

Zhaoming Wang

List of Publications by Year in Descending Order

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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.

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	Genome-wide association studies identify novel genetic loci for epigenetic age acceleration among survivors of childhood cancer.. <i>Genome Medicine</i> , 2022 , 14, 32	14.4	1
209	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	4	7
208	Racial and Ethnic Disparities in Health Outcomes Among Long-Term Survivors of Childhood Cancer: A Scoping Review. <i>Frontiers in Public Health</i> , 2021 , 9, 741334	6	2
207	Cohort Profile: The St. Jude Lifetime Cohort Study (SJLIFE) for paediatric cancer survivors. <i>International Journal of Epidemiology</i> , 2021 , 50, 39-49	7.8	18
206	Progression of Frailty in Survivors of Childhood Cancer: A St. Jude Lifetime Cohort Report. <i>Journal of the National Cancer Institute</i> , 2021 , 113, 1415-1421	9.7	3
205	Short sleep duration and physical and psychological health outcomes among adult survivors of childhood cancer. <i>Pediatric Blood and Cancer</i> , 2021 , 68, e28988	3	2
204	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	11	7
203	Persistent variations of blood DNA methylation associated with treatment exposures and risk for cardiometabolic outcomes in long-term survivors of childhood cancer in the St. Jude Lifetime Cohort. <i>Genome Medicine</i> , 2021 , 13, 53	14.4	3
202	The Association of Mitochondrial Copy Number With Sarcopenia in Adult Survivors of Childhood Cancer. <i>Journal of the National Cancer Institute</i> , 2021 , 113, 1570-1580	9.7	1
201	A Pedigree Analysis and Clonal Correlations of the Coexistence of B-Cell Lymphoma and Histiocytic/Dendritic Cell Tumor. <i>International Journal of Surgical Pathology</i> , 2021 , 29, 906-914	1.2	
200	Accelerated aging and mortality in long-term survivors of childhood cancer: A report from the St. Jude Lifetime Cohort (SJLIFE).. <i>Journal of Clinical Oncology</i> , 2021 , 39, 10045-10045	2.2	0
199	Mendelian randomization analyses suggest a role for cholesterol in the development of endometrial cancer. <i>International Journal of Cancer</i> , 2021 , 148, 307-319	7.5	13
198	Epigenetic Age Acceleration and Chronic Health Conditions Among Adult Survivors of Childhood Cancer. <i>Journal of the National Cancer Institute</i> , 2021 , 113, 597-605	9.7	5
197	The Role of Gallstones in Gallbladder Cancer in India: A Mendelian Randomization Study. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021 , 30, 396-403	4	2
196	Genetic Variants Associated with Therapy-Related Cardiomyopathy among Childhood Cancer Survivors of African Ancestry. <i>Cancer Research</i> , 2021 , 81, 2556-2565	10.1	8
195	Genome-wide Association Studies Reveal Novel Locus With Sex-/Therapy-Specific Fracture Risk Effects in Childhood Cancer Survivors. <i>Journal of Bone and Mineral Research</i> , 2021 , 36, 685-695	6.3	2
194	The clinicopathological and molecular features of sinusoidal large B-cell lymphoma. <i>Modern Pathology</i> , 2021 , 34, 922-933	9.8	3

193	SequencErr: measuring and suppressing sequencer errors in next-generation sequencing data. <i>Genome Biology</i> , 2021 , 22, 37	18.3	3
192	Sub-multiplicative interaction between polygenic risk score and household coal use in relation to lung adenocarcinoma among never-smoking women in Asia. <i>Environment International</i> , 2021 , 147, 105975	12.9	5
191	Inflammation-related genes S100s, RNASE3, and CYBB and risk of leukemic transformation in patients with myelodysplastic syndrome with myelofibrosis. <i>Biomarker Research</i> , 2021 , 9, 53	8	
190	Genetic analyses of gynecological disease identify genetic relationships between uterine fibroids and endometrial cancer, and a novel endometrial cancer genetic risk region at the WNT4 1p36.12 locus. <i>Human Genetics</i> , 2021 , 140, 1353-1365	6.3	5
189	Polygenic Risk Score Improves Risk Stratification and Prediction of Subsequent Thyroid Cancer after Childhood Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021 , 30, 2096-2104	4	3
188	Online Platform to Assess Complex Social Relationships and Patient-Reported Outcomes Among Adolescent and Young Adult Cancer Survivors. <i>JCO Clinical Cancer Informatics</i> , 2021 , 5, 859-871	5.2	2
187	St. Jude Cloud: A Pediatric Cancer Genomic Data-Sharing Ecosystem. <i>Cancer Discovery</i> , 2021 , 11, 1082-1094	29.4	32
186	Generalizability of "GWAS Hits" in Clinical Populations: Lessons from Childhood Cancer Survivors. <i>American Journal of Human Genetics</i> , 2020 , 107, 636-653	11	7
185	Pathogenic Germline Mutations in DNA Repair Genes in Combination With Cancer Treatment Exposures and Risk of Subsequent Neoplasms Among Long-Term Survivors of Childhood Cancer. <i>Journal of Clinical Oncology</i> , 2020 , 38, 2728-2740	2.2	11
184	A Novel Locus Predicts Spermatogenic Recovery among Childhood Cancer Survivors Exposed to Alkylating Agents. <i>Cancer Research</i> , 2020 , 80, 3755-3764	10.1	3
183	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> , 2020 , 26, 2362-2371	12.9	18
182	Cardiomyopathy risk among childhood cancer survivors of African ancestry and its molecular mechanisms. <i>Journal of Clinical Oncology</i> , 2020 , 38, 10514-10514	2.2	0
181	Estimated number of adult survivors of childhood cancer in United States with cancer-predisposing germline variants. <i>Pediatric Blood and Cancer</i> , 2020 , 67, e28047	3	7
180	Molecular Mechanism of Telomere Length Dynamics and Its Prognostic Value in Pediatric Cancers. <i>Journal of the National Cancer Institute</i> , 2020 , 112, 756-764	9.7	4
179	Meta-analysis of genome-wide association studies and functional assays decipher susceptibility genes for gastric cancer in Chinese populations. <i>Gut</i> , 2020 , 69, 641-651	19.2	18
178	Tuberculosis infection and lung adenocarcinoma: Mendelian randomization and pathway analysis of genome-wide association study data from never-smoking Asian women. <i>Genomics</i> , 2020 , 112, 1223-1232	4.3	8
177	Pediatric Cancer Variant Pathogenicity Information Exchange (PeCanPIE): a cloud-based platform for curating and classifying germline variants. <i>Genome Research</i> , 2019 , 29, 1555-1565	9.7	16
176	Genome-Wide Association Study in Irradiated Childhood Cancer Survivors Identifies HTR2A for Subsequent Basal Cell Carcinoma. <i>Journal of Investigative Dermatology</i> , 2019 , 139, 2042-2045.e8	4.3	9

175	Subsequent Breast Cancer in Female Childhood Cancer Survivors in the St Jude Lifetime Cohort Study (SJLIFE). <i>Journal of Clinical Oncology</i> , 2019 , 37, 1647-1656	2.2	13
174	Analysis of error profiles in deep next-generation sequencing data. <i>Genome Biology</i> , 2019 , 20, 50	18.3	98
173	Genetically Determined Height and Risk of Non-hodgkin Lymphoma. <i>Frontiers in Oncology</i> , 2019 , 9, 1539-1553	5.3	1
172	Sex-specific gene and pathway modeling of inherited glioma risk. <i>Neuro-Oncology</i> , 2019 , 21, 71-82	1	19
171	Association of Germline BRCA2 Mutations With the Risk of Pediatric or Adolescent Non-Hodgkin Lymphoma. <i>JAMA Oncology</i> , 2019 , 5, 1362-1364	13.4	10
170	Sex specific associations in genome wide association analysis of renal cell carcinoma. <i>European Journal of Human Genetics</i> , 2019 , 27, 1589-1598	5.3	15
169	Enrichment of heterozygous germline loss-of-function variants in pediatric osteosarcoma. <i>Journal of Physical Education and Sports Management</i> , 2019 , 5,	2.8	12
168	Genome-wide association study using whole-genome sequencing to identify a novel locus associated with cardiomyopathy risk in adult survivors of childhood cancer: Utility of a two-stage analytic approach.. <i>Journal of Clinical Oncology</i> , 2019 , 37, 1516-1516	2.2	
167	Polygenic risk of subsequent thyroid cancer after childhood cancer: A report from St. Jude lifetime cohort (SJLIFE) and Childhood Cancer Survivor Study (CCSS).. <i>Journal of Clinical Oncology</i> , 2019 , 37, 10060-10060	2.2	10060
166	Whole-Genome Sequencing of Childhood Cancer Survivors Treated with Cranial Radiation Therapy Identifies 5p15.33 Locus for Stroke: A Report from the St. Jude Lifetime Cohort Study. <i>Clinical Cancer Research</i> , 2019 , 25, 6700-6708	12.9	6
165	Agnostic Pathway/Gene Set Analysis of Genome-Wide Association Data Identifies Associations for Pancreatic Cancer. <i>Journal of the National Cancer Institute</i> , 2019 , 111, 557-567	9.7	16
164	Clinical Significance of Prognostic Nutritional Index for Patients with Diffuse Large B-cell Lymphoma. <i>Nutrition and Cancer</i> , 2019 , 71, 569-574	2.8	8
163	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
162	Pan-cancer genome and transcriptome analyses of 1,699 paediatric leukaemias and solid tumours. <i>Nature</i> , 2018 , 555, 371-376	50.4	380
161	Subphenotype meta-analysis of testicular cancer genome-wide association study data suggests a role for RBFOX family genes in cryptorchidism susceptibility. <i>Human Reproduction</i> , 2018 , 33, 967-977	5.7	7
160	A High-risk Haplotype for Premature Menopause in Childhood Cancer Survivors Exposed to Gonadotoxic Therapy. <i>Journal of the National Cancer Institute</i> , 2018 , 110, 895-904	9.7	13
159	BOLD MRI to evaluate early development of renal injury in a rat model of diabetes. <i>Journal of International Medical Research</i> , 2018 , 46, 1391-1403	1.4	5
158	Genome-wide meta-analysis identifies five new susceptibility loci for pancreatic cancer. <i>Nature Communications</i> , 2018 , 9, 556	17.4	103

157	Atypical presentation of blastic plasmacytoid dendritic cell neoplasm: A potential diagnostic pitfall in nasal cavity. <i>Oral Surgery, Oral Medicine, Oral Pathology and Oral Radiology</i> , 2018 , 126, e212-e214	2	2
156	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> , 2018 , 8, 7352	4.9	30
155	Identification of nine new susceptibility loci for endometrial cancer. <i>Nature Communications</i> , 2018 , 9, 3166	17.4	70
154	Hepatic carcinoma with indolent T-lymphoblastic proliferation (iT-LBP). <i>International Journal of Clinical and Experimental Pathology</i> , 2018 , 11, 1674-1678	1.4	0
153	Expression of hypoxia-inducible factor-1a predicts benefit from rituximab plus cyclophosphamide, doxorubicin, vincristine, and prednisone in diffuse large B-cell lymphoma. <i>International Journal of Clinical and Experimental Pathology</i> , 2018 , 11, 4472-4482	1.4	
152	Morphological classification of intraductal papillary neoplasm of the bile duct. <i>European Radiology</i> , 2018 , 28, 1568-1578	8	19
151	Genetic Risk for Subsequent Neoplasms Among Long-Term Survivors of Childhood Cancer. <i>Journal of Clinical Oncology</i> , 2018 , 36, 2078-2087	2.2	60
150	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> , 2018 , 36, 2206-2215	2.2	51
149	Two high-risk susceptibility loci at 6p25.3 and 14q32.13 for Waldenström macroglobulinemia. <i>Nature Communications</i> , 2018 , 9, 4182	17.4	8
148	Polygenic Determinants for Subsequent Breast Cancer Risk in Survivors of Childhood Cancer: The St Jude Lifetime Cohort Study (SJLIFE). <i>Clinical Cancer Research</i> , 2018 , 24, 6230-6235	12.9	11
147	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> , 2018 , 143, 2359-2366	7.5	13
146	Identification of new susceptibility loci for gastric non-cardia adenocarcinoma: pooled results from two Chinese genome-wide association studies. <i>Gut</i> , 2017 , 66, 581-587	19.2	51
145	Genome-wide association study of prostate-specific antigen levels identifies novel loci independent of prostate cancer. <i>Nature Communications</i> , 2017 , 8, 14248	17.4	40
144	Exome Array Analysis Identifies Variants in SPOCD1 and BTN3A2 That Affect Risk for Gastric Cancer. <i>Gastroenterology</i> , 2017 , 152, 2011-2021	13.3	32
143	A functional SNP rs1892901 in FOSL1 is associated with gastric cancer in Chinese population. <i>Scientific Reports</i> , 2017 , 7, 41737	4.9	5
142	Genome-wide association analysis implicates dysregulation of immunity genes in chronic lymphocytic leukaemia. <i>Nature Communications</i> , 2017 , 8, 14175	17.4	54
141	Common genetic variation and risk of gallbladder cancer in India: a case-control genome-wide association study. <i>Lancet Oncology</i> , 2017 , 18, 535-544	21.7	50
140	Functional characterization of a multi-cancer risk locus on chr5p15.33 reveals regulation of TERT by ZNF148. <i>Nature Communications</i> , 2017 , 8, 15034	17.4	26

139	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> , 2017 , 109,	9.7	46
138	Potential Susceptibility Loci Identified for Renal Cell Carcinoma by Targeting Obesity-Related Genes. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017 , 26, 1436-1442	4	0
137	ICT1 predicts a poor survival and correlated with cell proliferation in diffuse large B-cell lymphoma. <i>Gene</i> , 2017 , 627, 255-262	3.8	6
136	Genome-wide association study identifies multiple risk loci for renal cell carcinoma. <i>Nature Communications</i> , 2017 , 8, 15724	17.4	50
135	Meta-analysis of five genome-wide association studies identifies multiple new loci associated with testicular germ cell tumor. <i>Nature Genetics</i> , 2017 , 49, 1141-1147	36.3	85
134	Genome-wide association study of glioma subtypes identifies specific differences in genetic susceptibility to glioblastoma and non-glioblastoma tumors. <i>Nature Genetics</i> , 2017 , 49, 789-794	36.3	163
133	Association analysis identifies 65 new breast cancer risk loci. <i>Nature</i> , 2017 , 551, 92-94	50.4	643
132	Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. <i>Nature Genetics</i> , 2017 , 49, 1767-1778	36.3	186
131	Genetic Variants Related to Longer Telomere Length are Associated with Increased Risk of Renal Cell Carcinoma. <i>European Urology</i> , 2017 , 72, 747-754	10.2	27
130	Lupus-related single nucleotide polymorphisms and risk of diffuse large B-cell lymphoma. <i>Lupus Science and Medicine</i> , 2017 , 4, e000187	4.6	10
129	GWAS follow-up study of esophageal squamous cell carcinoma identifies potential genetic loci associated with family history of upper gastrointestinal cancer. <i>Scientific Reports</i> , 2017 , 7, 4642	4.9	7
128	The genomic landscape of pediatric and young adult T-lineage acute lymphoblastic leukemia. <i>Nature Genetics</i> , 2017 , 49, 1211-1218	36.3	430
127	Increasing mapping precision of genome-wide association studies: to genotype and impute, sequence, or both?. <i>Genome Biology</i> , 2017 , 18, 118	18.3	8
126	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> , 2017 , 195, 663-673	10.2	16
125	The OncoArray Consortium: A Network for Understanding the Genetic Architecture of Common Cancers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017 , 26, 126-135	4	183
124	Two Novel Susceptibility Loci for Prostate Cancer in Men of African Ancestry. <i>Journal of the National Cancer Institute</i> , 2017 , 109,	9.7	38
123	A high-risk genetic profile for premature menopause (PM) in childhood cancer survivors (CCS) exposed to gonadotoxic therapy: A report from the St. Jude Lifetime Cohort (SJLIFE) and Childhood Cancer Survivor Study (CCSS).. <i>Journal of Clinical Oncology</i> , 2017 , 35, 10502-10502	2.2	1
122	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> , 2017 , 26, 454-465	5.6	40

121	Genome-wide association study of gastric adenocarcinoma in Asia: a comparison of associations between cardia and non-cardia tumours. <i>Gut</i> , 2016 , 65, 1611-8	19.2	76
120	Female chromosome X mosaicism is age-related and preferentially affects the inactivated X chromosome. <i>Nature Communications</i> , 2016 , 7, 11843	17.4	59
119	Identification of four novel susceptibility loci for oestrogen receptor negative breast cancer. <i>Nature Communications</i> , 2016 , 7, 11375	17.4	64
118	Meta-analysis of genome-wide association studies discovers multiple loci for chronic lymphocytic leukemia. <i>Nature Communications</i> , 2016 , 7, 10933	17.4	70
117	Meta-analysis of genome-wide association studies identifies multiple lung cancer susceptibility loci in never-smoking Asian women. <i>Human Molecular Genetics</i> , 2016 , 25, 620-9	5.6	32
116	Diverse and Targetable Kinase Alterations Drive Histiocytic Neoplasms. <i>Cancer Discovery</i> , 2016 , 6, 154-65	24.4	269
115	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
114	GWAS meta-analysis of 16 852 women identifies new susceptibility locus for endometrial cancer. <i>Human Molecular Genetics</i> , 2016 , 25, 2612-2620	5.6	15
113	Pathway, in silico and tissue-specific expression quantitative analyses of oesophageal squamous cell carcinoma genome-wide association studies data. <i>International Journal of Epidemiology</i> , 2016 , 45, 206-20	7.8	10
112	Prostate Cancer Susceptibility in Men of African Ancestry at 8q24. <i>Journal of the National Cancer Institute</i> , 2016 , 108,	9.7	72
111	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, 1203-14	5.6	20
110	Three new pancreatic cancer susceptibility signals identified on chromosomes 1q32.1, 5p15.33 and 8q24.21. <i>Oncotarget</i> , 2016 , 7, 66328-66343	3.3	66
109	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
108	Cross-Cancer Genome-Wide Analysis of Lung, Ovary, Breast, Prostate, and Colorectal Cancer Reveals Novel Pleiotropic Associations. <i>Cancer Research</i> , 2016 , 76, 5103-14	10.1	66
107	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> , 2016 , 25, 4726-4738	5.6	17
106	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> , 2015 , 134, 333-41	6.3	24
105	Directional dominance on stature and cognition in diverse human populations. <i>Nature</i> , 2015 , 523, 459-462	20.4	119
104	Integration of multiethnic fine-mapping and genomic annotation to prioritize candidate functional SNPs at prostate cancer susceptibility regions. <i>Human Molecular Genetics</i> , 2015 , 24, 5603-18	5.6	35

103	A Genome-Wide Scan Identifies Variants in NFIB Associated with Metastasis in Patients with Osteosarcoma. <i>Cancer Discovery</i> , 2015 , 5, 920-31	24.4	71
102	Two susceptibility loci identified for prostate cancer aggressiveness. <i>Nature Communications</i> , 2015 , 6, 6889	17.4	75
101	Common genetic variants in epigenetic machinery genes and risk of upper gastrointestinal cancers. <i>International Journal of Epidemiology</i> , 2015 , 44, 1341-52	7.8	7
100	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> , 2015 , 36, 684-8	4.7	12
99	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> , 2015 , 137, 311-9	7.5	55
98	TERT gene harbors multiple variants associated with pancreatic cancer susceptibility. <i>International Journal of Cancer</i> , 2015 , 137, 2175-83	7.5	46
97	Analysis of Heritability and Shared Heritability Based on Genome-Wide Association Studies for Thirteen Cancer Types. <i>Journal of the National Cancer Institute</i> , 2015 , 107, djv279	9.7	107
96	Multilevel-analysis identify a cis-expression quantitative trait locus associated with risk of renal cell carcinoma. <i>Oncotarget</i> , 2015 , 6, 4097-109	3.3	1
95	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> , 2015 , 11, e1005378	6	220
94	Methodological Considerations in Estimation of Phenotype Heritability Using Genome-Wide SNP Data, Illustrated by an Analysis of the Heritability of Height in a Large Sample of African Ancestry Adults. <i>PLoS ONE</i> , 2015 , 10, e0131106	3.7	2
93	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
92	Characterization of large structural genetic mosaicism in human autosomes. <i>American Journal of Human Genetics</i> , 2015 , 96, 487-97	11	77
91	Noninvasive evaluation of renal oxygenation in primary nephrotic syndrome with blood oxygen level dependent magnetic resonance imaging: Initial experience. <i>Journal of International Medical Research</i> , 2015 , 43, 356-63	1.4	4
90	A large multiethnic genome-wide association study of prostate cancer identifies novel risk variants and substantial ethnic differences. <i>Cancer Discovery</i> , 2015 , 5, 878-91	24.4	94
89	Distinct Genomic Alterations in Prostate Tumors from African American Men. <i>EBioMedicine</i> , 2015 , 2, 1858-8		1
88	Genetic studies of body mass index yield new insights for obesity biology. <i>Nature</i> , 2015 , 518, 197-206	50.4	2687
87	Common variation at 1q24.1 (ALDH9A1) is a potential risk factor for renal cancer. <i>PLoS ONE</i> , 2015 , 10, e0122589	3.7	15
86	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> , 2015 , 8, 7482-7	1.4	12

85	Genome-wide interaction study of smoking and bladder cancer risk. <i>Carcinogenesis</i> , 2014 , 35, 1737-44	4.6	33
84	Genome-wide association study identifies multiple loci associated with bladder cancer risk. <i>Human Molecular Genetics</i> , 2014 , 23, 1387-98	5.6	101
83	A genome-wide association study of prostate cancer in West African men. <i>Human Genetics</i> , 2014 , 133, 509-21	6.3	56
82	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> , 2014 , 23, 47-54	4	28
81	Rare missense variants in POT1 predispose to familial cutaneous malignant melanoma. <i>Nature Genetics</i> , 2014 , 46, 482-6	36.3	220
80	A genome-wide association study of renal cell carcinoma among African Americans. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014 , 23, 209-14	4	17
79	Genetic variants in fas signaling pathway genes and risk of gastric cancer. <i>International Journal of Cancer</i> , 2014 , 134, 822-31	7.5	21
78	The 19q12 bladder cancer GWAS signal: association with cyclin E function and aggressive disease. <i>Cancer Research</i> , 2014 , 74, 5808-18	10.1	19
77	Genetic polymorphisms in the 9p21 region associated with risk of multiple cancers. <i>Carcinogenesis</i> , 2014 , 35, 2698-705	4.6	54
76	Joint analysis of three genome-wide association studies of esophageal squamous cell carcinoma in Chinese populations. <i>Nature Genetics</i> , 2014 , 46, 1001-1006	36.3	112
75	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
74	Characterizing the genetic basis of methylome diversity in histologically normal human lung tissue. <i>Nature Communications</i> , 2014 , 5, 3365	17.4	103
73	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
72	Genome-wide association study identifies multiple susceptibility loci for pancreatic cancer. <i>Nature Genetics</i> , 2014 , 46, 994-1000	36.3	226
71	A meta-analysis of 87,040 individuals identifies 23 new susceptibility loci for prostate cancer. <i>Nature Genetics</i> , 2014 , 46, 1103-9	36.3	331
70	Defining the role of common variation in the genomic and biological architecture of adult human height. <i>Nature Genetics</i> , 2014 , 46, 1173-86	36.3	1339
69	Pathway-based analysis of GWAs data identifies association of sex determination genes with susceptibility to testicular germ cell tumors. <i>Human Molecular Genetics</i> , 2014 , 23, 6061-8	5.6	25
68	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

67	Gastric cancer following a liver transplantation for glycogen storage disease type Ia (von Gierke disease): A case report. <i>Oncology Letters</i> , 2014 , 8, 2803-2805	2.6	5
66	A fast and powerful tree-based association test for detecting complex joint effects in case-control studies. <i>Bioinformatics</i> , 2014 , 30, 2171-8	7.2	3
65	DNA methylation levels at chromosome 8q24 in peripheral blood are associated with 8q24 cancer susceptibility loci. <i>Cancer Prevention Research</i> , 2014 , 7, 1282-92	3.2	9
64	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
63	HHV-6-associated acute lymphadenitis in immunocompetent patients: a case report and review of literature. <i>International Journal of Clinical and Experimental Pathology</i> , 2014 , 7, 3413-7	1.4	7
62	Meta-analysis of gene-level associations for rare variants based on single-variant statistics. <i>American Journal of Human Genetics</i> , 2013 , 93, 236-48	11	49
61	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
60	Association between MDR1 C3435T polymorphism and risk of breast cancer. <i>Gene</i> , 2013 , 532, 94-9	3.8	16
59	Common genetic polymorphisms modify the effect of smoking on absolute risk of bladder cancer. <i>Cancer Research</i> , 2013 , 73, 2211-20	10.1	82
58	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> , 2013 , 132, 2464-8	7.5	20
57	An integrated transcriptome and epigenome analysis identifies a novel candidate gene for pancreatic cancer. <i>BMC Medical Genomics</i> , 2013 , 6, 33	3.7	28
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