

# Zhaoming Wang

## List of Publications by Citations

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210  
papers

21,353  
citations

65  
h-index

145  
g-index

230  
ext. papers

25,719  
ext. citations

12.5  
avg, IF

4.97  
L-index

#	Paper	IF	Citations
210	Genetic studies of body mass index yield new insights for obesity biology. <i>Nature</i> , <b>2015</b> , 518, 197-206	50.4	2687
209	Defining the role of common variation in the genomic and biological architecture of adult human height. <i>Nature Genetics</i> , <b>2014</b> , 46, 1173-86	36.3	1339
208	A genome-wide association study identifies alleles in FGFR2 associated with risk of sporadic postmenopausal breast cancer. <i>Nature Genetics</i> , <b>2007</b> , 39, 870-4	36.3	1240
207	Genome-wide association study of prostate cancer identifies a second risk locus at 8q24. <i>Nature Genetics</i> , <b>2007</b> , 39, 645-9	36.3	979
206	Multiple loci identified in a genome-wide association study of prostate cancer. <i>Nature Genetics</i> , <b>2008</b> , 40, 310-5	36.3	787
205	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , <b>2017</b> , 551, 92-94	50.4	643
204	A genome-wide association study identifies pancreatic cancer susceptibility loci on chromosomes 13q22.1, 1q32.1 and 5p15.33. <i>Nature Genetics</i> , <b>2010</b> , 42, 224-8	36.3	463
203	A multistage genome-wide association study in breast cancer identifies two new risk alleles at 1p11.2 and 14q24.1 (RAD51L1). <i>Nature Genetics</i> , <b>2009</b> , 41, 579-84	36.3	452
202	A genome-wide association study of lung cancer identifies a region of chromosome 5p15 associated with risk for adenocarcinoma. <i>American Journal of Human Genetics</i> , <b>2009</b> , 85, 679-91	11	442
201	Genome-wide meta-analysis identifies 11 new loci for anthropometric traits and provides insights into genetic architecture. <i>Nature Genetics</i> , <b>2013</b> , 45, 501-12	36.3	437
200	The genomic landscape of pediatric and young adult T-lineage acute lymphoblastic leukemia. <i>Nature Genetics</i> , <b>2017</b> , 49, 1211-1218	36.3	430
199	Detectable clonal mosaicism and its relationship to aging and cancer. <i>Nature Genetics</i> , <b>2012</b> , 44, 651-8	36.3	409
198	A multi-stage genome-wide association study of bladder cancer identifies multiple susceptibility loci. <i>Nature Genetics</i> , <b>2010</b> , 42, 978-84	36.3	408
197	A shared susceptibility locus in PLCE1 at 10q23 for gastric adenocarcinoma and esophageal squamous cell carcinoma. <i>Nature Genetics</i> , <b>2010</b> , 42, 764-7	36.3	403
196	Pan-cancer genome and transcriptome analyses of 1,699 paediatric leukaemias and solid tumours. <i>Nature</i> , <b>2018</b> , 555, 371-376	50.4	380
195	A meta-analysis of 87,040 individuals identifies 23 new susceptibility loci for prostate cancer. <i>Nature Genetics</i> , <b>2014</b> , 46, 1103-9	36.3	331
194	Sex-stratified genome-wide association studies including 270,000 individuals show sexual dimorphism in genetic loci for anthropometric traits. <i>PLoS Genetics</i> , <b>2013</b> , 9, e1003500	6	277

193	Diverse and Targetable Kinase Alterations Drive Histiocytic Neoplasms. <i>Cancer Discovery</i> , <b>2016</b> , 6, 154-65	4.4	269
192	The landscape of recombination in African Americans. <i>Nature</i> , <b>2011</b> , 476, 170-5	50.4	243
191	Genome-wide association analysis identifies new lung cancer susceptibility loci in never-smoking women in Asia. <i>Nature Genetics</i> , <b>2012</b> , 44, 1330-5	36.3	237
190	Rare variants of large effect in BRCA2 and CHEK2 affect risk of lung cancer. <i>Nature Genetics</i> , <b>2014</b> , 46, 736-41	36.3	228
189	Genome-wide association study identifies multiple susceptibility loci for pancreatic cancer. <i>Nature Genetics</i> , <b>2014</b> , 46, 994-1000	36.3	226
188	Rare missense variants in POT1 predispose to familial cutaneous malignant melanoma. <i>Nature Genetics</i> , <b>2014</b> , 46, 482-6	36.3	220
187	The Influence of Age and Sex on Genetic Associations with Adult Body Size and Shape: A Large-Scale Genome-Wide Interaction Study. <i>PLoS Genetics</i> , <b>2015</b> , 11, e1005378	6	220
186	Identification of a new prostate cancer susceptibility locus on chromosome 8q24. <i>Nature Genetics</i> , <b>2009</b> , 41, 1055-7	36.3	201
185	Genome-wide association study of renal cell carcinoma identifies two susceptibility loci on 2p21 and 11q13.3. <i>Nature Genetics</i> , <b>2011</b> , 43, 60-5	36.3	199
184	Genome-wide association analyses of esophageal squamous cell carcinoma in Chinese identify multiple susceptibility loci and gene-environment interactions. <i>Nature Genetics</i> , <b>2012</b> , 44, 1090-7	36.3	196
183	A meta-analysis identifies new loci associated with body mass index in individuals of African ancestry. <i>Nature Genetics</i> , <b>2013</b> , 45, 690-6	36.3	192
182	Genome-wide association study of glioma and meta-analysis. <i>Human Genetics</i> , <b>2012</b> , 131, 1877-88	6.3	191
181	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , <b>2017</b> , 49, 1767-1778	36.3	186
180	Distribution of allele frequencies and effect sizes and their interrelationships for common genetic susceptibility variants. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2011</b> , 108, 18026-31	11.5	185
179	The OncoArray Consortium: A Network for Understanding the Genetic Architecture of Common Cancers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2017</b> , 26, 126-135	4	183
178	Genome-wide association study of prostate cancer in men of African ancestry identifies a susceptibility locus at 17q21. <i>Nature Genetics</i> , <b>2011</b> , 43, 570-3	36.3	171
177	Genome-wide association study of glioma subtypes identifies specific differences in genetic susceptibility to glioblastoma and non-glioblastoma tumors. <i>Nature Genetics</i> , <b>2017</b> , 49, 789-794	36.3	163
176	Genome-wide association study identifies two susceptibility loci for osteosarcoma. <i>Nature Genetics</i> , <b>2013</b> , 45, 799-803	36.3	156

175	Genome-wide association study identifies multiple risk loci for chronic lymphocytic leukemia. <i>Nature Genetics</i> , <b>2013</b> , 45, 868-76	36.3	147
174	A meta-analysis of genome-wide association studies of breast cancer identifies two novel susceptibility loci at 6q14 and 20q11. <i>Human Molecular Genetics</i> , <b>2012</b> , 21, 5373-84	5.6	143
173	Genome-wide association study identifies new prostate cancer susceptibility loci. <i>Human Molecular Genetics</i> , <b>2011</b> , 20, 3867-75	5.6	143
172	Meta-analysis identifies four new loci associated with testicular germ cell tumor. <i>Nature Genetics</i> , <b>2013</b> , 45, 680-5	36.3	132
171	Directional dominance on stature and cognition in diverse human populations. <i>Nature</i> , <b>2015</b> , 523, 459-462	60.4	119
170	Joint analysis of three genome-wide association studies of esophageal squamous cell carcinoma in Chinese populations. <i>Nature Genetics</i> , <b>2014</b> , 46, 1001-1006	36.3	112
169	Genome-wide association study identifies multiple susceptibility loci for diffuse large B cell lymphoma. <i>Nature Genetics</i> , <b>2014</b> , 46, 1233-8	36.3	108
168	Analysis of Heritability and Shared Heritability Based on Genome-Wide Association Studies for Thirteen Cancer Types. <i>Journal of the National Cancer Institute</i> , <b>2015</b> , 107, djv279	9.7	107
167	Genome-wide meta-analysis identifies five new susceptibility loci for pancreatic cancer. <i>Nature Communications</i> , <b>2018</b> , 9, 556	17.4	103
166	Characterizing the genetic basis of methylome diversity in histologically normal human lung tissue. <i>Nature Communications</i> , <b>2014</b> , 5, 3365	17.4	103
165	Genome-wide analysis of BMI in adolescents and young adults reveals additional insight into the effects of genetic loci over the life course. <i>Human Molecular Genetics</i> , <b>2013</b> , 22, 3597-607	5.6	103
164	Genome-wide association study identifies multiple loci associated with bladder cancer risk. <i>Human Molecular Genetics</i> , <b>2014</b> , 23, 1387-98	5.6	101
163	Population substructure and control selection in genome-wide association studies. <i>PLoS ONE</i> , <b>2008</b> , 3, e2551	3.7	99
162	Analysis of error profiles in deep next-generation sequencing data. <i>Genome Biology</i> , <b>2019</b> , 20, 50	18.3	98
161	A large multiethnic genome-wide association study of prostate cancer identifies novel risk variants and substantial ethnic differences. <i>Cancer Discovery</i> , <b>2015</b> , 5, 878-91	24.4	94
160	Meta-analysis of five genome-wide association studies identifies multiple new loci associated with testicular germ cell tumor. <i>Nature Genetics</i> , <b>2017</b> , 49, 1141-1147	36.3	85
159	Common genetic polymorphisms modify the effect of smoking on absolute risk of bladder cancer. <i>Cancer Research</i> , <b>2013</b> , 73, 2211-20	10.1	82
158	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> , <b>2011</b> , 20, 4282-9	5.6	82

157	Characterization of large structural genetic mosaicism in human autosomes. <i>American Journal of Human Genetics</i> , <b>2015</b> , 96, 487-97	11	77
156	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> , <b>2014</b> , 23, 6616-33	5.6	77
155	Genome-wide association study of gastric adenocarcinoma in Asia: a comparison of associations between cardia and non-cardia tumours. <i>Gut</i> , <b>2016</b> , 65, 1611-8	19.2	76
154	Two susceptibility loci identified for prostate cancer aggressiveness. <i>Nature Communications</i> , <b>2015</b> , 6, 6889	17.4	75
153	Genome-wide association study identifies five susceptibility loci for follicular lymphoma outside the HLA region. <i>American Journal of Human Genetics</i> , <b>2014</b> , 95, 462-71	11	74
152	A genome-wide association study identifies a novel susceptibility locus for renal cell carcinoma on 12p11.23. <i>Human Molecular Genetics</i> , <b>2012</b> , 21, 456-62	5.6	74
151	Prostate Cancer Susceptibility in Men of African Ancestry at 8q24. <i>Journal of the National Cancer Institute</i> , <b>2016</b> , 108,	9.7	72
150	A Genome-Wide Scan Identifies Variants in NFIB Associated with Metastasis in Patients with Osteosarcoma. <i>Cancer Discovery</i> , <b>2015</b> , 5, 920-31	24.4	71
149	Meta-analysis of genome-wide association studies discovers multiple loci for chronic lymphocytic leukemia. <i>Nature Communications</i> , <b>2016</b> , 7, 10933	17.4	70
148	Identification of nine new susceptibility loci for endometrial cancer. <i>Nature Communications</i> , <b>2018</b> , 9, 3166	17.4	70
147	Three new pancreatic cancer susceptibility signals identified on chromosomes 1q32.1, 5p15.33 and 8q24.21. <i>Oncotarget</i> , <b>2016</b> , 7, 66328-66343	3.3	66
146	Cross-Cancer Genome-Wide Analysis of Lung, Ovary, Breast, Prostate, and Colorectal Cancer Reveals Novel Pleiotropic Associations. <i>Cancer Research</i> , <b>2016</b> , 76, 5103-14	10.1	66
145	Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. <i>Nature Communications</i> , <b>2016</b> , 7, 11375	17.4	64
144	Genetic Risk for Subsequent Neoplasms Among Long-Term Survivors of Childhood Cancer. <i>Journal of Clinical Oncology</i> , <b>2018</b> , 36, 2078-2087	2.2	60
143	Female chromosome X mosaicism is age-related and preferentially affects the inactivated X chromosome. <i>Nature Communications</i> , <b>2016</b> , 7, 11843	17.4	59
142	A genome-wide association study of prostate cancer in West African men. <i>Human Genetics</i> , <b>2014</b> , 133, 509-21	6.3	56
141	Genetic variants associated with longer telomere length are associated with increased lung cancer risk among never-smoking women in Asia: a report from the female lung cancer consortium in Asia. <i>International Journal of Cancer</i> , <b>2015</b> , 137, 311-9	7.5	55
140	Genotypic variants at 2q33 and risk of esophageal squamous cell carcinoma in China: a meta-analysis of genome-wide association studies. <i>Human Molecular Genetics</i> , <b>2012</b> , 21, 2132-41	5.6	55

139	Genome-wide association analysis implicates dysregulation of immunity genes in chronic lymphocytic leukaemia. <i>Nature Communications</i> , <b>2017</b> , 8, 14175	17.4	54
138	Genetic polymorphisms in the 9p21 region associated with risk of multiple cancers. <i>Carcinogenesis</i> , <b>2014</b> , 35, 2698-705	4.6	54
137	A comprehensive candidate gene approach identifies genetic variation associated with osteosarcoma. <i>BMC Cancer</i> , <b>2011</b> , 11, 209	4.8	54
136	Common variation in genes related to innate immunity and risk of adult glioma. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2009</b> , 18, 1651-8	4	54
135	Testicular germ cell tumor susceptibility associated with the UCK2 locus on chromosome 1q23. <i>Human Molecular Genetics</i> , <b>2013</b> , 22, 2748-53	5.6	53
134	Identification of new susceptibility loci for gastric non-cardia adenocarcinoma: pooled results from two Chinese genome-wide association studies. <i>Gut</i> , <b>2017</b> , 66, 581-587	19.2	51
133	Premature Physiologic Aging as a Paradigm for Understanding Increased Risk of Adverse Health Across the Lifespan of Survivors of Childhood Cancer. <i>Journal of Clinical Oncology</i> , <b>2018</b> , 36, 2206-2215	2.2	51
132	Common genetic variation and risk of gallbladder cancer in India: a case-control genome-wide association study. <i>Lancet Oncology</i> , <b>2017</b> , 18, 535-544	21.7	50
131	Genome-wide association study identifies multiple risk loci for renal cell carcinoma. <i>Nature Communications</i> , <b>2017</b> , 8, 15724	17.4	50
130	Meta-analysis of gene-level associations for rare variants based on single-variant statistics. <i>American Journal of Human Genetics</i> , <b>2013</b> , 93, 236-48	11	49
129	Common variation at 2q22.3 (ZEB2) influences the risk of renal cancer. <i>Human Molecular Genetics</i> , <b>2013</b> , 22, 825-31	5.6	49
128	Genome-Wide Association Study to Identify Susceptibility Loci That Modify Radiation-Related Risk for Breast Cancer After Childhood Cancer. <i>Journal of the National Cancer Institute</i> , <b>2017</b> , 109,	9.7	46
127	TERT gene harbors multiple variants associated with pancreatic cancer susceptibility. <i>International Journal of Cancer</i> , <b>2015</b> , 137, 2175-83	7.5	46
126	Inherited variation at chromosome 12p13.33, including RAD52, influences the risk of squamous cell lung carcinoma. <i>Cancer Discovery</i> , <b>2012</b> , 2, 131-9	24.4	46
125	A genome-wide association study of marginal zone lymphoma shows association to the HLA region. <i>Nature Communications</i> , <b>2015</b> , 6, 5751	17.4	44
124	Fine mapping the KLK3 locus on chromosome 19q13.33 associated with prostate cancer susceptibility and PSA levels. <i>Human Genetics</i> , <b>2011</b> , 129, 675-85	6.3	41
123	Improved imputation of common and uncommon SNPs with a new reference set. <i>Nature Genetics</i> , <b>2011</b> , 44, 6-7	36.3	41
122	Genome-wide association study of prostate-specific antigen levels identifies novel loci independent of prostate cancer. <i>Nature Communications</i> , <b>2017</b> , 8, 14248	17.4	40

121	Association between GWAS-identified lung adenocarcinoma susceptibility loci and EGFR mutations in never-smoking Asian women, and comparison with findings from Western populations. <i>Human Molecular Genetics</i> , <b>2017</b> , 26, 454-465	5.6	40
120	Genetically predicted longer telomere length is associated with increased risk of B-cell lymphoma subtypes. <i>Human Molecular Genetics</i> , <b>2016</b> , 25, 1663-76	5.6	39
119	Two Novel Susceptibility Loci for Prostate Cancer in Men of African Ancestry. <i>Journal of the National Cancer Institute</i> , <b>2017</b> , 109,	9.7	38
118	The influence of obesity-related factors in the etiology of renal cell carcinoma-A mendelian randomization study. <i>PLoS Medicine</i> , <b>2019</b> , 16, e1002724	11.6	38
117	Using principal components of genetic variation for robust and powerful detection of gene-gene interactions in case-control and case-only studies. <i>American Journal of Human Genetics</i> , <b>2010</b> , 86, 331-42 <sup>11</sup>		37
116	Integration of multiethnic fine-mapping and genomic annotation to prioritize candidate functional SNPs at prostate cancer susceptibility regions. <i>Human Molecular Genetics</i> , <b>2015</b> , 24, 5603-18	5.6	35
115	Genome-wide interaction study of smoking and bladder cancer risk. <i>Carcinogenesis</i> , <b>2014</b> , 35, 1737-44	4.6	33
114	Large-scale pathway-based analysis of bladder cancer genome-wide association data from five studies of European background. <i>PLoS ONE</i> , <b>2012</b> , 7, e29396	3.7	33
113	Exome Array Analysis Identifies Variants in SPOCD1 and BTN3A2 That Affect Risk for Gastric Cancer. <i>Gastroenterology</i> , <b>2017</b> , 152, 2011-2021	13.3	32
112	Meta-analysis of genome-wide association studies identifies multiple lung cancer susceptibility loci in never-smoking Asian women. <i>Human Molecular Genetics</i> , <b>2016</b> , 25, 620-9	5.6	32
111	St. Jude Cloud: A Pediatric Cancer Genomic Data-Sharing Ecosystem. <i>Cancer Discovery</i> , <b>2021</b> , 11, 1082-1099	11.4	32
110	Genome-wide association study confirms lung cancer susceptibility loci on chromosomes 5p15 and 15q25 in an African-American population. <i>Lung Cancer</i> , <b>2016</b> , 98, 33-42	5.9	31
109	Sex-specific glioma genome-wide association study identifies new risk locus at 3p21.31 in females, and finds sex-differences in risk at 8q24.21. <i>Scientific Reports</i> , <b>2018</b> , 8, 7352	4.9	30
108	Personal history of diabetes, genetic susceptibility to diabetes, and risk of brain glioma: a pooled analysis of observational studies. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2014</b> , 23, 47-54	4	28
107	An integrated transcriptome and epigenome analysis identifies a novel candidate gene for pancreatic cancer. <i>BMC Medical Genomics</i> , <b>2013</b> , 6, 33	3.7	28
106	Genetic Variants Related to Longer Telomere Length are Associated with Increased Risk of Renal Cell Carcinoma. <i>European Urology</i> , <b>2017</b> , 72, 747-754	10.2	27
105	Functional characterization of a multi-cancer risk locus on chr5p15.33 reveals regulation of TERT by ZNF148. <i>Nature Communications</i> , <b>2017</b> , 8, 15034	17.4	26
104	Telomere length and variation in telomere biology genes in individuals with osteosarcoma. <i>International Journal of Molecular Epidemiology and Genetics</i> , <b>2011</b> , 2, 19-29	0.9	26

103	Pathway-based analysis of GWAs data identifies association of sex determination genes with susceptibility to testicular germ cell tumors. <i>Human Molecular Genetics</i> , <b>2014</b> , 23, 6061-8	5.6	25
102	Interactions between household air pollution and GWAS-identified lung cancer susceptibility markers in the Female Lung Cancer Consortium in Asia (FLCCA). <i>Human Genetics</i> , <b>2015</b> , 134, 333-41	6.3	24
101	Association between adult height, genetic susceptibility and risk of glioma. <i>International Journal of Epidemiology</i> , <b>2012</b> , 41, 1075-85	7.8	24
100	Large-scale fine mapping of the HNF1B locus and prostate cancer risk. <i>Human Molecular Genetics</i> , <b>2011</b> , 20, 3322-9	5.6	22
99	Genetic variants in fas signaling pathway genes and risk of gastric cancer. <i>International Journal of Cancer</i> , <b>2014</b> , 134, 822-31	7.5	21
98	Intraspinal rhabdoid meningioma metastasis to the liver. <i>Journal of Clinical Neuroscience</i> , <b>2011</b> , 18, 714-6	6.2	21
97	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> , <b>2016</b> , 25, 1203-14	5.6	20
96	Known glioma risk loci are associated with glioma with a family history of brain tumours -- a case-control gene association study. <i>International Journal of Cancer</i> , <b>2013</b> , 132, 2464-8	7.5	20
95	Sex-specific gene and pathway modeling of inherited glioma risk. <i>Neuro-Oncology</i> , <b>2019</b> , 21, 71-82	1	19
94	The 19q12 bladder cancer GWAS signal: association with cyclin E function and aggressive disease. <i>Cancer Research</i> , <b>2014</b> , 74, 5808-18	10.1	19
93	Clinical outcomes of Ex Vivo liver resection and liver autotransplantation for hepatic alveolar echinococcosis. <i>Journal of Huazhong University of Science and Technology [Medical Sciences]</i> , <b>2012</b> , 32, 598-600		19
92	Morphological classification of intraductal papillary neoplasm of the bile duct. <i>European Radiology</i> , <b>2018</b> , 28, 1568-1578	8	19
91	Shortened Leukocyte Telomere Length Associates with an Increased Prevalence of Chronic Health Conditions among Survivors of Childhood Cancer: A Report from the St. Jude Lifetime Cohort. <i>Clinical Cancer Research</i> , <b>2020</b> , 26, 2362-2371	12.9	18
90	Insight in glioma susceptibility through an analysis of 6p22.3, 12p13.33-12.1, 17q22-23.2 and 18q23 SNP genotypes in familial and non-familial glioma. <i>Human Genetics</i> , <b>2012</b> , 131, 1507-17	6.3	18
89	Cohort Profile: The St. Jude Lifetime Cohort Study (SJLIFE) for paediatric cancer survivors. <i>International Journal of Epidemiology</i> , <b>2021</b> , 50, 39-49	7.8	18
88	Meta-analysis of genome-wide association studies and functional assays decipher susceptibility genes for gastric cancer in Chinese populations. <i>Gut</i> , <b>2020</b> , 69, 641-651	19.2	18
87	A genome-wide association study of renal cell carcinoma among African Americans. <i>Cancer Epidemiology Biomarkers and Prevention</i> , <b>2014</b> , 23, 209-14	4	17
86	Genetic variants in epidermal growth factor receptor pathway genes and risk of esophageal squamous cell carcinoma and gastric cancer in a Chinese population. <i>PLoS ONE</i> , <b>2013</b> , 8, e68999	3.7	17



85	Characterization of SNPs associated with prostate cancer in men of Ashkenazic descent from the set of GWAS identified SNPs: impact of cancer family history and cumulative SNP risk prediction. <i>PLoS ONE</i> , <b>2013</b> , 8, e60083	3.7	17
84	Functional characterization of a chr13q22.1 pancreatic cancer risk locus reveals long-range interaction and allele-specific effects on DIS3 expression. <i>Human Molecular Genetics</i> , <b>2016</b> , 25, 4726-4738 <sup>5,6</sup>	5.6	17
83	Pediatric Cancer Variant Pathogenicity Information Exchange (PeCanPIE): a cloud-based platform for curating and classifying germline variants. <i>Genome Research</i> , <b>2019</b> , 29, 1555-1565	9.7	16
82	Association between MDR1 C3435T polymorphism and risk of breast cancer. <i>Gene</i> , <b>2013</b> , 532, 94-9	3.8	16
81	Genetic Modifiers of Progression-Free Survival in Never-Smoking Lung Adenocarcinoma Patients Treated with First-Line Tyrosine Kinase Inhibitors. <i>American Journal of Respiratory and Critical Care Medicine</i> , <b>2017</b> , 195, 663-673	10.2	16
80	Agnostic Pathway/Gene Set Analysis of Genome-Wide Association Data Identifies Associations for Pancreatic Cancer. <i>Journal of the National Cancer Institute</i> , <b>2019</b> , 111, 557-567	9.7	16
79	GWAS meta-analysis of 16 852 women identifies new susceptibility locus for endometrial cancer. <i>Human Molecular Genetics</i> , <b>2016</b> , 25, 2612-2620	5.6	15
78	Sex specific associations in genome wide association analysis of renal cell carcinoma. <i>European Journal of Human Genetics</i> , <b>2019</b> , 27, 1589-1598	5.3	15
77	A flexible Bayesian model for studying gene-environment interaction. <i>PLoS Genetics</i> , <b>2012</b> , 8, e1002482	6	15
76	Common variation at 1q24.1 (ALDH9A1) is a potential risk factor for renal cancer. <i>PLoS ONE</i> , <b>2015</b> , 10, e0122589	3.7	15
75	Genetic admixture and population substructure in Guanacaste Costa Rica. <i>PLoS ONE</i> , <b>2010</b> , 5, e13336	3.7	14
74	Subsequent Breast Cancer in Female Childhood Cancer Survivors in the St Jude Lifetime Cohort Study (SJLIFE). <i>Journal of Clinical Oncology</i> , <b>2019</b> , 37, 1647-1656	2.2	13
73	A High-risk Haplotype for Premature Menopause in Childhood Cancer Survivors Exposed to Gonadotoxic Therapy. <i>Journal of the National Cancer Institute</i> , <b>2018</b> , 110, 895-904	9.7	13
72	Mendelian randomization analyses suggest a role for cholesterol in the development of endometrial cancer. <i>International Journal of Cancer</i> , <b>2021</b> , 148, 307-319	7.5	13
71	Age-specific genome-wide association study in glioblastoma identifies increased proportion of slower grade glioma-like features associated with younger age. <i>International Journal of Cancer</i> , <b>2018</b> , 143, 2359-2366	7.5	13
70	Enrichment of heterozygous germline loss-of-function variants in pediatric osteosarcoma. <i>Journal of Physical Education and Sports Management</i> , <b>2019</b> , 5,	2.8	12
69	Further Confirmation of Germline Glioma Risk Variant rs78378222 in TP53 and Its Implication in Tumor Tissues via Integrative Analysis of TCGA Data. <i>Human Mutation</i> , <b>2015</b> , 36, 684-8	4.7	12
68	A two-platform design for next generation genome-wide association studies. <i>Genetic Epidemiology</i> , <b>2012</b> , 36, 400-8	2.6	12

67	Nodular pulmonary amyloidosis and obvious ossification due to primary pulmonary MALT lymphoma with extensive plasmacytic differentiation: Report of a rare case and review of the literature. <i>International Journal of Clinical and Experimental Pathology</i> , <b>2015</b> , 8, 7482-7	1.4	12
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14	Uterine leiomyoma with indolent B-lymphoblastic proliferation. <i>International Journal of Clinical and Experimental Pathology</i> , <b>2013</b> , 6, 1422-6	1.4	1

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