206	Network-Based Integration of GWAS and Gene Expression Identifies a <i>HOX</i> -Centric Network Associated with Serous Ovarian Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 1574-1584.	2.5	28
207	Cohort Profile: The Mexican American Mano a Mano Cohort. <i>International Journal of Epidemiology</i> , 2017, 46, e3-e3.	1.9	28
208	Modification of lung cancer susceptibility by green tea extract as measured by the comet assay. <i>Cancer Detection and Prevention</i> , 2002, 26, 411-418.	2.1	27
209	Interplay between mutagen sensitivity and epidemiological factors in modulating lung cancer risk. <i>International Journal of Cancer</i> , 2007, 120, 2687-2695.	5.1	27
210	Genetic polymorphisms in double-strand break DNA repair genes associated with risk of oral premalignant lesions. <i>European Journal of Cancer</i> , 2008, 44, 1603-1611.	2.8	27
211	Genetic Variations in the Regulator of G-Protein Signaling Genes Are Associated with Survival in Late-Stage Non-Small Cell Lung Cancer. <i>PLoS ONE</i> , 2011, 6, e21120.	2.5	27
212	Polymorphisms in the <i>XRCC1</i> gene modify survival of bladder cancer patients treated with chemotherapy. <i>International Journal of Cancer</i> , 2013, 133, 2004-2009.	5.1	27
213	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
214	Identification of 22 susceptibility loci associated with testicular germ cell tumors. <i>Nature Communications</i> , 2021, 12, 4487.	12.8	27
215	<i>XRCC3</i> genetic polymorphism, smoking, and lung carcinoma risk in minority populations. <i>Cancer</i> , 2003, 98, 1701-1706.	4.1	26
216	Predictors of esophageal cancer risk: Assessment of susceptibility to DNA damage using comet assay. <i>Genes Chromosomes and Cancer</i> , 2005, 44, 415-422.	2.8	26

#	ARTICLE	IF	CITATIONS
217	Systematic evaluation of apoptotic pathway gene polymorphisms and lung cancer risk. <i>Carcinogenesis</i> , 2012, 33, 1699-1706.	2.8	26
218	Intake of red meat and heterocyclic amines, metabolic pathway genes and bladder cancer risk. <i>International Journal of Cancer</i> , 2012, 131, 1892-1903.	5.1	26
219	Serum miR-331-3p predicts tumor recurrence in esophageal adenocarcinoma. <i>Scientific Reports</i> , 2018, 8, 14006.	3.3	26
220	Population-based targeted sequencing of 54 candidate genes identifies <i>PALB2</i> as a susceptibility gene for high-grade serous ovarian cancer. <i>Journal of Medical Genetics</i> , 2021, 58, 305-313.	3.2	26
221	Mitochondrial DNA copy number in peripheral blood leukocytes and the aggressiveness of localized prostate cancer. <i>Oncotarget</i> , 2015, 6, 41988-41996.	1.8	26
222	Ethnic differences in poly(ADP-ribose) polymerase pseudogene genotype distribution and association with lung cancer risk. <i>Carcinogenesis</i> , 1999, 20, 1465-1470.	2.8	25
223	Roles of tumor suppressor and telomere maintenance genes in cancer and aging—an epidemiological study. <i>Carcinogenesis</i> , 2005, 26, 1741-1747.	2.8	25
224	Genetic susceptibility to bladder cancer with an emphasis on gene-gene and gene-environmental interactions. <i>Current Opinion in Urology</i> , 2008, 18, 493-498.	1.8	25
225	Risk of urinary bladder cancer: a case-control analysis of industry and occupation. <i>BMC Cancer</i> , 2009, 9, 443.	2.6	25
226	Mitochondrial DNA copy number in peripheral blood leukocytes and the risk of clear cell renal cell carcinoma. <i>Carcinogenesis</i> , 2015, 36, 249-255.	2.8	25
227	Prognostic significance of promoter CpG island methylation of obesity-related genes in patients with nonmetastatic renal cell carcinoma. <i>Cancer</i> , 2017, 123, 3617-3627.	4.1	25
228	Genetic interaction analysis among oncogenesis-related genes revealed novel genes and networks in lung cancer development. <i>Oncotarget</i> , 2019, 10, 1760-1774.	1.8	25
229	Common Genetic Variation in Circadian Rhythm Genes and Risk of Epithelial Ovarian Cancer (EOC). <i>Journal of Genetics and Genome Research</i> , 2015, 2, .	0.3	25
230	The 19q12 Bladder Cancer GWAS Signal: Association with Cyclin E Function and Aggressive Disease. <i>Cancer Research</i> , 2014, 74, 5808-5818.	0.9	24
231	The Relationship between Native American Ancestry, Body Mass Index and Diabetes Risk among Mexican-Americans. <i>PLoS ONE</i> , 2015, 10, e0141260.	2.5	24
232	Common variants at the <i>CHEK2</i> gene locus and risk of epithelial ovarian cancer. <i>Carcinogenesis</i> , 2015, 36, 1341-1353.	2.8	24
233	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
234	Prospective analysis of DNA damage and repair markers of lung cancer risk from the Prostate, Lung, Colorectal and Ovarian (PLCO) Cancer Screening Trial. <i>Carcinogenesis</i> , 2011, 32, 69-73.	2.8	23

#	ARTICLE	IF	CITATIONS
235	Hypertension Susceptibility Loci are Associated with Anthracycline-related Cardiotoxicity in Long-term Childhood Cancer Survivors. <i>Scientific Reports</i> , 2017, 7, 9698.	3.3	23
236	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
237	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
238	BPDE Induced Lymphocytic Chromosome 3p Deletions May Predict Renal Cell Carcinoma Risk. <i>Journal of Urology</i> , 2008, 179, 2416-2421.	0.4	22
239	Cyclooxygenase-2 gene polymorphisms reduce the risk of oral premalignant lesions. <i>Cancer</i> , 2009, 115, 1498-1506.	4.1	22
240	Genome-wide association studies in bladder cancer: first results and potential relevance. <i>Current Opinion in Urology</i> , 2009, 19, 540-546.	1.8	22
241	Genetic variation in DNA-repair pathways and response to radiochemotherapy in esophageal adenocarcinoma: a retrospective cohort study of the Eastern Cooperative Oncology Group. <i>BMC Cancer</i> , 2011, 11, 176.	2.6	22
242	γ-H2AX level in peripheral blood lymphocytes as a risk predictor for bladder cancer. <i>Carcinogenesis</i> , 2013, 34, 2543-2547.	2.8	22
243	Epithelial-Mesenchymal Transition (EMT) Gene Variants and Epithelial Ovarian Cancer (EOC) Risk. <i>Genetic Epidemiology</i> , 2015, 39, 689-697.	1.3	22
244	Determination of proline in human serum by a robust LC-MS/MS method: application to identification of human metabolites as candidate biomarkers for esophageal cancer early detection and risk stratification. <i>Biomedical Chromatography</i> , 2015, 29, 570-577.	1.7	22
245	Genomic DNA Hypomethylation and Risk of Renal Cell Carcinoma: A Case-Control Study. <i>Clinical Cancer Research</i> , 2016, 22, 2074-2082.	7.0	22
246	Low serum testosterone is associated with tumor aggressiveness and poor prognosis in prostate cancer. <i>Oncology Letters</i> , 2017, 13, 1949-1957.	1.8	22
247	Chromosome 5 aberrations and genetic predisposition to lung cancer. , 1998, 79, 490-493.		21
248	Genetic variations in cell-cycle pathway and the risk of oral premalignant lesions. <i>Cancer</i> , 2008, 113, 2488-2495.	4.1	21
249	Germline prognostic markers for urinary bladder cancer: Obstacles and opportunities. <i>Urologic Oncology: Seminars and Original Investigations</i> , 2012, 30, 524-532.	1.6	21
250	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
251	Acculturation and Diabetes Risk in the Mexican American Mano a Mano Cohort. <i>American Journal of Public Health</i> , 2016, 106, 547-549.	2.7	21
252	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

#	ARTICLE	IF	CITATIONS
253	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
254	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
255	Genetic variations in regulator of G-protein signaling genes as susceptibility loci for second primary tumor/recurrence in head and neck squamous cell carcinoma. <i>Carcinogenesis</i> , 2010, 31, 1755-1761.	2.8	20
256	A Genome-Wide Association Study of Renal Cell Carcinoma among African Americans. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 209-214.	2.5	20
257	Increased leukocyte mitochondrial DNA copy number is associated with oral premalignant lesions: an epidemiology study. <i>Carcinogenesis</i> , 2014, 35, 1760-1764.	2.8	20
258	Coinherited genetics of multiple myeloma and its precursor, monoclonal gammopathy of undetermined significance. <i>Blood Advances</i> , 2020, 4, 2789-2797.	5.2	20
259	Genetic Variations in the Transforming Growth Factor Beta Pathway as Predictors of Bladder Cancer Risk. <i>PLoS ONE</i> , 2012, 7, e51758.	2.5	20
260	Two-stage induced differentiation of OCT4+/Nanog+ stem-like cells in lung adenocarcinoma. <i>Oncotarget</i> , 2016, 7, 68360-68370.	1.8	20
261	Methyl-CpG-binding domain 2. <i>Cancer</i> , 2004, 100, 1853-1858.	4.1	19
262	Genetic variants of the Wnt signaling pathway as predictors of recurrence and survival in early-stage non-small cell lung cancer patients. <i>Carcinogenesis</i> , 2014, 35, 1284-1291.	2.8	19
263	Assessing the genetic architecture of epithelial ovarian cancer histological subtypes. <i>Human Genetics</i> , 2016, 135, 741-756.	3.8	19
264	Lower mitochondrial DNA copy number in peripheral blood leukocytes increases the risk of endometrial cancer. <i>Molecular Carcinogenesis</i> , 2016, 55, 1111-1117.	2.7	19
265	D-mannose: a Novel Prognostic Biomarker for Patients with Esophageal Adenocarcinoma. <i>Carcinogenesis</i> , 2017, 38, bgw207.	2.8	19
266	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
267	Prevalence of Aflatoxin-Associated TP53R249S Mutation in Hepatocellular Carcinoma in Hispanics in South Texas. <i>Cancer Prevention Research</i> , 2018, 11, 103-112.	1.5	19
268	Mutagen sensitivity in humans: A comparison between two nomenclature systems for recording chromatid breaks. <i>Cancer Genetics and Cytogenetics</i> , 1996, 87, 127-132.	1.0	18
269	Three Measures of Mutagen Sensitivity in a Cancer-Free Population. <i>Cancer Genetics and Cytogenetics</i> , 1999, 110, 65-69.	1.0	18
270	Case-control analysis of nucleotide excision repair pathway and the risk of renal cell carcinoma. <i>Carcinogenesis</i> , 2008, 29, 2112-2119.	2.8	18



#	ARTICLE	IF	CITATIONS
271	Joint Association of Genome-Wide Association Study-Identified Susceptibility Loci and Dietary Patterns in Risk of Renal Cell Carcinoma Among Non-Hispanic Whites. <i>American Journal of Epidemiology</i> , 2014, 180, 499-507.	3.4	18
272	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
273	Epigenetic analysis of microRNA genes in tumors from surgically resected lung cancer patients and association with survival. <i>Molecular Carcinogenesis</i> , 2015, 54, E45-51.	2.7	18
274	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
275	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>Oncolmmunology</i> , 2017, 6, e1265719.	4.6	18
276	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
277	A miR-SNP biomarker linked to an increased lung cancer survival by miRNA-mediated down-regulation of FZD4 expression and Wnt signaling. <i>Scientific Reports</i> , 2017, 7, 9029.	3.3	18
278	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
279	Discovery and fine-mapping of height loci via high-density imputation of GWASs in individuals of African ancestry. <i>American Journal of Human Genetics</i> , 2021, 108, 564-582.	6.2	18
280	Genetic Variants in Telomere-Maintenance Genes and Bladder Cancer Risk. <i>PLoS ONE</i> , 2012, 7, e30665.	2.5	18
281	Mutagen sensitivity as a marker of cancer susceptibility. <i>Journal of Cellular Biochemistry</i> , 1996, 63, 80-84.	2.6	17
282	Joint Effect of Mutagen Sensitivity and Insulin-Like Growth Factors in Predicting the Risk of Developing Secondary Primary Tumors and Tumor Recurrence in Patients with Head and Neck Cancer. <i>Clinical Cancer Research</i> , 2006, 12, 7194-7201.	7.0	17
283	Genetic variations in regulator of G-protein signaling (RGS) confer risk of bladder cancer. <i>Cancer</i> , 2013, 119, 1643-1651.	4.1	17
284	Mutational Profiles Reveal an Aberrant TGF- $\beta$ 2-CEA Regulated Pathway in Colon Adenomas. <i>PLoS ONE</i> , 2016, 11, e0153933.	2.5	17
285	Circulating obesity-driven biomarkers are associated with risk of clear cell renal cell carcinoma: a two-stage, case-control study. <i>Carcinogenesis</i> , 2019, 40, 1191-1197.	2.8	17
286	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
287	Functional variants in DCAF4 associated with lung cancer risk in European populations. <i>Carcinogenesis</i> , 2017, 38, 541-551.	2.8	16
288	Socio-demographic, Clinical, and Genetic Determinants of Quality of Life in Lung Cancer Patients. <i>Scientific Reports</i> , 2018, 8, 10640.	3.3	16

#	ARTICLE	IF	CITATIONS
289	Sleep duration and risk of cancer in the Mexican American Mano-a-Mano Cohort. <i>Sleep Health</i> , 2019, 5, 78-83.	2.5	16
290	Heritability of prostate cancer: a tale of rare variants and common single nucleotide polymorphisms. <i>Annals of Translational Medicine</i> , 2016, 4, 206-206.	1.7	16
291	Ionizing Radiation-Induced $\gamma$ -H2AX Activity in Whole Blood Culture and the Risk of Lung Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013, 22, 443-451.	2.5	15
292	Risk Assessment of Esophageal Adenocarcinoma Using $\gamma$ -H2AX Assay. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2013, 22, 1797-1804.	2.5	15
293	Evaluating the ovarian cancer gonadotropin hypothesis: A candidate gene study. <i>Gynecologic Oncology</i> , 2015, 136, 542-548.	1.4	15
294	Identification of lung cancer histology-specific variants applying Bayesian framework variant prioritization approaches within the TRICL and ILCCO consortia. <i>Carcinogenesis</i> , 2015, 36, 1314-1326.	2.8	15
295	Plasma MicroRNA signature predicting weight gain among Mexican-American women. <i>Obesity</i> , 2017, 25, 958-964.	3.0	15
296	Measurement of DNA damage in peripheral blood by the $\gamma$ -H2AX assay as predictor of colorectal cancer risk. <i>DNA Repair</i> , 2017, 53, 24-30.	2.8	15
297	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
298	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
299	Adult height is associated with increased risk of ovarian cancer: a Mendelian randomisation study. <i>British Journal of Cancer</i> , 2018, 118, 1123-1129.	6.4	15
300	Different Transmission Dynamics of Coronavirus Disease 2019 (COVID-19) and Influenza Suggest the Relative Efficiency of Isolation/Quarantine and Social Distancing Against COVID-19 in China. <i>Clinical Infectious Diseases</i> , 2021, 73, e4305-e4311.	5.8	15
301	Low-Glycemic Index Diets as an Intervention in Metabolic Diseases: A Systematic Review and Meta-Analysis. <i>Nutrients</i> , 2022, 14, 307.	4.1	15
302	Risk assessment of renal cell carcinoma using alkaline comet assay. <i>Cancer</i> , 2007, 110, 282-288.	4.1	14
303	Identification of polymorphisms in ultraconserved elements associated with clinical outcomes in locally advanced colorectal adenocarcinoma. <i>Cancer</i> , 2012, 118, 6188-6198.	4.1	14
304	Deficiency of cell cycle checkpoints and DNA repair system predispose individuals to esophageal cancer. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2006, 602, 143-150.	1.0	13
305	Predictors of Survival in Never-Smokers with Non-Small Cell Lung Cancer: A Large-Scale, Two-Phase Genetic Study. <i>Clinical Cancer Research</i> , 2012, 18, 5983-5991.	7.0	13
306	Association of leukocyte telomere length in peripheral blood leukocytes with endometrial cancer risk in Caucasian Americans. <i>Carcinogenesis</i> , 2015, 36, 1327-1332.	2.8	13

#	ARTICLE	IF	CITATIONS
307	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
308	Genetic polymorphisms in genes related to risk-taking behaviours predicting body mass index trajectory among Mexican American adolescents. <i>Pediatric Obesity</i> , 2017, 12, 356-362.	2.8	13
309	Genetic variants in telomere-maintenance genes are associated with ovarian cancer risk and outcome. <i>Journal of Cellular and Molecular Medicine</i> , 2017, 21, 510-518.	3.6	13
310	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
311	The impact of health education videos on general public's mental health and behavior during COVID-19. <i>Global Health Research and Policy</i> , 2021, 6, 37.	3.6	13
312	Inherited variants affecting RNA editing may contribute to ovarian cancer susceptibility: results from a large-scale collaboration. <i>Oncotarget</i> , 2016, 7, 72381-72394.	1.8	13
313	Benzo(a)pyrene Diol Epoxide-Induced Chromosome 9p21 Aberrations Are Associated with Increased Risk of Bladder Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2008, 17, 2445-2450.	2.5	12
314	Re: Short Telomere Length, Cancer Survival, and Cancer Risk in 47 102 Individuals. <i>Journal of the National Cancer Institute</i> , 2013, 105, 1157-1157.	6.3	12
315	Genetic variations in base excision repair pathway and risk of bladder cancer: A case-control study in the United States. <i>Molecular Carcinogenesis</i> , 2015, 54, 50-57.	2.7	12
316	Cohort Profile: The MD Anderson Cancer Patients and Survivors Cohort (MDA-CPSC). <i>International Journal of Epidemiology</i> , 2016, 45, 713-713f.	1.9	12
317	Cross-Cancer Genome-Wide Association Study of Endometrial Cancer and Epithelial Ovarian Cancer Identifies Genetic Risk Regions Associated with Risk of Both Cancers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 217-228.	2.5	12
318	Smoking and nasopharyngeal cancer: individual data meta-analysis of six prospective studies on 334 935 men. <i>International Journal of Epidemiology</i> , 2021, 50, 975-986.	1.9	12
319	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
320	Myeloperoxidase Promoter Region Polymorphism and Lung Cancer Risk. , 2003, 75, 121-134.		11
321	Gamma-radiation sensitivity and polymorphisms in RAD51L1 modulate glioma risk. <i>Carcinogenesis</i> , 2010, 31, 1762-1769.	2.8	11
322	A genetic variant near the PMAIP1/Noxa gene is associated with increased bleomycin sensitivity. <i>Human Molecular Genetics</i> , 2011, 20, 820-826.	2.9	11
323	Common genetic variants in cell cycle pathway are associated with survival in stage III-IV non-small-cell lung cancer. <i>Carcinogenesis</i> , 2011, 32, 1867-1871.	2.8	11
324	Development and validation of risk models and molecular diagnostics to permit personalized management of cancer. <i>Cancer</i> , 2014, 120, 11-19.	4.1	11

#	ARTICLE	IF	CITATIONS
325	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
326	Acculturation, sociodemographic and lifestyle factors associated with compliance with physical activity recommendations in the Mexican-American Mano A Mano cohort. <i>BMJ Open</i> , 2015, 5, e008302.	1.9	11
327	Investigation of Leukocyte Telomere Length and Genetic Variants in Chromosome 5p15.33 as Prognostic Markers in Lung Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2019, 28, 1228-1237.	2.5	11
328	Cross-Cancer Pleiotropic Associations with Lung Cancer Risk in African Americans. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2019, 28, 715-723.	2.5	11
329	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
330	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
331	Serum microRNAs as predictors of risk for non-muscle invasive bladder cancer. <i>Oncotarget</i> , 2018, 9, 14895-14908.	1.8	11
332	Association study of nicotinic acetylcholine receptor genes identifies a novel lung cancer susceptibility locus near CHRNA1 in African-Americans. <i>Oncotarget</i> , 2012, 3, 1428-1438.	1.8	11
333	BPDE-induced lymphocytic 3p21.3 aberrations may predict head and neck carcinoma risk. <i>Cancer</i> , 2002, 95, 563-568.	4.1	10
334	Benzo[a]pyrene diol epoxide-induced 9p21 aberrations associated with genetic predisposition to bladder cancer. <i>Genes Chromosomes and Cancer</i> , 2004, 41, 330-338.	2.8	10
335	Mutagen-Induced Chromatid Breakage as a Marker of Cancer Risk. , 2005, 291, 059-068.		10
336	High Mutagen Sensitivity in Peripheral Blood Lymphocytes Predicts Poor Overall and Disease-Specific Survival in Patients with Stage III Non-Small Cell Lung Cancer Treated with Radiotherapy and Chemotherapy. <i>Clinical Cancer Research</i> , 2005, 11, 2894-2898.	7.0	10
337	Differential Induction in Telomerase Activity among Bladder Cancer Patients and Controls on $\beta$ -Radiation. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007, 16, 606-609.	2.5	10
338	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
339	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
340	Associations between language acculturation, age of immigration, and obesity in the Mexican American Mano A Mano cohort. <i>Obesity Research and Clinical Practice</i> , 2017, 11, 544-557.	1.8	10
341	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
342	Susceptibility loci of CNOT6 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

#	ARTICLE	IF	CITATIONS
343	Glycemic index, glycemic load and carbohydrate intake in association with risk of renal cell carcinoma. <i>Carcinogenesis</i> , 2017, 38, 1129-1135.	2.8	10
344	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
345	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
346	Germline Genetic Variants in the Wnt/ $\beta$ -Catenin Pathway as Predictors of Colorectal Cancer Risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 540-546.	2.5	9
347	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
348	Variants in genes encoding small GTPases and association with epithelial ovarian cancer susceptibility. <i>PLoS ONE</i> , 2018, 13, e0197561.	2.5	9
349	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
350	A case-control analysis of lymphocytic chromosome 9 aberrations in lung cancer. <i>International Journal of Cancer</i> , 2002, 102, 536-540.	5.1	7
351	Specific chromosome aberrations in peripheral blood lymphocytes are associated with risk of bladder cancer. <i>Genes Chromosomes and Cancer</i> , 2004, 41, 379-389.	2.8	7
352	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
353	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
354	Methylation of subtelomeric repeat D4Z4 in peripheral blood leukocytes is associated with biochemical recurrence in localized prostate cancer patients. <i>Carcinogenesis</i> , 2017, 38, 821-826.	2.8	7
355	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
356	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>Oncot Immunology</i> , 2019, 8, e1483303.	4.6	7
357	Reply to "Mosaic loss of chromosome Y in leukocytes matters". <i>Nature Genetics</i> , 2019, 51, 7-9.	21.4	7
358	Integration of multiomic annotation data to prioritize and characterize inflammation and immune-related risk variants in squamous cell lung cancer. <i>Genetic Epidemiology</i> , 2021, 45, 99-114.	1.3	7
359	High NAFLD fibrosis score in non-alcoholic fatty liver disease as a predictor of carotid plaque development: a retrospective cohort study based on regular health check-up data in China. <i>Annals of Medicine</i> , 2021, 53, 1621-1631.	3.8	7
360	Epidemiology of Renal Cell Carcinoma. , 2017, , 1-18.		7

#	ARTICLE	IF	CITATIONS
361	LINE-1 methylation in peripheral blood leukocytes and clinical characteristics and prognosis of prostate cancer patients. <i>Oncotarget</i> , 2017, 8, 94020-94027.	1.8	7
362	Pharmacogenetics in Esophageal Cancer. <i>Seminars in Oncology</i> , 2005, 32, 87-89.	2.2	6
363	The D2 Dopamine Receptor Gene and Nicotine Dependence Among Bladder Cancer Patients and Controls. <i>Behavior Genetics</i> , 2010, 40, 49-58.	2.1	6
364	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
365	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
366	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
367	Systematic analyses of regulatory variants in DNase I hypersensitive sites identified two novel lung cancer susceptibility loci. <i>Carcinogenesis</i> , 2019, 40, 432-440.	2.8	5
368	Assessment of variation in immunosuppressive pathway genes reveals TGFBR2 to be associated with risk of clear cell ovarian cancer. <i>Oncotarget</i> , 2016, 7, 69097-69110.	1.8	5
369	Lung carcinoma patients with a family history of cancer and lymphocyte primary chromosome 9 aberrations. , 1997, 79, 1527-1532.		4
370	The Role of Physical Activity in Harm Reduction among Betel Quid Chewers from a Prospective Cohort of 419,378 Individuals. <i>PLoS ONE</i> , 2016, 11, e0152246.	2.5	4
371	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
372	Converting health risks into loss of life years - a paradigm shift in clinical risk communication. <i>Aging</i> , 2021, 13, 21513-21525.	3.1	4
373	rs495139 in the TYMS-ENOSF1 Region and Risk of Ovarian Carcinoma of Mucinous Histology. <i>International Journal of Molecular Sciences</i> , 2018, 19, 2473.	4.1	3
374	Genetic variants in the inflammation pathway as predictors of recurrence and progression in non-muscle invasive bladder cancer treated with Bacillus Calmette-Guérin. <i>Oncotarget</i> , 2017, 8, 88782-88791.	1.8	3
375	Mobile Phone Use and its Association With Sitting Time and Meeting Physical Activity Recommendations in a Mexican American Cohort. <i>JMIR MHealth and UHealth</i> , 2016, 4, e54.	3.7	3
376	Irradiation-induced telomerase activity and the risk of lung cancer. <i>Cancer</i> , 2007, 109, 1157-1163.	4.1	2
377	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
378	Strategies to Identify Pharmacogenomic Biomarkers: Candidate Gene, Pathway-Based, and Genome-Wide Approaches. , 2008, , 353-370.		2

#	ARTICLE	IF	CITATIONS
379	Mutagen Sensitivity. , 2015, , 2957-2961.		2
380	Genetic variations in apoptosis pathway and the risk of ovarian cancer. Oncotarget, 2016, 7, 56737-56745.	1.8	2
381	“Sugar-Sweetened Beverages” Is an Independent Risk From Pancreatic Cancer: Based on Half a Million Asian Cohort Followed for 25 Years. Frontiers in Oncology, 2022, 12, 835901.	2.8	2
382	Cohort profile: The National Colorectal Cancer Cohort (NCRCC) study in China. BMJ Open, 2021, 11, e051397.	1.9	2
383	Multilevel-analysis identify a cis-expression quantitative trait locus associated with risk of renal cell carcinoma. Oncotarget, 2015, 6, 4097-4109.	1.8	1
384	Validation of plasma metabolites associated with breast cancer risk among Mexican Americans. Cancer Epidemiology, 2020, 69, 101826.	1.9	1
385	Trend Analysis and Intervention Effect Starting Point Detection of COVID-19 Epidemics Using Recalibrated Time Series Models “ Worldwide, 2020. China CDC Weekly, 2021, 3, 417-422.	2.3	1
386	Land use mix and leukocyte telomere length in Mexican Americans. Scientific Reports, 2021, 11, 19742.	3.3	1
387	Pharmacogenetics in Cancer Chemotherapy. , 2008, , 113-128.		1
388	Mutagen Sensitivity as Measured by Induced Chromatid Breakage as a Marker of Cancer Risk. Methods in Molecular Biology, 2014, 1105, 183-192.	0.9	1
389	Predictive Utility of Mortality by Aging Measures at Different Hierarchical Levels and the Response to Modifiable Life Style Factors: Implications for Geroprotective Programs. Frontiers in Medicine, 2022, 9, 831260.	2.6	1
390	Assessment of Insulin-Like Growth Factors and Mutagen Sensitivity as Predictors of Lung Cancer Risk. , 2003, 75, 279-288.		0
391	Biomarkers for Assessing Risk of Cancer. , 2015, , 317-330.e3.		0
392	Response. Journal of the National Cancer Institute, 2017, 109, .	6.3	0
393	Genetic variants in epithelial“mesenchymal transition genes as predictors of clinical outcomes in localized prostate cancer. Carcinogenesis, 2020, 41, 1057-1064.	2.8	0
394	Genetic determinants of multiple myeloma risk within the Wnt/beta-catenin signaling pathway. Cancer Epidemiology, 2021, 73, 101972.	1.9	0
395	Pharmacogenetics of Lung Cancer. , 2010, , 87-106.		0
396	Mutagen Sensitivity. , 2011, , 2409-2412.		0

#	ARTICLE	IF	CITATIONS
397	Mutagen Sensitivity. , 2015, , 1-4.		0
398	Clonal Hematopoiesis Increases Risk of Therapy-Related Myeloid Neoplasms. Blood, 2016, 128, 38-38.	1.4	0
399	A Public Health Perspective on Preventing and Controlling the Spread of Coronavirus Disease 2019. China CDC Weekly, 2020, 2, 237-240.	2.3	0
400	A whole-exome case-control association study to characterize the contribution of rare coding variation to pancreatic cancer risk. Human Genetics and Genomics Advances, 2022, 3, 100078.	1.7	0