John K Cowell

List of Publications by Year in descending order

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214 papers 9,979 citations

51 h-index 89 g-index

219 all docs

219 docs citations

219 times ranked

11467 citing authors

#	Article	IF	CITATIONS
1	Investigation of LGI1 as the antigen in limbic encephalitis previously attributed to potassium channels: a case series. Lancet Neurology, The, 2010, 9, 776-785.	4.9	947
2	Insulin-like growth factor-II gene expression in Wilms' tumour and embryonic tissues. Nature, 1985, 317, 260-262.	13.7	419
3	Monocytic and granulocytic myeloid derived suppressor cells differentially regulate spatiotemporal tumour plasticity during metastatic cascade. Nature Communications, 2017, 8, 14979.	5.8	292
4	The TACC domain identifies a family of centrosomal proteins that can interact with microtubules. Proceedings of the National Academy of Sciences of the United States of America, 2000, 97, 14352-14357.	3.3	250
5	Molecular Study of Malignant Gliomas Treated with Epidermal Growth Factor Receptor Inhibitors: Tissue Analysis from North American Brain Tumor Consortium Trials 01-03 and 00-01. Clinical Cancer Research, 2005, 11, 7841-7850.	3.2	238
6	Targeted therapy of human malignant glioma in a mouse model by 2-5A antisense directed against telomerase RNA. Oncogene, 1998, 16, 3323-3330.	2.6	194
7	Expression of oligodendrocyte progenitor cell antigens by gliomas: Implications for the histogenesis of brain tumors. Proceedings of the National Academy of Sciences of the United States of America, 1999, 96, 10361-10366.	3.3	175
8	A novel gene, LGI1, from 10q24 is rearranged and downregulated in malignant brain tumors. Oncogene, 1998, 17, 2873-2881.	2.6	169
9	Antisense telomerase treatment: induction of two distinct pathways, apoptosis and differentiation. FASEB Journal, 1998, 12, 801-811.	0.2	166
10	Inhibition of telomerase increases the susceptibility of human malignant glioblastoma cells to cisplatin-induced apoptosis. Oncogene, 1998, 16, 2243-2248.	2.6	159
11	Manipulation of nonsense mediated decay identifies gene mutations in colon cancer Cells with microsatellite instability. Oncogene, 2004, 23, 639-645.	2.6	154
12	Evaluating human cancer cell metastasis in zebrafish. BMC Cancer, 2013, 13, 453.	1.1	151
13	Oncogenic point mutations in exon 20 of the RB1 gene in families showing incomplete penetrance and mild expression of the retinoblastoma phenotype Proceedings of the National Academy of Sciences of the United States of America, 1992, 89, 6177-6181.	3.3	122
14	The Third Member of the Transforming Acidic Coiled Coil-Containing Gene Family, TACC3, Maps in 4p16, Close to Translocation Breakpoints in Multiple Myeloma, and Is Upregulated in Various Cancer Cell Lines. Genomics, 1999, 58, 165-170.	1.3	121
15	Constitutional mutations in the WT1 gene in patients with Denys-Drash syndrome. Human Molecular Genetics, 1992, 1, 301-305.	1.4	118
16	Cloning of TACC1, an embryonically expressed, potentially transforming coiled coil containing gene, from the 8p11 breast cancer amplicon. Oncogene, 1999, 18, 4032-4038.	2.6	111
17	Ubiquitin-conjugating enzyme UBE2C: molecular biology, role in tumorigenesis, and potential as a biomarker. Tumor Biology, 2012, 33, 723-730.	0.8	108
18	Lgi1 null mutant mice exhibit myoclonic seizures and CA1 neuronal hyperexcitability. Human Molecular Genetics, 2010, 19, 1702-1711.	1.4	106

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19	Genome-wide aberrations in pancreatic adenocarcinoma. Cancer Genetics and Cytogenetics, 2005, 161, 36-50.	1.0	104
20	Down-Regulation of WAVE3, a Metastasis Promoter Gene, Inhibits Invasion and Metastasis of Breast Cancer Cells. American Journal of Pathology, 2007, 170, 2112-2121.	1.9	103
21	WAVE3 promotes cell motility and invasion through the regulation of MMP-1, MMP-3, and MMP-9 expression. Experimental Cell Research, 2005, 308, 135-145.	1.2	99
22	A role for p300/CREB binding protein genes in promoting cancer progression in colon cancer cell lines with microsatellite instability. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 1273-1278.	3.3	98
23	The involvement of JAK-STAT3 in cell motility, invasion, and metastasis. Jak-stat, 2014, 3, e28086.	2.2	98
24	A photographic representation of the variability in the G-banded structure of the chromosomes in the mouse karyotype. Chromosoma, 1984, 89, 294-320.	1.0	95
25	WAVE3-mediated Cell Migration and Lamellipodia Formation Are Regulated Downstream of Phosphatidylinositol 3-Kinase. Journal of Biological Chemistry, 2005, 280, 21748-21755.	1.6	94
26	LGI1, a Putative Tumor Metastasis Suppressor Gene, Controls in Vitro Invasiveness and Expression of Matrix Metalloproteinases in Glioma Cells through the ERK1/2 Pathway. Journal of Biological Chemistry, 2004, 279, 23151-23157.	1.6	93
27	Identifying candidate colon cancer tumor suppressor genes using inhibition of nonsense-mediated mRNA decay in colon cancer cells. Oncogene, 2007, 26, 2873-2884.	2.6	90
28	Radiation-Induced Meningioma: A Distinct Molecular Genetic Pattern?. Journal of Neuropathology and Experimental Neurology, 2000, 59, 614-620.	0.9	85
29	Development of Autoimmunity in IL-14α-Transgenic Mice. Journal of Immunology, 2006, 177, 5676-5686.	0.4	84
30	2-5A Antisense Directed against Telomerase RNA Produces Apoptosis in Ovarian Cancer Cells. Gynecologic Oncology, 2000, 76, 183-192.	0.6	83
31	Homozygous deletion of a DNA marker from chromosome 11p13 in sporadic Wilms tumor. Genomics, 1988, 3, 25-31.	1.3	81
32	c-Abl-mediated Phosphorylation of WAVE3 Is Required for Lamellipodia Formation and Cell Migration. Journal of Biological Chemistry, 2007, 282, 26257-26265.	1.6	81
33	Primary tumor-induced immunity eradicates disseminated tumor cells in syngeneic mouse model. Nature Communications, 2019, 10, 1430.	5.8	77
34	The neural progenitor-restricted isoform of the MARK4 gene in 19q13.2 is upregulated in human gliomas and overexpressed in a subset of glioblastoma cell lines. Oncogene, 2003, 22, 2581-2591.	2.6	76
35	Copy Number and Gene Expression Alterations in Radiation-Induced Papillary Thyroid Carcinoma from Chernobyl Pediatric Patients. Thyroid, 2010, 20, 475-487.	2.4	76
36	Ponatinib suppresses the development of myeloid and lymphoid malignancies associated with FGFR1 abnormalities. Leukemia, 2013, 27, 32-40.	3.3	75

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37	HSP90 and HSP70 Proteins Are Essential for Stabilization and Activation of WASF3 Metastasis-promoting Protein. Journal of Biological Chemistry, 2012, 287, 10051-10059.	1.6	74
38	Cytogenetic changes in Wilms' tumors. Cancer Genetics and Cytogenetics, 1988, 34, 223-234.	1.0	68
39	Suppression of the cell proliferation and invasion phenotypes in glioma cells by the LGI1 gene. Oncogene, 2003, 22, 3985-3991.	2.6	63
40	Molecular characterization of the t(3;9) associated with immortalization in the MCF10A cell line. Cancer Genetics and Cytogenetics, 2005, 163, 23-29.	1.0	63
41	CLCA2 tumour suppressor gene in 1p31 is epigenetically regulated in breast cancer. Oncogene, 2004, 23, 1474-1480.	2.6	61
42	THE EXPRESSION PATTERN OF WILMS' TUMOUR GENE (WT1) PRODUCT IN NORMAL TISSUES AND PAEDIATRIC RENAL TUMOURS. , 1996, 179, 162-168.		60
43	WAVE3, an actin-polymerization gene, is truncated and inactivated as a result of a constitutional $t(1;13)(q21;q12)$ chromosome translocation in a patient with ganglioneuroblastoma. Oncogene, 2002, 21, 5967-5974.	2.6	59
44	Array CGH analysis of pediatric medulloblastomas. Genes Chromosomes and Cancer, 2006, 45, 290-303.	1.5	59
45	Defining the expression pattern of the LGI1 gene in BAC transgenic mice. Mammalian Genome, 2007, 18, 328-337.	1.0	59
46	Application of bacterial artificial chromosome array-based comparative genomic hybridization and spectral karyotyping to the analysis of glioblastoma multiforme. Cancer Genetics and Cytogenetics, 2004, 151, 36-51.	1.0	58
47	Inactivation of the WASF3 gene in prostate cancer cells leads to suppression of tumorigenicity and metastases. British Journal of Cancer, 2010, 103, 1066-1075.	2.9	57
48	The WASF3–NCKAP1–CYFIP1 Complex Is Essential for Breast Cancer Metastasis. Cancer Research, 2016, 76, 5133-5142.	0.4	57
49	NB4S, a Member of the TBC1 Domain Family of Genes, is Truncated as a Result of a Constitutional $t(1;10)(p22;q21)$ Chromosome Translocation in a Patient with Stage 4S Neuroblastoma. Human Molecular Genetics, 1998, 7, 1169-1178.	1.4	56
50	Novel FGFR inhibitor ponatinib suppresses the growth of non-small cell lung cancer cells overexpressing FGFR1. Oncology Reports, 2013, 29, 2181-2190.	1.2	55
51	The Application of Microarray Technology to the Analysis of the Cancer Genome. Current Molecular Medicine, 2007, 7, 103-120.	0.6	53
52	The EVI5 TBC domain provides the GTPase-activating protein motif for RAB11. Oncogene, 2007, 26, 2804-2808.	2.6	53
53	Knockdown of zebrafish Lgi1a results in abnormal development, brain defects and a seizure-like behavioral phenotype. Human Molecular Genetics, 2010, 19, 4409-4420.	1.4	53
54	Functional interrelationship between the WASF3 and KISS1 metastasisâ€associated genes in breast cancer cells. International Journal of Cancer, 2011, 129, 2825-2835.	2.3	52

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55	Colon carcinoma cells harboring PIK3CA mutations display resistance to growth factor deprivation induced apoptosis. Molecular Cancer Therapeutics, 2007, 6, 1143-1150.	1.9	51
56	Comprehensive analysis of loss of heterozygosity events in glioblastoma using the 100K SNP mapping arrays and comparison with copy number abnormalities defined by BAC array comparative genomic hybridization. Genes Chromosomes and Cancer, 2008, 47, 221-237.	1.5	51
57	Characterization of the 1p/19q Chromosomal Loss in Oligodendrogliomas Using Comparative Genomic Hybridization Arrays (CGHa). Journal of Neuropathology and Experimental Neurology, 2004, 63, 151-158.	0.9	49
58	Molecular Characterization of the $t(8;13)(p11;q12)$ Translocation Associated With an Atypical Myeloproliferative Disorder: Evidence for Three Discrete Loci Involved in Myeloid Leukemias on $8p11$. Blood, 1997 , 90 , 3136 - 3141 .	0.6	47
59	A novel member of the WD-repeat gene family, WDR11, maps to the 10q26 region and is disrupted by a chromosome translocation in human glioblastoma cells. Oncogene, 2001, 20, 5378-5392.	2.6	47
60	Genetic analysis of Down syndrome-associated heart defects in mice. Human Genetics, 2011, 130, 623-632.	1.8	47
61	Transgelin increases metastatic potential of colorectal cancer cells in vivo and alters expression of genes involved in cell motility. BMC Cancer, 2016, 16, 55.	1.1	46
62	COP1 and GSK3 \hat{l}^2 Cooperate to Promote c-Jun Degradation and Inhibit Breast Cancer Cell Tumorigenesis. Neoplasia, 2013, 15, 1075-IN11.	2.3	45
63	Targeting the WASF3–CYFIP1 Complex Using Stapled Peptides Suppresses Cancer Cell Invasion. Cancer Research, 2016, 76, 965-973.	0.4	45
64	Genomic organization and expression profile of the human and mouse WAVE gene family. Mammalian Genome, 2003, 14, 314-322.	1.0	44
65	High-Resolution Analysis of Genetic Events in Cancer Cells Using Bacterial Artificial Chromosome Arrays and Comparative Genome Hybridization. Advances in Cancer Research, 2003, 90, 91-125.	1.9	44
66	Loss of heterozygosity for the short arm of chromosome 7 in sporadic Wilms tumour. Oncogene, 1998, 17, 395-400.	2.6	43
67	Interaction of the transforming acidic coiled-coil 1 (TACC1) protein with ch-TOG and GAS41/NuBI1 suggests multiple TACC1-containing protein complexes in human cells. Biochemical Journal, 2002, 363, 195-200.	1.7	43
68	Frequent Constitutional C to T Mutations in CGA-Arginine Codons in the RB1 Gene Produce Premature Stop Codons in Patients with Bilateral (Hereditary) Retinoblastoma. European Journal of Human Genetics, 1994, 2, 281-290.	1.4	43
69	Loss of Zebrafish lgi1b Leads to Hydrocephalus and Sensitization to Pentylenetetrazol Induced Seizure-Like Behavior. PLoS ONE, 2011, 6, e24596.	1.1	43
70	Genetics and cytogenetics of retinoblastoma. Cancer Genetics and Cytogenetics, 1992, 64, 1-11.	1.0	42
71	Arsenic Trioxide Affects Signal Transducer and Activator of Transcription Proteins through Alteration of Protein Tyrosine Kinase Phosphorylation. Clinical Cancer Research, 2006, 12, 6817-6825.	3.2	42
72	Mutation in the FGFR1 tyrosine kinase domain or inactivation of PTEN is associated with acquired resistance to FGFR inhibitors in FGFR1â€driven leukemia/lymphomas. International Journal of Cancer, 2017, 141, 1822-1829.	2.3	42

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73	Molecular alterations in the neurofibromatosis Type 2 gene and its protein rarely occurring in meningothelial meningiomas. Journal of Neurosurgery, 2001, 94, 111-117.	0.9	41
74	Novel amplicons on the short arm of chromosome 7 identified using high resolution array CGH contain over expressed genes in addition toEGFR in glioblastoma multiforme. Genes Chromosomes and Cancer, 2005, 44, 392-404.	1.5	41
75	Ploidy status and copy number aberrations in primary glioblastomas defined by integrated analysis of allelic ratios, signal ratios and loss of heterozygosity using 500K SNP Mapping Arrays. BMC Genomics, 2008, 9, 489.	1.2	40
76	Genetic fingerprinting of the development and progression of T-cell lymphoma in a murine model of atypical myeloproliferative disorder initiated by the ZNF198–fibroblast growth factor receptor-1 chimeric tyrosine kinase. Blood, 2009, 114, 1576-1584.	0.6	40
77	Constitutive Notch pathway activation in murine ZMYM2-FGFR1–induced T-cell lymphomas associated with atypical myeloproliferative disease. Blood, 2011, 117, 6837-6847.	0.6	40
78	Interaction of the transforming acidic coiled-coil 1 (TACC1) protein with ch-TOG and GAS41/NuBI1 suggests multiple TACC1-containing protein complexes in human cells. Biochemical Journal, 2002, 363, 195.	1.7	38
79	Identification of inactivating mutations in the JAK1, SYNJ2, and CLPTM1 genes in prostate cancer cells using inhibition of nonsense-mediated decay and microarray analysis. Cancer Genetics and Cytogenetics, 2005, 161, 97-103.	1.0	38
80	Development of a murine model for blastoid variant mantle-cell lymphoma. Blood, 2007, 109, 4899-4906.	0.6	38
81	Critical role of the WASF3 gene in JAK2/STAT3 regulation of cancer cell motility. Carcinogenesis, 2013, 34, 1994-1999.	1.3	38
82	Development of a Blastoid Variant, Mantle Cell Lymphoma Model in Transgenic Mice Blood, 2005, 106, 419-419.	0.6	37
83	Identification of consistent novel submegabase deletions in low-grade oligodendrogliomas using array-based comparative genomic hybridization. Genes Chromosomes and Cancer, 2005, 44, 85-96.	1.5	36
84	Gain of $1q$ Is a Potential Univariate Negative Prognostic Marker for Survival in Medulloblastoma. Clinical Cancer Research, 2007, 13, 7022-7028.	3.2	36
85	The Oncogenic Fusion Protein-tyrosine Kinase ZNF198/Fibroblast Growth Factor Receptor-1 Has Signaling Function Comparable with Interleukin-6 Cytokine Receptors. Journal of Biological Chemistry, 2003, 278, 16198-16208.	1.6	35
86	Occurrence and evolution of homogeneously staining regions may be due to breakage-fusion-bridge cycles following telomere loss. Chromosoma, 1983, 88, 216-221.	1.0	34
87	Genomic profiling of myeloid sarcoma by array comparative genomic hybridization. Genes Chromosomes and Cancer, 2005, 44, 373-383.	1.5	34
88	Genome Wide Copy Number Abnormalities in Pediatric Medulloblastomas as Assessed by Array Comparative Genome Hybridization. Brain Pathology, 2007, 17, 282-296.	2.1	34
89	Changes in DNA Content During In Vitro Transformation of Mouse Salivary Gland Epithelium2. Journal of the National Cancer Institute, 1980, 64, 1443-1449.	3.0	33
90	Molecular Definition of Chromosome Translocations Involving 10q24 and 19q13 in Human Malignant Glioma Cells. Cancer Genetics and Cytogenetics, 1998, 105, 60-68.	1.0	33

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91	The t(8;13) Atypical Myeloproliferative Disorder: Further Analysis of the ZNF198 Gene and Lack of Evidence for Multiple Genes Disrupted on Chromosome 13. Blood, 1998, 92, 1456-1458.	0.6	33
92	ZNF198, a zinc finger protein rearranged in myeloproliferative disease, localizes to the PML nuclear bodies and interacts with SUMO-1 and PML. Experimental Cell Research, 2006, 312, 3739-3751.	1.2	32
93	Identification and characterisation of constitutional chromosome abnormalities using arrays of bacterial artificial chromosomes. British Journal of Cancer, 2004, 90, 860-865.	2.9	31
94	Src Activation Plays an Important Key Role in Lymphomagenesis Induced by FGFR1 Fusion Kinases. Cancer Research, 2011, 71, 7312-7322.	0.4	31
95	Aberrant expression of the tumour suppressor gene p53 is very frequent in Wilms' tumours. Journal of Pathology, 1992, 168, 237-242.	2.1	29
96	Germline mutations in the RB1 gene in patients with hereditary retinoblastoma. Genes Chromosomes and Cancer, 1995, 14, 277-284.	1.5	29
97	HIF1A induces expression of the WASF3 metastasisâ€associated gene under hypoxic conditions. International Journal of Cancer, 2012, 131, E905-15.	2.3	29
98	Celecoxib Ameliorates Seizure Susceptibility in Autosomal Dominant Lateral Temporal Epilepsy. Journal of Neuroscience, 2018, 38, 3346-3357.	1.7	29
99	Chromosome analysis of human neuroblastoma cell line TR14 showing double minutes and an aberration involving chromosome 1. Cancer Genetics and Cytogenetics, 1983, 9, 273-280.	1.0	28
100	Characterization of FAM10A4, a Member of the ST13 Tumor Suppressor Gene Family That Maps to the 13q14.3 Region Associated with B-Cell Leukemia, Multiple Myeloma, and Prostate Cancer. Genomics, 2002, 80, 5-7.	1.3	28
101	Truncating mutations in the ACVR2 gene attenuates activin signaling in prostate cancer cells. Cancer Genetics and Cytogenetics, 2005, 163, 123-129.	1.0	27
102	Induction of the plasminogen activator inhibitor-2 in cells expressing the ZNF198/FGFR1 fusion kinase that is involved in atypical myeloproliferative disease. Blood, 2006, 107, 3693-3699.	0.6	27
103	Identification of genes involved in squamous cell carcinoma of the lung using synchronized data from DNA copy number and transcript expression profiling analysis. Lung Cancer, 2008, 59, 315-331.	0.9	27
104	Dysregulated signaling pathways in the development of CNTRL-FGFR1–induced myeloid and lymphoid malignancies associated with FGFR1 in human and mouse models. Blood, 2013, 122, 1007-1016.	0.6	27
105	miR-339 Promotes Development of Stem Cell Leukemia/Lymphoma Syndrome via Downregulation of the <i>BCL2L11</i> and <i>BAX</i> Proapoptotic Genes. Cancer Research, 2018, 78, 3522-3531.	0.4	27
106	The co-chaperone UNC45A is essential for the expression of mitotic kinase NEK7 and tumorigenesis. Journal of Biological Chemistry, 2019, 294, 5246-5260.	1.6	27
107	Genetic and Cytogenetic Analysis of Patients Showing Reduced Esterase-D Levels and Mental Retardation from a Survey of 500 Individuals with Retinoblastoma. Ophthalmic Paediatrics and Genetics, 1989, 10, 117-127.	0.4	26
108	Identification of YAC clones for human chromosome 1p32 and physical mapping of the infantile neuronal ceroid lipofuscinosis (INCL) locus. Genomics, 1995, 25, 404-412.	1.3	26

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109	Candidate glioblastoma development gene identification using concordance between copy number abnormalities and gene expression level changes. Genes Chromosomes and Cancer, 2007, 46, 875-894.	1.5	26
110	Identification of the promoter, genomic structure, and mouse ortholog of LGI1. Mammalian Genome, 2000, 11, 622-627.	1.0	25
111	Obtaining DNA from a geographically dispersed cohort of current and former smokers: Use of mail-based mouthwash collection and monetary incentives. Nicotine and Tobacco Research, 2004, 6, 439-446.	1.4	25
112	Molecular characterization of the 1p22 breakpoint region spanning the constitutional translocation breakpoint in a neuroblastoma patient with a $t(1;10)(p22;q21)$. Cancer Genetics and Cytogenetics, 1998, 100, 10-20.	1.0	24
113	ZNF198 protein, involved in rearrangement in myeloproliferative disease, forms complexes with the DNA repair-associated HHR6A/6B and RAD18 proteins. Oncogene, 2003, 22, 3417-3423.	2.6	24
114	The temporal and spatial expression pattern of the LGI1 epilepsy predisposition gene during mouse embryonic cranial development. BMC Neuroscience, 2011, 12, 43.	0.8	24
115	Acute Progression of BCR-FGFR1 Induced Murine B-Lympho/Myeloproliferative Disorder Suggests Involvement of Lineages at the Pro-B Cell Stage. PLoS ONE, 2012, 7, e38265.	1.1	24
116	EVI5 protein associates with the INCENP-aurora B kinase-survivin chromosomal passenger complex and is involved in the completion of cytokinesis. Experimental Cell Research, 2006, 312, 2325-2335.	1.2	23
117	Differential expression of the LGI and SLIT families of genes in human cancer cells. Gene, 2005, 356, 85-90.	1.0	22
118	Long tandem repeats as a form of genomic copy number variation: structure and length polymorphism of a chromosome 5p repeat in control and schizophrenia populations. Psychiatric Genetics, 2009, 19, 64-71.	0.6	22
119	Pediatric primary intramedullary spinal cord glioblastoma. Rare Tumors, 2010, 2, 135-141.	0.3	22
120	Reexpression of LGI1 in glioma cells results in dysregulation of genes implicated in the canonical axon guidance pathway. Genomics, 2010, 95, 93-100.	1.3	22
121	Evaluation of phosphatidylinositol-3-kinase catalytic subunit (PIK3CA) and epidermal growth factor receptor (EGFR) gene mutations in pancreaticobiliary adenocarcinoma. Journal of Gastrointestinal Oncology, 2013, 4, 20-9.	0.6	22
122	A Yeast Artificial Chromosome Contig That Spans the RB1-D13S31 Interval on Human Chromosome 13 and Encompasses the Frequently Deleted Region in B-cell Chronic Lymphocytic Leukemia. Genomics, 1995, 30, 425-430.	1.3	21
123	Cloning of the human Gfi-1 gene and its mapping to chromosome region 1p22. Oncogene, 1997, 14, 1003-1005.	2.6	21
124	EVI5 is a novel centrosomal protein that binds to \hat{l}_{\pm} - and \hat{l}_{3} -tubulin. Genomics, 2005, 86, 594-605.	1.3	21
125	Homozygous inactivation of the <i>LGI1</i> gene results in hypomyelination in the peripheral and central nervous systems. Journal of Neuroscience Research, 2010, 88, 3328-3336.	1.3	21
126	Analysis of Wilms Tumors Using SNP Mapping Array-Based Comparative Genomic Hybridization. PLoS ONE, 2011, 6, e18941.	1.1	21

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127	The pleiotropic effects of TNFî± in breast cancer subtypes is regulated by TNFAIP3/A20. Oncogene, 2019, 38, 469-482.	2.6	21
128	Molecular characterization of a $(1;10)(p22;q21)$ constitutional translocation from a patient with neuroblastoma. Cancer Genetics and Cytogenetics, 1995, 81, 151-157.	1.0	20
129	Telomerase activity in pancreatic endocrine tumors. American Journal of Gastroenterology, 2002, 97, 1022-1030.	0.2	20
130	Application of spectral karyotyping to the analysis of the human chromosome complement of interspecies somatic cell hybrids. Cancer Genetics and Cytogenetics, 2003, 142, 30-35.	1.0	20
131	Mass spectroscopy identifies the splicing-associated proteins, PSF, hnRNP H3, hnRNP A2/B1, and TLS/FUS as interacting partners of the ZNF198 protein associated with rearrangement in myeloproliferative disease. Experimental Cell Research, 2005, 309, 78-85.	1.2	20
132	Mass Spectrometry Identifies LGI1-Interacting Proteins that Are Involved in Synaptic Vesicle Function in the Human Brain. Journal of Molecular Neuroscience, 2009, 39, 137-143.	1.1	20
133	Targeting FGFR1 to suppress leukemogenesis in syndromic and <i>de novo</i> AML in murine models. Oncotarget, 2016, 7, 49733-49742.	0.8	20
134	FGFR1 fusion kinase regulation of MYC expression drives development of stem cell leukemia/lymphoma syndrome. Leukemia, 2018, 32, 2363-2373.	3.3	20
135	The ability of normal mouse cells to reduce the malignant potential of transformed mouse bladder epithelial cells depends on their somatic origin. International Journal of Cancer, 1984, 33, 657-667.	2.3	19
136	The aniridia-Wilms' tumour association: molecular and genetic analysis of chromosome deletions on the short arm of chromosome 11. Human Genetics, 1989, 82, 123-126.	1.8	19
137	Overlay analysis of the oligonucleotide array gene expression profiles and copy number abnormalities as determined by array comparative genomic hybridization in medulloblastomas. Genes Chromosomes and Cancer, 2007, 46, 53-66.	1.5	19
138	The promise of zebrafish as a chemical screening tool in cancer therapy. Future Medicinal Chemistry, 2015, 7, 1395-1405.	1.1	19
139	A chromosomal breakpoint that separates the esterase D and retinoblastoma predisposition loci in a patient with $del(13)(q14q31)$. Cancer Genetics and Cytogenetics, 1987, 27, 27-31.	1.0	18
140	TTC4,a Novel Human Gene Containing the Tetratricopeptide Repeat and Mapping to the Region of Chromosome 1p31 That Is Frequently Deleted in Sporadic Breast Cancer. Genomics, 1999, 55, 157-163.	1.3	18
141	Essential roles of leucine-rich glioma inactivated 1 in the development of embryonic and postnatal cerebellum. Scientific Reports, 2015, 5, 7827.	1.6	18
142	A novel missense mutation in patients from a retinoblastoma pedigree showing only mild expression of the tumor phenotype. Oncogene, 1998, 16, 3211-3213.	2.6	17
143	Development of ZMYM2â€FGFR1 driven AML in human CD34+ cells in immunocompromised mice. International Journal of Cancer, 2016, 139, 836-840.	2.3	17
144	FGFR1OP2-FGFR1 induced myeloid leukemia and T-cell lymphoma in a mouse model. Haematologica, 2016, 101, e91-e94.	1.7	17

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145	LGI1: From zebrafish to human epilepsy. Progress in Brain Research, 2014, 213, 159-179.	0.9	16
146	Homozygous Deletion of the <scp>LGI</scp> 1 Gene in Mice Leads to Developmental Abnormalities Resulting in Cortical Dysplasia. Brain Pathology, 2015, 25, 587-597.	2.1	16
147	Suppression of Breast Cancer Metastasis Using Stapled Peptides Targeting the WASF Regulatory Complex. Cancer Growth and Metastasis, 2017, 10, 117906441771319.	3.5	16
148	A transcription map of the minimally deleted region from 13q14 in B-cell chronic lymphocytic leukemia as defined by large scale sequencing of the 650 kb critical region. Oncogene, 2000, 19, 5772-5780.	2.6	15
149	Phosphorylation of the SSBP2 and ABL proteins by the ZNF198â€FGFR1 fusion kinase seen in atypical myeloproliferative disorders as revealed by phosphopeptideâ€specific MS. Proteomics, 2009, 9, 3979-3988.	1.3	15
150	Deletion of chromosome region 13q14 is transmissible and does not always predispose to retinoblastoma. Human Genetics, 1988, 80, 43-45.	1.8	14
151	Insertional inactivation of the WT1 gene in tumour cells from a patient with WAGR syndrome. Human Genetics, 1993, 92, 83-86.	1.8	14
152	Characterization of the breakpoints in a $t(8;13)(p11;q12)$ translocation from a patient with myeloproliferative disease using fluorescence in situ hybridization., 1998, 21, 160-165.		14
153	Identification of a Novel Gene (ADPRTL1) Encoding a Potential Poly(ADP-ribosyl)transferase Protein. Genomics, 1999, 62, 533-536.	1.3	14
154	Molecular characterization of a $7p15-21$ homozygous deletion in a Wilms tumor. Genes Chromosomes and Cancer, 2003, 36, 1-6.	1.5	14
155	Defined genetic events associated with the spontaneous in vitro transformation of ElA/Ras-expressing human IMR90 fibroblasts. Carcinogenesis, 2006, 27, 350-359.	1.3	14
156	Array comparative genome hybridization analysis of acute lymphoblastic leukaemia and acute megakaryoblastic leukaemia in patients with Down syndrome. British Journal of Haematology, 2008, 142, 934-945.	1.2	14
157	The miR-17/92 cluster is involved in the molecular etiology of the SCLL syndrome driven by the BCR-FGFR1 chimeric kinase. Oncogene, 2018, 37, 1926-1938.	2.6	14
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