

Bin Tean Teh

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.

284
papers

22,795
citations

77
h-index

144
g-index

306
ext. papers

26,209
ext. citations

10.1
avg, IF

5.82
L-index

#	Paper	IF	Citations
284	Patterns of somatic mutation in human cancer genomes. <i>Nature</i> , 2007 , 446, 153-8	50.4	2400
283	Exome sequencing identifies frequent mutation of the SWI/SNF complex gene PBRM1 in renal carcinoma. <i>Nature</i> , 2011 , 469, 539-42	50.4	943
282	Systematic sequencing of renal carcinoma reveals inactivation of histone modifying genes. <i>Nature</i> , 2010 , 463, 360-3	50.4	927
281	Somatic mutations of the histone H3K27 demethylase gene UTX in human cancer. <i>Nature Genetics</i> , 2009 , 41, 521-3	36.3	627
280	HRPT2, encoding parafibromin, is mutated in hyperparathyroidism-jaw tumor syndrome. <i>Nature Genetics</i> , 2002 , 32, 676-80	36.3	575
279	Signatures of mutation and selection in the cancer genome. <i>Nature</i> , 2010 , 463, 893-8	50.4	538
278	Exome sequencing of gastric adenocarcinoma identifies recurrent somatic mutations in cell adhesion and chromatin remodeling genes. <i>Nature Genetics</i> , 2012 , 44, 570-4	36.3	486
277	Somatic mutations of the protein kinase gene family in human lung cancer. <i>Cancer Research</i> , 2005 , 65, 7591-5	10.1	392
276	Whole-Genome and Epigenomic Landscapes of Etiologically Distinct Subtypes of Cholangiocarcinoma. <i>Cancer Discovery</i> , 2017 , 7, 1116-1135	24.4	368
275	A SUMOylation-defective MITF germline mutation predisposes to melanoma and renal carcinoma. <i>Nature</i> , 2011 , 480, 94-8	50.4	365
274	Exome sequencing of liver fluke-associated cholangiocarcinoma. <i>Nature Genetics</i> , 2012 , 44, 690-3	36.3	348
273	Interleukin-8 mediates resistance to antiangiogenic agent sunitinib in renal cell carcinoma. <i>Cancer Research</i> , 2010 , 70, 1063-71	10.1	342
272	Exome sequencing identifies distinct mutational patterns in liver fluke-related and non-infection-related bile duct cancers. <i>Nature Genetics</i> , 2013 , 45, 1474-8	36.3	334
271	Gene expression signatures delineate biological and prognostic subgroups in peripheral T-cell lymphoma. <i>Blood</i> , 2014 , 123, 2915-23	2.2	299
270	An antioxidant response phenotype shared between hereditary and sporadic type 2 papillary renal cell carcinoma. <i>Cancer Cell</i> , 2011 , 20, 511-23	24.3	299
269	Identification of molecular subtypes of gastric cancer with different responses to PI3-kinase inhibitors and 5-fluorouracil. <i>Gastroenterology</i> , 2013 , 145, 554-65	13.3	288
268	Origins and functional consequences of somatic mitochondrial DNA mutations in human cancer. <i>ELife</i> , 2014 , 3,	8.9	229

267	Preparing the "soil": the primary tumor induces vasculature reorganization in the sentinel lymph node before the arrival of metastatic cancer cells. <i>Cancer Research</i> , 2006 , 66, 10365-76	10.1	227
266	VHL promotes E2 box-dependent E-cadherin transcription by HIF-mediated regulation of SIP1 and snail. <i>Molecular and Cellular Biology</i> , 2007 , 27, 157-69	4.8	217
265	Renal translocation carcinomas: clinicopathologic, immunohistochemical, and gene expression profiling analysis of 31 cases with a review of the literature. <i>American Journal of Surgical Pathology</i> , 2008 , 32, 656-70	6.7	196
264	Genome-wide mutational signatures of aristolochic acid and its application as a screening tool. <i>Science Translational Medicine</i> , 2013 , 5, 197ra101	17.5	194
263	Janus kinase 3-activating mutations identified in natural killer/T-cell lymphoma. <i>Cancer Discovery</i> , 2012 , 2, 591-7	24.4	193
262	Loss of parafibromin immunoreactivity is a distinguishing feature of parathyroid carcinoma. <i>Clinical Cancer Research</i> , 2004 , 10, 6629-37	12.9	193
261	Molecular classification of patients with unexplained hamartomatous and hyperplastic polyposis. <i>JAMA - Journal of the American Medical Association</i> , 2005 , 294, 2465-73	27.4	189
260	A molecular classification of papillary renal cell carcinoma. <i>Cancer Research</i> , 2005 , 65, 5628-37	10.1	185
259	Aristolochic acids and their derivatives are widely implicated in liver cancers in Taiwan and throughout Asia. <i>Science Translational Medicine</i> , 2017 , 9,	17.5	184
258	Sunitinib acts primarily on tumor endothelium rather than tumor cells to inhibit the growth of renal cell carcinoma. <i>Cancer Research</i> , 2010 , 70, 1053-62	10.1	182
257	Molecular subclassification of kidney tumors and the discovery of new diagnostic markers. <i>Oncogene</i> , 2003 , 22, 6810-8	9.2	176
256	Inhibition of MAPK kinase signaling pathways suppressed renal cell carcinoma growth and angiogenesis in vivo. <i>Cancer Research</i> , 2008 , 68, 81-8	10.1	165
255	Birt-Hogg-Dubé syndrome: mapping of a novel hereditary neoplasia gene to chromosome 17p12-q11.2. <i>Oncogene</i> , 2001 , 20, 5239-42	9.2	162
254	Regulation of endocytosis via the oxygen-sensing pathway. <i>Nature Medicine</i> , 2009 , 15, 319-24	50.5	158
253	Glypican 3: a novel marker in testicular germ cell tumors. <i>American Journal of Surgical Pathology</i> , 2006 , 30, 1570-5	6.7	150
252	Expression of alpha-methylacyl-CoA racemase in papillary renal cell carcinoma. <i>American Journal of Surgical Pathology</i> , 2004 , 28, 69-76	6.7	150
251	Thymic carcinoids in multiple endocrine neoplasia type 1. <i>Annals of Surgery</i> , 1998 , 228, 99-105	7.8	143
250	Reversible epithelial to mesenchymal transition and acquired resistance to sunitinib in patients with renal cell carcinoma: evidence from a xenograft study. <i>Molecular Cancer Therapeutics</i> , 2010 , 9, 1525-35	6.1	140

249	Exome sequencing identifies highly recurrent MED12 somatic mutations in breast fibroadenoma. <i>Nature Genetics</i> , 2014 , 46, 877-80	36.3	138
248	Methylation subtypes and large-scale epigenetic alterations in gastric cancer. <i>Science Translational Medicine</i> , 2012 , 4, 156ra140	17.5	134
247	Genomic landscapes of breast fibroepithelial tumors. <i>Nature Genetics</i> , 2015 , 47, 1341-5	36.3	129
246	Tubulocystic carcinoma of the kidney: clinicopathologic and molecular characterization. <i>American Journal of Surgical Pathology</i> , 2008 , 32, 177-87	6.7	127
245	Detection of DNA copy number changes and oncogenic signaling abnormalities from gene expression data reveals MYC activation in high-grade papillary renal cell carcinoma. <i>Cancer Research</i> , 2007 , 67, 3171-6	10.1	124
244	Excessive fatty acid oxidation induces muscle atrophy in cancer cachexia. <i>Nature Medicine</i> , 2016 , 22, 666-71	5.5	123
243	Characterization of the mouse Men1 gene and its expression during development. <i>Oncogene</i> , 1998 , 17, 2485-93	9.2	122
242	MicroRNA profiling of human kidney cancer subtypes. <i>International Journal of Oncology</i> , 2009 , 35, 109-14		121
241	Loss of tumor suppressor KDM6A amplifies PRC2-regulated transcriptional repression in bladder cancer and can be targeted through inhibition of EZH2. <i>Science Translational Medicine</i> , 2017 , 9,	17.5	114
240	Genomic and Epigenomic Profiling of High-Risk Intestinal Metaplasia Reveals Molecular Determinants of Progression to Gastric Cancer. <i>Cancer Cell</i> , 2018 , 33, 137-150.e5	24.3	113
239	The <i>Opisthorchis viverrini</i> genome provides insights into life in the bile duct. <i>Nature Communications</i> , 2014 , 5, 4378	17.4	113
238	Serglycin is a theranostic target in nasopharyngeal carcinoma that promotes metastasis. <i>Cancer Research</i> , 2011 , 71, 3162-72	10.1	113
237	Deficiency of FLCN in mouse kidney led to development of polycystic kidneys and renal neoplasia. <i>PLoS ONE</i> , 2008 , 3, e3581	3.7	113
236	Functional importance of Dicer protein in the adaptive cellular response to hypoxia. <i>Journal of Biological Chemistry</i> , 2012 , 287, 29003-20	5.4	111
235	Cystic renal neoplasia following conditional inactivation of apc in mouse renal tubular epithelium. <i>Journal of Biological Chemistry</i> , 2005 , 280, 3938-45	5.4	111
234	Whole-genome reconstruction and mutational signatures in gastric cancer. <i>Genome Biology</i> , 2012 , 13, R115	18.3	110
233	As an independent unfavorable prognostic factor, IL-8 promotes metastasis of nasopharyngeal carcinoma through induction of epithelial-mesenchymal transition and activation of AKT signaling. <i>Carcinogenesis</i> , 2012 , 33, 1302-9	4.6	108
232	Accuracy of combined protein gene product 9.5 and parafibromin markers for immunohistochemical diagnosis of parathyroid carcinoma. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2009 , 94, 434-41	5.6	105

231	Functional characterization of JMJD2A, a histone deacetylase- and retinoblastoma-binding protein. <i>Journal of Biological Chemistry</i> , 2005 , 280, 28507-18	5.4	105
230	Hypoxia promotes ligand-independent EGF receptor signaling via hypoxia-inducible factor-mediated upregulation of caveolin-1. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012 , 109, 4892-7	11.5	101
229	Prospective clinical trial of preoperative sunitinib in patients with renal cell carcinoma. <i>Journal of Urology</i> , 2010 , 184, 859-64	2.5	97
228	The draft genome of tropical fruit durian (<i>Durio zibethinus</i>). <i>Nature Genetics</i> , 2017 , 49, 1633-1641	36.3	96
227	Clinicopathologic studies of thymic carcinoids in multiple endocrine neoplasia type 1. <i>Medicine (United States)</i> , 1997 , 76, 21-9	1.8	96
226	Novel FH mutations in families with hereditary leiomyomatosis and renal cell cancer (HLRCC) and patients with isolated type 2 papillary renal cell carcinoma. <i>Journal of Medical Genetics</i> , 2011 , 48, 226-34	5.8	94
225	Comparative genomic hybridization reveals frequent losses of chromosomes 1p and 3q in pheochromocytomas and abdominal paragangliomas, suggesting a common genetic etiology. <i>American Journal of Pathology</i> , 2000 , 156, 651-9	5.8	94
224	Robust classification of renal cell carcinoma based on gene expression data and predicted cytogenetic profiles. <i>Cancer Research</i> , 2004 , 64, 4117-21	10.1	90
223	ADAR-Mediated RNA Editing Predicts Progression and Prognosis of Gastric Cancer. <i>Gastroenterology</i> , 2016 , 151, 637-650.e10	13.3	90
222	Genomic and Transcriptomic Profiling of Combined Hepatocellular and Intrahepatic Cholangiocarcinoma Reveals Distinct Molecular Subtypes. <i>Cancer Cell</i> , 2019 , 35, 932-947.e8	24.3	89
221	Next-generation sequencing of translocation renal cell carcinoma reveals novel RNA splicing partners and frequent mutations of chromatin-remodeling genes. <i>Clinical Cancer Research</i> , 2014 , 20, 4129-40	12.9	89
220	Gene expression profiling of renal medullary carcinoma: potential clinical relevance. <i>Cancer</i> , 2004 , 100, 976-85	6.4	88
219	Genetic drivers of oncogenic pathways in molecular subgroups of peripheral T-cell lymphoma. <i>Blood</i> , 2019 , 133, 1664-1676	2.2	87
218	A genotypic and histopathological study of a large Dutch kindred with hyperparathyroidism-jaw tumor syndrome. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2000 , 85, 1449-54	5.6	86
217	Abscisic acid perception and signaling: structural mechanisms and applications. <i>Acta Pharmacologica Sinica</i> , 2014 , 35, 567-84	8	85
216	Gene expression of parathyroid tumors: molecular subclassification and identification of the potential malignant phenotype. <i>Cancer Research</i> , 2004 , 64, 7405-11	10.1	85
215	The t(1;3) breakpoint-spanning genes LSAMP and NORE1 are involved in clear cell renal cell carcinomas. <i>Cancer Cell</i> , 2003 , 4, 405-13	24.3	85
214	HRPT2 gene alterations in ossifying fibroma of the jaws. <i>Oral Oncology</i> , 2006 , 42, 735-9	4.4	80

213	In-depth characterization of the cisplatin mutational signature in human cell lines and in esophageal and liver tumors. <i>Genome Research</i> , 2018 , 28, 654-665	9.7	79
212	MSIseq: Software for Assessing Microsatellite Instability from Catalogs of Somatic Mutations. <i>Scientific Reports</i> , 2015 , 5, 13321	4.9	79
211	Inactivation of BHD in sporadic renal tumors. <i>Cancer Research</i> , 2003 , 63, 4583-7	10.1	79
210	Epigenomic profiling of primary gastric adenocarcinoma reveals super-enhancer heterogeneity. <i>Nature Communications</i> , 2016 , 7, 12983	17.4	78
209	Parafibromin, a component of the human PAF complex, regulates growth factors and is required for embryonic development and survival in adult mice. <i>Molecular and Cellular Biology</i> , 2008 , 28, 2930-40	4.8	78
208	C-kit expression in renal oncocytomas and chromophobe renal cell carcinomas. <i>Human Pathology</i> , 2005 , 36, 262-8	3.7	78
207	A family with isolated hyperparathyroidism segregating a missense MEN1 mutation and showing loss of the wild-type alleles in the parathyroid tumors. <i>American Journal of Human Genetics</i> , 1998 , 63, 1544-9	11	77
206	Human kidney injury molecule-1 (hKIM-1): a useful immunohistochemical marker for diagnosing renal cell carcinoma and ovarian clear cell carcinoma. <i>American Journal of Surgical Pathology</i> , 2007 , 31, 371-81	6.7	76
205	Genetic and expression profiles of squamous cell carcinoma of the head and neck correlate with cisplatin sensitivity and resistance in cell lines and patients. <i>Clinical Cancer Research</i> , 2004 , 10, 8204-13	12.9	76
204	Integrated epigenomics identifies BMP4 as a modulator of cisplatin sensitivity in gastric cancer. <i>Gut</i> , 2013 , 62, 22-33	19.2	75
203	Two distinct types of blood vessels in clear cell renal cell carcinoma have contrasting prognostic implications. <i>Clinical Cancer Research</i> , 2007 , 13, 161-9	12.9	75
202	Alternative genetic pathways in parathyroid tumorigenesis. <i>Journal of Clinical Endocrinology and Metabolism</i> , 1999 , 84, 3775-80	5.6	75
201	Genomic expression and single-nucleotide polymorphism profiling discriminates chromophobe renal cell carcinoma and oncocytoma. <i>BMC Cancer</i> , 2010 , 10, 196	4.8	74
200	VHL substrate transcription factor ZHX2 as an oncogenic driver in clear cell renal cell carcinoma. <i>Science</i> , 2018 , 361, 290-295	33.3	73
199	Complexity of tumor vasculature in clear cell renal cell carcinoma. <i>Cancer</i> , 2009 , 115, 2282-9	6.4	73
198	The investigational Aurora kinase A inhibitor MLN8237 induces defects in cell viability and cell-cycle progression in malignant bladder cancer cells in vitro and in vivo. <i>Clinical Cancer Research</i> , 2013 , 19, 1717-28	12.9	72
197	Chromosomal amplification of leucine-rich repeat kinase-2 (LRRK2) is required for oncogenic MET signaling in papillary renal and thyroid carcinomas. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011 , 108, 1439-44	11.5	72
196	Adult cystic nephroma and mixed epithelial and stromal tumor of the kidney are the same disease entity: molecular and histologic evidence. <i>American Journal of Surgical Pathology</i> , 2009 , 33, 72-80	6.7	70

195	Evaluation of retinoblastoma and Ki-67 immunostaining as diagnostic markers of benign and malignant parathyroid disease. <i>World Journal of Surgery</i> , 1999 , 23, 68-74	3.3	70
194	Deficiency Drives Enhancer Activation of Oncogenes in Clear Cell Renal Cell Carcinoma. <i>Cancer Discovery</i> , 2017 , 7, 1284-1305	24.4	69
193	Genetic and structural variation in the gastric cancer kinome revealed through targeted deep sequencing. <i>Cancer Research</i> , 2011 , 71, 29-39	10.1	69
192	Parafibromin inhibits cancer cell growth and causes G1 phase arrest. <i>Biochemical and Biophysical Research Communications</i> , 2006 , 350, 17-24	3.4	69
191	Whole-exome sequencing studies of parathyroid carcinomas reveal novel PRUNE2 mutations, distinctive mutational spectra related to APOBEC-catalyzed DNA mutagenesis and mutational enrichment in kinases associated with cell migration and invasion. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2015 , 100, E360-4	5.6	68
190	Phenotype and phenocopy: the relationship between genotype and clinical phenotype in a single large family with multiple endocrine neoplasia type 1 (MEN 1). <i>Clinical Endocrinology</i> , 2000 , 53, 205-11	3.4	68
189	Differential loss of heterozygosity in familial, sporadic, and uremic hyperparathyroidism. <i>Human Genetics</i> , 1997 , 99, 342-9	6.3	67
188	Genome-scale mutational signatures of aflatoxin in cells, mice, and human tumors. <i>Genome Research</i> , 2017 , 27, 1475-1486	9.7	64
187	Mutational landscapes of tongue carcinoma reveal recurrent mutations in genes of therapeutic and prognostic relevance. <i>Genome Medicine</i> , 2015 , 7, 98	14.4	64
186	Classification of renal neoplasms based on molecular signatures. <i>Journal of Urology</i> , 2006 , 175, 2302-6	2.5	64
185	Mutation signatures implicate aristolochic acid in bladder cancer development. <i>Genome Medicine</i> , 2015 , 7, 38	14.4	63
184	Birt-Hogg-Dubé renal tumors are genetically distinct from other renal neoplasias and are associated with up-regulation of mitochondrial gene expression. <i>BMC Medical Genomics</i> , 2010 , 3, 59	3.7	63
183	Facilitating a culture of responsible and effective sharing of cancer genome data. <i>Nature Medicine</i> , 2016 , 22, 464-71	50.5	63
182	Tissue Microbiome Profiling Identifies an Enrichment of Specific Enteric Bacteria in Opisthorchis viverrini Associated Cholangiocarcinoma. <i>EBioMedicine</i> , 2016 , 8, 195-202	8.8	61
181	Genetic risk of extranodal natural killer T-cell lymphoma: a genome-wide association study. <i>Lancet Oncology</i> , 2016 , 17, 1240-7	21.7	58
180	Nanoscale chromatin profiling of gastric adenocarcinoma reveals cancer-associated cryptic promoters and somatically acquired regulatory elements. <i>Nature Communications</i> , 2014 , 5, 4361	17.4	55
179	Familial isolated hyperparathyroidism as a variant of multiple endocrine neoplasia type 1 in a large Danish pedigree. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2000 , 85, 165-7	5.6	55
178	Activation of the PI3K/AKT pathway induces urothelial carcinoma of the renal pelvis: identification in human tumors and confirmation in animal models. <i>Cancer Research</i> , 2009 , 69, 8256-64	10.1	54

177	Gene expression profiling of favorable histology Wilms tumors and its correlation with clinical features. <i>Cancer Research</i> , 2002 , 62, 6598-605	10.1	54
176	Renal papillary adenoma--a putative precursor of papillary renal cell carcinoma. <i>Human Pathology</i> , 2007 , 38, 239-46	3.7	53
175	Downregulation of CASR expression and global loss of parafibromin staining are strong negative determinants of prognosis in parathyroid carcinoma. <i>Modern Pathology</i> , 2011 , 24, 688-97	9.8	52
174	Familial non-VHL non-papillary clear-cell renal cancer. <i>Lancet, The</i> , 1997 , 349, 848-9	4.0	52
173	Somatic pairing of chromosome 19 in renal oncocytoma is associated with deregulated EGLN2-mediated [corrected] oxygen-sensing response. <i>PLoS Genetics</i> , 2008 , 4, e1000176	6	52
172	Mapping of the second locus for the Van der Woude syndrome to chromosome 1p34. <i>European Journal of Human Genetics</i> , 2001 , 9, 747-52	5.3	52
171	Prognostic value of tumor necrosis factor-related apoptosis-inducing ligand (TRAIL) and TRAIL receptors in renal cell cancer. <i>Clinical Cancer Research</i> , 2009 , 15, 650-9	12.9	51
170	Mutation signatures of carcinogen exposure: genome-wide detection and new opportunities for cancer prevention. <i>Genome Medicine</i> , 2014 , 6, 24	14.4	50
169	MED12 is frequently mutated in breast phyllodes tumours: a study of 112 cases. <i>Journal of Clinical Pathology</i> , 2015 , 68, 685-91	3.9	50
168	Genomic profiling reveals mutational landscape in parathyroid carcinomas. <i>JCI Insight</i> , 2017 , 2, e92061	9.9	50
167	A large multiple endocrine neoplasia type 1 family with clinical expression suggestive of anticipation. <i>Journal of Clinical Endocrinology and Metabolism</i> , 1997 , 82, 3487-92	5.6	49
166	Expression of S-100 protein in renal cell neoplasms. <i>Human Pathology</i> , 2006 , 37, 462-70	3.7	48
165	The phospholipase C beta 3 gene located in the MEN1 region shows loss of expression in endocrine tumours. <i>Human Molecular Genetics</i> , 1994 , 3, 1775-81	5.6	48
164	Radiosensitization by inhibiting STAT1 in renal cell carcinoma. <i>International Journal of Radiation Oncology Biology Physics</i> , 2009 , 73, 288-95	4	47
163	Familial hyperparathyroidism: surgical outcome after 30 years of follow-up in three families with germline HRPT2 mutations. <i>Surgery</i> , 2008 , 143, 630-40	3.6	47
162	Exome-wide Sequencing Shows Low Mutation Rates and Identifies Novel Mutated Genes in Seminomas. <i>European Urology</i> , 2015 , 68, 77-83	10.2	46
161	Expression of the PTTG1 oncogene is associated with aggressive clear cell renal cell carcinoma. <i>Cancer Research</i> , 2012 , 72, 4361-71	10.1	46
160	An eleven gene molecular signature for extra-capsular spread in oral squamous cell carcinoma serves as a prognosticator of outcome in patients without nodal metastases. <i>Oral Oncology</i> , 2015 , 51, 355-62	4.4	45

159	Gene expression profiling of early- and late-relapse nonseminomatous germ cell tumor and primitive neuroectodermal tumor of the testis. <i>Clinical Cancer Research</i> , 2004 , 10, 2368-78	12.9	45
158	Methylation profiles reveal distinct subgroup of hepatocellular carcinoma patients with poor prognosis. <i>PLoS ONE</i> , 2014 , 9, e104158	3.7	44
157	Genetic studies of a family with hereditary hyperparathyroidism-jaw tumour syndrome. <i>Clinical Endocrinology</i> , 1999 , 50, 191-6	3.4	44
156	Comparative gene expression profiling analysis of urothelial carcinoma of the renal pelvis and bladder. <i>BMC Medical Genomics</i> , 2010 , 3, 58	3.7	43
155	Kinase targets in renal-cell carcinomas: reassessing the old and discovering the new. <i>Lancet Oncology</i> , 2010 , 11, 571-8	21.7	42
154	p38delta/MAPK13 as a diagnostic marker for cholangiocarcinoma and its involvement in cell motility and invasion. <i>International Journal of Cancer</i> , 2010 , 126, 2353-61	7.5	42
153	Folliculin interacts with p0071 (plakophilin-4) and deficiency is associated with disordered RhoA signalling, epithelial polarization and cytokinesis. <i>Human Molecular Genetics</i> , 2012 , 21, 5268-79	5.6	42
152	Beyond fitness tracking: The use of consumer-grade wearable data from normal volunteers in cardiovascular and lipidomics research. <i>PLoS Biology</i> , 2018 , 16, e2004285	9.7	42
151	Selective loss of chromosome 11 in pheochromocytomas associated with the VHL syndrome. <i>Oncogene</i> , 2002 , 21, 1117-22	9.2	41
150	Pericyte coverage of differentiated vessels inside tumor vasculature is an independent unfavorable prognostic factor for patients with clear cell renal cell carcinoma. <i>Cancer</i> , 2013 , 119, 313-24	6.4	40
149	Renal medullary carcinoma: molecular, pathological and clinical evidence for treatment with topoisomerase-inhibiting therapy. <i>BJU International</i> , 2010 , 106, 62-5	5.6	40
148	Lymphoblastoid cell line with B1 cell characteristics established from a chronic lymphocytic leukemia clone by in vitro EBV infection. <i>Oncolmmunology</i> , 2012 , 1, 18-27	7.2	40
147	Malignant melanoma in patients with multiple endocrine neoplasia type 1 and involvement of the MEN1 gene in sporadic melanoma. <i>International Journal of Cancer</i> , 2000 , 87, 463-7	7.5	40
146	Expression of RON Proto-oncogene in Renal Oncocytoma and Chromophobe Renal Cell Carcinoma. <i>American Journal of Surgical Pathology</i> , 2004 , 28, 1045-50	6.7	39
145	High-depth sequencing of over 750 genes supports linear progression of primary tumors and metastases in most patients with liver-limited metastatic colorectal cancer. <i>Genome Biology</i> , 2015 , 16, 32	18.3	38
144	The Succinated Proteome of FH-Mutant Tumours. <i>Metabolites</i> , 2014 , 4, 640-54	5.6	38
143	Identification of a new exon and complex splicing alterations in familial erythrocytosis or von Hippel-Lindau disease. <i>Blood</i> , 2018 , 132, 469-483	2.2	37
142	Germline Mutations in Cancer Predisposition Genes are Frequent in Sporadic Sarcomas. <i>Scientific Reports</i> , 2017 , 7, 10660	4.9	37

141	Dual modulation of MCL-1 and mTOR determines the response to sunitinib. <i>Journal of Clinical Investigation</i> , 2017 , 127, 153-168	15.9	37
140	Functional and genetic characterization of three cell lines derived from a single tumor of an <i>Opisthorchis viverrini</i> -associated cholangiocarcinoma patient. <i>Human Cell</i> , 2020 , 33, 695-708	4.5	36
139	All the World's a Stage: Facilitating Discovery Science and Improved Cancer Care through the Global Alliance for Genomics and Health. <i>Cancer Discovery</i> , 2015 , 5, 1133-6	24.4	35
138	Genetics of <i>Opisthorchis viverrini</i> -related cholangiocarcinoma. <i>Current Opinion in Gastroenterology</i> , 2015 , 31, 258-63	3	35
137	Innovations and challenges in renal cancer: consensus statement from the first international conference. <i>Clinical Cancer Research</i> , 2004 , 10, 6277S-81S	12.9	35
136	VX680/MK-0457, a potent and selective Aurora kinase inhibitor, targets both tumor and endothelial cells in clear cell renal cell carcinoma. <i>American Journal of Translational Research (discontinued)</i> , 2010 , 2, 296-308	3	35
135	Epigenomic Promoter Alterations Amplify Gene Isoform and Immunogenic Diversity in Gastric Adenocarcinoma. <i>Cancer Discovery</i> , 2017 , 7, 630-651	24.4	34
134	Molecular classification of breast phyllodes tumors: validation of the histologic grading scheme and insights into malignant progression. <i>Breast Cancer Research and Treatment</i> , 2011 , 129, 319-29	4.4	34
133	Gene expression profiling of renal cell carcinoma. <i>Clinical Cancer Research</i> , 2004 , 10, 6315S-21S	12.9	32
132	SETD2 histone modifier loss in aggressive GI stromal tumours. <i>Gut</i> , 2016 , 65, 1960-1972	19.2	30
131	Cholangiocarcinoma. <i>Nature Reviews Disease Primers</i> , 2021 , 7, 65	51.1	30
130	Pathogenesis of cholangiocarcinoma: From genetics to signalling pathways. <i>Baillieres Best Practice and Research in Clinical Gastroenterology</i> , 2015 , 29, 233-44	2.5	29
129	Deregulation of E2-EPF ubiquitin carrier protein in papillary renal cell carcinoma. <i>American Journal of Pathology</i> , 2011 , 178, 853-60	5.8	29
128	CDC73/HRPT2 CpG island hypermethylation and mutation of 5' untranslated sequence are uncommon mechanisms of silencing parafibromin in parathyroid tumors. <i>Endocrine-Related Cancer</i> , 2010 , 17, 273-82	5.7	29
127	Single base mutation in the hormone binding domain of the thyroid hormone receptor beta gene in generalised thyroid hormone resistance demonstrated by single stranded conformation polymorphism analysis. <i>Biochemical and Biophysical Research Communications</i> , 1991 , 178, 606-12	3.4	29
126	The tumor suppressor parafibromin is required for posttranscriptional processing of histone mRNA. <i>Molecular Carcinogenesis</i> , 2010 , 49, 215-23	5	28
125	Gene expression profiling of renal cell carcinoma and its implications in diagnosis, prognosis, and therapeutics. <i>Advances in Cancer Research</i> , 2003 , 89, 157-81	5.9	28
124	Independent genetic events associated with the development of multiple parathyroid tumors in patients with primary hyperparathyroidism. <i>American Journal of Pathology</i> , 2002 , 161, 1299-306	5.8	28

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