

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

398 papers	20,262 citations	72 h-index	126 g-index
410 ext. papers	23,365 ext. citations	7.3 avg, IF	6.18 L-index

#	Paper	IF	Citations
398	Minimum amount of physical activity for reduced mortality and extended life expectancy: a prospective cohort study. <i>Lancet, The</i> , 2011 , 378, 1244-53	40	1123
397	Genome-wide association scan of tag SNPs identifies a susceptibility locus for lung cancer at 15q25.1. <i>Nature Genetics</i> , 2008 , 40, 616-22	36.3	1036
396	Intratumor heterogeneity in localized lung adenocarcinomas delineated by multiregion sequencing. <i>Science</i> , 2014 , 346, 256-9	33.3	659
395	Common 5p15.33 and 6p21.33 variants influence lung cancer risk. <i>Nature Genetics</i> , 2008 , 40, 1407-9	36.3	465
394	Detectable clonal mosaicism and its relationship to aging and cancer. <i>Nature Genetics</i> , 2012 , 44, 651-8	36.3	409
393	A multi-stage genome-wide association study of bladder cancer identifies multiple susceptibility loci. <i>Nature Genetics</i> , 2010 , 42, 978-84	36.3	408
392	Telomere dysfunction: a potential cancer predisposition factor. <i>Journal of the National Cancer Institute</i> , 2003 , 95, 1211-8	9.7	381
391	A risk model for prediction of lung cancer. <i>Journal of the National Cancer Institute</i> , 2007 , 99, 715-26	9.7	299
390	Genetic variation in the prostate stem cell antigen gene PSCA confers susceptibility to urinary bladder cancer. <i>Nature Genetics</i> , 2009 , 41, 991-5	36.3	270
389	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	36.3	246
388	Bladder cancer predisposition: a multigenic approach to DNA-repair and cell-cycle-control genes. <i>American Journal of Human Genetics</i> , 2006 , 78, 464-79	11	234
387	Rare variants of large effect in BRCA2 and CHEK2 affect risk of lung cancer. <i>Nature Genetics</i> , 2014 , 46, 736-41	36.3	228
386	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	6.4	223
385	Focus on bladder cancer. <i>Cancer Cell</i> , 2004 , 6, 111-6	24.3	221
384	Evaluation of genetic variants in microRNA-related genes and risk of bladder cancer. <i>Cancer Research</i> , 2008 , 68, 2530-7	10.1	215
383	Mitochondrial DNA content: its genetic heritability and association with renal cell carcinoma. <i>Journal of the National Cancer Institute</i> , 2008 , 100, 1104-12	9.7	206
382	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, 358-63	9.7	202

381	Genome-wide association study of renal cell carcinoma identifies two susceptibility loci on 2p21 and 11q13.3. <i>Nature Genetics</i> , 2011 , 43, 60-5	36.3	199
380	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, 1552-6	9.7	191
379	p53 Genotypes and Haplotypes Associated With Lung Cancer Susceptibility and Ethnicity. <i>Journal of the National Cancer Institute</i> , 2002 , 94, 681-90	9.7	191
378	Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. <i>Nature Genetics</i> , 2017 , 49, 680-691	36.3	190
377	Genetic variants and risk of lung cancer in never smokers: a genome-wide association study. <i>Lancet Oncology</i> , 2010 , 11, 321-30	21.7	190
376	Preleukaemic clonal haemopoiesis and risk of therapy-related myeloid neoplasms: a case-control study. <i>Lancet Oncology</i> , 2017 , 18, 100-111	21.7	189
375	Identification of six new susceptibility loci for invasive epithelial ovarian cancer. <i>Nature Genetics</i> , 2015 , 47, 164-71	36.3	177
374	From genotype to phenotype: correlating XRCC1 polymorphisms with mutagen sensitivity. <i>DNA Repair</i> , 2003 , 2, 901-8	4.3	167
373	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, 4980-95	5.6	159
372	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, 959-71	9.7	153
371	Genetic variations in radiation and chemotherapy drug action pathways predict clinical outcomes in esophageal cancer. <i>Journal of Clinical Oncology</i> , 2006 , 24, 3789-98	2.2	153
370	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, 857-71	2.2	142
369	Meta-analysis identifies four new loci associated with testicular germ cell tumor. <i>Nature Genetics</i> , 2013 , 45, 680-5	36.3	132
368	XPA polymorphism associated with reduced lung cancer risk and a modulating effect on nucleotide excision repair capacity. <i>Carcinogenesis</i> , 2003 , 24, 505-9	4.6	132
367	Genetic susceptibility to tobacco-related cancer. <i>Oncogene</i> , 2004 , 23, 6500-23	9.2	130
366	Familial aggregation of common sequence variants on 15q24-25.1 in lung cancer. <i>Journal of the National Cancer Institute</i> , 2008 , 100, 1326-30	9.7	126
365	Higher lung cancer risk for younger African-Americans with the Pro/Pro p53 genotype. <i>Carcinogenesis</i> , 1995 , 16, 2205-8	4.6	124
364	Polymorphisms in inflammation genes and bladder cancer: from initiation to recurrence, progression, and survival. <i>Journal of Clinical Oncology</i> , 2005 , 23, 5746-56	2.2	123

363	PALB2, CHEK2 and ATM rare variants and cancer risk: data from COGS. <i>Journal of Medical Genetics</i> , 2016 , 53, 800-811	5.8	121
362	Energy stress-induced lncRNA FILNC1 represses c-Myc-mediated energy metabolism and inhibits renal tumor development. <i>Nature Communications</i> , 2017 , 8, 783	17.4	120
361	Genetic susceptibility to lung cancer: the role of DNA damage and repair. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2003 , 12, 689-98	4	118
360	An expanded risk prediction model for lung cancer. <i>Cancer Prevention Research</i> , 2008 , 1, 250-4	3.2	114
359	Genetic variations in microRNA-related genes are associated with survival and recurrence in patients with renal cell carcinoma. <i>Carcinogenesis</i> , 2010 , 31, 1805-12	4.6	110
358	MicroRNA expression signatures in Barrett's esophagus and esophageal adenocarcinoma. <i>Clinical Cancer Research</i> , 2009 , 15, 5744-52	12.9	110
357	Genome-wide association study identifies multiple susceptibility loci for diffuse large B cell lymphoma. <i>Nature Genetics</i> , 2014 , 46, 1233-8	36.3	108
356	Genetic variants on 15q25.1, smoking, and lung cancer: an assessment of mediation and interaction. <i>American Journal of Epidemiology</i> , 2012 , 175, 1013-20	3.8	108
355	Genetic variants in MicroRNA biosynthesis pathways and binding sites modify ovarian cancer risk, survival, and treatment response. <i>Cancer Research</i> , 2010 , 70, 9765-76	10.1	105
354	Genome-wide association study identifies multiple loci associated with bladder cancer risk. <i>Human Molecular Genetics</i> , 2014 , 23, 1387-98	5.6	101
353	Polymorphisms in DNA repair genes, smoking, and bladder cancer risk: findings from the international consortium of bladder cancer. <i>Cancer Research</i> , 2009 , 69, 6857-64	10.1	94
352	An evolutionary perspective on single-nucleotide polymorphism screening in molecular cancer epidemiology. <i>Cancer Research</i> , 2004 , 64, 2251-7	10.1	90
351	Sulfotransferase (SULT) 1A1 polymorphism as a predisposition factor for lung cancer: a case-control analysis. <i>Lung Cancer</i> , 2002 , 35, 137-42	5.9	88
350	Mosaic loss of chromosome Y is associated with common variation near TCL1A. <i>Nature Genetics</i> , 2016 , 48, 563-8	36.3	87
349	Prognostic significance of pretreatment serum levels of albumin, LDH and total bilirubin in patients with non-metastatic breast cancer. <i>Carcinogenesis</i> , 2015 , 36, 243-8	4.6	83
348	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	3	83
347	Nucleotide excision repair gene polymorphisms and recurrence after treatment for superficial bladder cancer. <i>Clinical Cancer Research</i> , 2005 , 11, 1408-15	12.9	83
346	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, 4282-9	5.6	82

345	Hepatocellular carcinoma risk prediction model for the general population: the predictive power of transaminases. <i>Journal of the National Cancer Institute</i> , 2012 , 104, 1599-611	9.7	81
344	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, 84-94	11.4	81
343	Genetic variation in MicroRNA genes and risk of oral premalignant lesions. <i>Molecular Carcinogenesis</i> , 2010 , 49, 183-9	5	80
342	Molecular epidemiology, cancer-related symptoms, and cytokines pathway. <i>Lancet Oncology</i> , 2008 , 9, 777-85	21.7	79
341	Pathway-based serum microRNA profiling and survival in patients with advanced stage non-small cell lung cancer. <i>Cancer Research</i> , 2013 , 73, 4801-9	10.1	78
340	MicroRNA in the pathogenesis and prognosis of esophageal cancer. <i>Current Pharmaceutical Design</i> , 2013 , 19, 1292-300	3.3	78
339	Characterization of large structural genetic mosaicism in human autosomes. <i>American Journal of Human Genetics</i> , 2015 , 96, 487-97	11	77
338	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, 6616-33	5.6	77
337	MicroRNA expression signatures during malignant progression from Barrett's esophagus to esophageal adenocarcinoma. <i>Cancer Prevention Research</i> , 2013 , 6, 196-205	3.2	77
336	Association of vitamin D levels and risk of ovarian cancer: a Mendelian randomization study. <i>International Journal of Epidemiology</i> , 2016 , 45, 1619-1630	7.8	77
335	Genetic instability in bladder cancer assessed by the comet assay. <i>Journal of the National Cancer Institute</i> , 2003 , 95, 540-7	9.7	76
334	Genetic variants of myeloperoxidase and lung cancer risk. <i>Carcinogenesis</i> , 2000 , 21, 1163-1166	4.6	76
333	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	3.7	75
332	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, 801-7	4	75
331	Genome-wide association study identifies five susceptibility loci for follicular lymphoma outside the HLA region. <i>American Journal of Human Genetics</i> , 2014 , 95, 462-71	11	74
330	A genome-wide association study identifies a novel susceptibility locus for renal cell carcinoma on 12p11.23. <i>Human Molecular Genetics</i> , 2012 , 21, 456-62	5.6	74
329	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, 817-25	9.7	73
328	A myeloperoxidase polymorphism associated with reduced risk of lung cancer. <i>Lung Cancer</i> , 2002 , 37, 35-40	5.9	73

327	Genetic susceptibility to esophageal cancer: the role of the nucleotide excision repair pathway. <i>Carcinogenesis</i> , 2009 , 30, 785-92	4.6	72
326	Meta-analysis of genome-wide association studies discovers multiple loci for chronic lymphocytic leukemia. <i>Nature Communications</i> , 2016 , 7, 10933	17.4	70
325	Genetic variations in PI3K-AKT-mTOR pathway and bladder cancer risk. <i>Carcinogenesis</i> , 2009 , 30, 2047-52	4.6	70
324	An association between NQO1 genetic polymorphism and risk of bladder cancer. <i>Mutation Research - Genetic Toxicology and Environmental Mutagenesis</i> , 2003 , 536, 131-7	3	70
323	Analysis of Genomes and Transcriptomes of Hepatocellular Carcinomas Identifies Mutations and Gene Expression Changes in the Transforming Growth Factor- β Pathway. <i>Gastroenterology</i> , 2018 , 154, 195-210	13.3	68
322	Mutagen sensitivity has high heritability: evidence from a twin study. <i>Cancer Research</i> , 2006 , 66, 5993-6	10.1	68
321	Opposing effects of emphysema, hay fever, and select genetic variants on lung cancer risk. <i>American Journal of Epidemiology</i> , 2005 , 161, 412-22	3.8	68
320	Household transmission of COVID-19-a systematic review and meta-analysis. <i>Journal of Infection</i> , 2020 , 81, 979-997	18.9	68
319	Telomere length in peripheral blood leukocytes and lung cancer risk: a large case-control study in Caucasians. <i>Cancer Research</i> , 2014 , 74, 2476-86	10.1	67
318	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, 1176-86	3.2	67
317	Comprehensive T cell repertoire characterization of non-small cell lung cancer. <i>Nature Communications</i> , 2020 , 11, 603	17.4	67
316	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, 2342-8	7.5	66
315	Telomere dysfunction in peripheral lymphocytes as a potential predisposition factor for renal cancer. <i>Journal of Urology</i> , 2007 , 178, 1492-6	2.5	66
314	Modulation of DNA damage/DNA repair capacity by XPC polymorphisms. <i>DNA Repair</i> , 2008 , 7, 141-8	4.3	64
313	Mutagen sensitivity: a genetic predisposition factor for cancer. <i>Cancer Research</i> , 2007 , 67, 3493-5	10.1	64
312	Projecting individualized probabilities of developing bladder cancer in white individuals. <i>Journal of Clinical Oncology</i> , 2007 , 25, 4974-81	2.2	63
311	Dietary isothiocyanates, GSTM1, GSTT1, NAT2 polymorphisms and bladder cancer risk. <i>International Journal of Cancer</i> , 2007 , 120, 2208-13	7.5	62
310	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, 459-65	3.2	61

309	Matrix metalloproteinase polymorphisms and bladder cancer risk. <i>Cancer Research</i> , 2006 , 66, 11644-8	10.1	61
308	High-order interactions among genetic polymorphisms in nucleotide excision repair pathway genes and smoking in modulating bladder cancer risk. <i>Carcinogenesis</i> , 2007 , 28, 2160-5	4.6	61
307	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, 3988-95	12.9	61
306	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, 514-21	3.2	60
305	Female chromosome X mosaicism is age-related and preferentially affects the inactivated X chromosome. <i>Nature Communications</i> , 2016 , 7, 11843	17.4	59
304	Cancer risk associated with chronic diseases and disease markers: prospective cohort study. <i>BMJ, The</i> , 2018 , 360, k134	5.9	57
303	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	6.4	57
302	Genetic variants in cell cycle control pathway confer susceptibility to lung cancer. <i>Clinical Cancer Research</i> , 2007 , 13, 5974-81	12.9	57
301	Epidemiology and genetic susceptibility to bladder cancer. <i>BJU International</i> , 2008 , 102, 1207-15	5.6	56
300	Obesity, metabolic factors and risk of different histological types of lung cancer: A Mendelian randomization study. <i>PLoS ONE</i> , 2017 , 12, e0177875	3.7	56
299	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	0.7	55
298	High serum iron is associated with increased cancer risk. <i>Cancer Research</i> , 2014 , 74, 6589-97	10.1	55
297	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, 1199-205	9.7	54
296	Testicular germ cell tumor susceptibility associated with the UCK2 locus on chromosome 1q23. <i>Human Molecular Genetics</i> , 2013 , 22, 2748-53	5.6	53
295	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, 1393-9	9.7	53
294	Genetic variations of the PI3K-AKT-mTOR pathway and clinical outcome in muscle invasive and metastatic bladder cancer patients. <i>Carcinogenesis</i> , 2010 , 31, 1387-91	4.6	52
293	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, 617-24	3.2	52
292	Novel genetic variants in the chromosome 5p15.33 region associate with lung cancer risk. <i>Carcinogenesis</i> , 2011 , 32, 1493-9	4.6	52

291	Diabetes with early kidney involvement may shorten life expectancy by 16 years. <i>Kidney International</i> , 2017 , 92, 388-396	9.9	51
290	Pharmacogenomics of platinum-based chemotherapy in NSCLC. <i>Expert Opinion on Drug Metabolism and Toxicology</i> , 2009 , 5, 745-55	5.5	51
289	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		51
288	Genome-wide association study identifies multiple risk loci for renal cell carcinoma. <i>Nature Communications</i> , 2017 , 8, 15724	17.4	50
287	Sulfotransferase 1A1 (SULT1A1) polymorphism and bladder cancer risk: a case-control study. <i>Cancer Letters</i> , 2003 , 202, 61-9	9.9	50
286	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		50
285	Common variation at 2q22.3 (ZEB2) influences the risk of renal cancer. <i>Human Molecular Genetics</i> , 2013 , 22, 825-31	5.6	49
284	Nucleotide excision repair pathway genes and oral premalignant lesions. <i>Clinical Cancer Research</i> , 2007 , 13, 3753-8	12.9	49
283	Genetic susceptibility to tobacco carcinogenesis. <i>Cancer Investigation</i> , 1999 , 17, 645-59	2.1	48
282	Shared heritability and functional enrichment across six solid cancers. <i>Nature Communications</i> , 2019 , 10, 431	17.4	45
281	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, 2118-23	4.6	45
280	Genetic variants in cell cycle control pathway confer susceptibility to bladder cancer. <i>Cancer</i> , 2008 , 112, 2467-74	6.4	45
279	Adult body mass index and risk of ovarian cancer by subtype: a Mendelian randomization study. <i>International Journal of Epidemiology</i> , 2016 , 45, 884-95	7.8	45
278	A genome-wide association study of marginal zone lymphoma shows association to the HLA region. <i>Nature Communications</i> , 2015 , 6, 5751	17.4	44
277	Profiling of genetic variations in inflammation pathway genes in relation to bladder cancer predisposition. <i>Clinical Cancer Research</i> , 2008 , 14, 2236-44	12.9	44
276	A validated miRNA profile predicts response to therapy in esophageal adenocarcinoma. <i>Cancer</i> , 2014 , 120, 3635-41	6.4	42
275	Genetic variations in the sonic hedgehog pathway affect clinical outcomes in non-muscle-invasive bladder cancer. <i>Cancer Prevention Research</i> , 2010 , 3, 1235-45	3.2	42
274	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	8.1	42

273	Cyclin D1 gene polymorphism as a risk factor for oral premalignant lesions. <i>Carcinogenesis</i> , 2006 , 27, 2034-7	4.6	41
272	Mutagen sensitivity and genetic variants in nucleotide excision repair pathway: genotype-phenotype correlation. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2007 , 16, 2065-71	4	41
271	Glutathione peroxidase 1 gene polymorphism and risk of recurrence in patients with superficial bladder cancer. <i>Urology</i> , 2005 , 66, 769-74	1.6	41
270	Cis-eQTL analysis and functional validation of candidate susceptibility genes for high-grade serous ovarian cancer. <i>Nature Communications</i> , 2015 , 6, 8234	17.4	40
269	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, 4028-38	10.1	40
268	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, 955-65	1.9	40
267	Association between asbestos exposure, cigarette smoking, myeloperoxidase (MPO) genotypes, and lung cancer risk. <i>American Journal of Industrial Medicine</i> , 2002 , 42, 29-37	2.7	40
266	Genetically predicted longer telomere length is associated with increased risk of B-cell lymphoma subtypes. <i>Human Molecular Genetics</i> , 2016 , 25, 1663-76	5.6	39
265	GWAS-identified colorectal cancer susceptibility loci associated with clinical outcomes. <i>Carcinogenesis</i> , 2012 , 33, 1327-31	4.6	39
264	Global and targeted serum metabolic profiling of colorectal cancer progression. <i>Cancer</i> , 2017 , 123, 4066-4074	10.7	38
263	MicroRNA-related genetic variants associated with clinical outcomes in early-stage non-small cell lung cancer patients. <i>Cancer Research</i> , 2013 , 73, 1867-75	10.1	38
262	Dietary intake of vegetables and fruits and the modification effects of GSTM1 and NAT2 genotypes on bladder cancer risk. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2009 , 18, 2090-7	4	38
261	The influence of obesity-related factors in the etiology of renal cell carcinoma-A mendelian randomization study. <i>PLoS Medicine</i> , 2019 , 16, e1002724	11.6	38
260	Radiation-induced chromatid breaks as a predictor of breast cancer risk. <i>International Journal of Radiation Oncology Biology Physics</i> , 2001 , 49, 533-7	4	37
259	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	12.9	36
258	Genotypes, haplotypes and diplotypes of XPC and risk of bladder cancer. <i>Carcinogenesis</i> , 2007 , 28, 698-703	4.3	36
257	Global and targeted circulating microRNA profiling of colorectal adenoma and colorectal cancer. <i>Cancer</i> , 2018 , 124, 785-796	6.4	36
256	Genetic susceptibility to bladder cancer risk and outcome. <i>Personalized Medicine</i> , 2011 , 8, 365-374	2.2	35

255	Metabolomics profiling in plasma samples from glioma patients correlates with tumor phenotypes. <i>Oncotarget</i> , 2016 , 7, 20486-95	3.3	35
254	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	7.8	34
253	Long telomeres in peripheral blood leukocytes are associated with an increased risk of soft tissue sarcoma. <i>Cancer</i> , 2013 , 119, 1885-91	6.4	34
252	Association of mitochondrial DNA copy number in peripheral blood leukocytes with risk of esophageal adenocarcinoma. <i>Carcinogenesis</i> , 2013 , 34, 2521-4	4.6	34
251	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	4.6	33
250	The somatic mutation landscape of premalignant colorectal adenoma. <i>Gut</i> , 2018 , 67, 1299-1305	19.2	33
249	Reduced mitochondrial DNA copy number in peripheral blood leukocytes increases the risk of soft tissue sarcoma. <i>Carcinogenesis</i> , 2013 , 34, 1039-43	4.6	33
248	Genetic susceptibility to renal cell carcinoma: the role of DNA double-strand break repair pathway. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2008 , 17, 2366-73	4	33
247	Smoking-related genomic signatures in non-small cell lung cancer. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2008 , 178, 1164-72	10.2	33
246	Case-control analysis of dietary folate and risk of bladder cancer. <i>Nutrition and Cancer</i> , 2005 , 53, 144-51	2.8	33
245	Genetic and intermediate phenotypic susceptibility markers of gastric cancer in Hispanic Americans: a case-control study. <i>Cancer</i> , 2014 , 120, 3040-8	6.4	32
244	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, 1746-9	4	32
243	Prognostic role of elevated mir-24-3p in breast cancer and its association with the metastatic process. <i>Oncotarget</i> , 2018 , 9, 12868-12878	3.3	32
242	Depressive symptoms and short telomere length are associated with increased mortality in bladder cancer patients. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015 , 24, 336-43	4	31
241	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	24.3	31
240	Dietary carotenoids and genetic instability modify bladder cancer risk. <i>Journal of Nutrition</i> , 2004 , 134, 3362-9	4.1	31
239	Microsomal epoxide hydrolase polymorphisms and lung cancer risk in non-Hispanic whites. <i>Molecular Carcinogenesis</i> , 2002 , 33, 99-104	5	31
238	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	5.9	31

237	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	4	31
236	Genetic Variants in the Wnt/ECatenin Signaling Pathway as Indicators of Bladder Cancer Risk. <i>Journal of Urology</i> , 2015 , 194, 1771-6	2.5	30
235	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, 1050-6	4.6	30
234	Cell cycle checkpoints, DNA damage/repair, and lung cancer risk. <i>Cancer Research</i> , 2005 , 65, 349-57	10.1	30
233	Identification of susceptibility pathways for the role of chromosome 15q25.1 in modifying lung cancer risk. <i>Nature Communications</i> , 2018 , 9, 3221	17.4	29
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