Johji Inazawa

List of Publications by Year
in descending order

135 papers	5,075 citations	93792 39 h-index	129628 63 g-index
138 all docs	138 docs citations	138 times ranked	9096 citing authors

#	Article	IF	Citations
1	Genomic landscape of a mouse model of diffuse-type gastric adenocarcinoma. Gastric Cancer, 2022, 25, 83-95.	2.7	3
2	Clinical Utility of Germline Genetic Testing in Japanese Men Undergoing Prostate Biopsy. JNCI Cancer Spectrum, 2022, 6, pkac001.	1.4	3
3	Potential for reversing miR-634-mediated cytoprotective processes to improve efficacy of chemotherapy against oral squamous cell carcinoma. Molecular Therapy - Oncolytics, 2022, 24, 897-908.	2.0	4
4	A missense variant in NUF2, a component of the kinetochore NDC80 complex, causes impaired chromosome segregation and aneuploidy associated with microcephaly and short stature. Human Genetics, 2021, 140, 1047-1060.	1.8	5
5	Isolation and characterisation of lymphatic endothelial cells from lung tissues affected by lymphangioleiomyomatosis. Scientific Reports, 2021, 11, 8406.	1.6	5
6	Identification of PDHX as a metabolic target for esophageal squamous cell carcinoma. Cancer Science, 2021, 112, 2792-2802.	1.7	8
7	Cancer-associated miRNAs and their therapeutic potential. Journal of Human Genetics, 2021, 66, 937-945.	1.1	51
8	Augmentation of lenvatinib efficacy by topical treatment of miR-634 ointment in anaplastic thyroid cancer. Biochemistry and Biophysics Reports, 2021, 26, 101009.	0.7	1
9	Integrative genomeâ€wide analyses reveal the transcriptional aberrations in Japanese esophageal squamous cell carcinoma. Cancer Science, 2021, 112, 4377-4392.	1.7	6
10	miR-766-5p Targets Super-Enhancers by Downregulating CBP and BRD4. Cancer Research, 2021, 81, 5190-5201.	0.4	11
11	Concurrent targeting of MAP3K3 and BRD4 by miR-3140-3p overcomes acquired resistance to BET inhibitors in neuroblastoma cells. Molecular Therapy - Nucleic Acids, 2021, 25, 83-92.	2.3	4
12	Linkage-specific deubiquitylation by OTUD5 defines an embryonic pathway intolerant to genomic variation. Science Advances, 2021, 7, .	4.7	25
13	Therapeutic Potential of LNP-Mediated Delivery of miR-634 for Cancer Therapy. Molecular Therapy - Nucleic Acids, 2020, 19, 330-338.	2.3	43
14	Improving the Efficacy of EGFR Inhibitors by Topical Treatment of Cutaneous Squamous Cell Carcinoma with miR-634 Ointment. Molecular Therapy - Oncolytics, 2020, 19, 294-307.	2.0	17
15	Clinical efficacy of haematopoietic stem cell transplantation for adult adrenoleukodystrophy. Brain Communications, 2020, 2, fcz048.	1.5	14
16	Large-scale genome-wide association study in a Japanese population identifies novel susceptibility loci across different diseases. Nature Genetics, 2020, 52, 669-679.	9.4	304
17	miR-1293, a Candidate for miRNA-Based Cancer Therapeutics, Simultaneously Targets BRD4 and the DNA Repair Pathway. Molecular Therapy, 2020, 28, 1494-1505.	3.7	22
18	Clinical impact of hemizygous deletion detection and panel-size in comprehensive genomic profiling.. Journal of Clinical Oncology, 2020, 38, e15671-e15671.	0.8	0

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19	Massive computational identification of somatic variants in exonic splicing enhancers using The Cancer Genome Atlas. Cancer Medicine, 2019, 8, 7372-7384.	1.3	1
20	Copy number variation analysis in 83 children with early-onset developmental and epileptic encephalopathy after targeted resequencing of a 109-epilepsy gene panel. Journal of Human Genetics, 2019, 64, 1097-1106.	1.1	8
21	12 new susceptibility loci for prostate cancer identified by genome-wide association study in Japanese population. Nature Communications, 2019, 10, 4422.	5.8	49
22	Array comparative genomic hybridization analysis discloses chromosome copy number alterations as indicators of patient outcome in lymph node-negative breast cancer. BMC Cancer, 2019, 19, 521.	1.1	10
23	GWAS identifies two novel colorectal cancer loci at 16q24.1 and 20q13.12. Carcinogenesis, 2018, 39, 652-660.	1.3	52
24	The Japanese Society of Pathology Guidelines on the handling of pathological tissue samples for genomic research: Standard operating procedures based on empirical analyses. Pathology International, 2018, 68, 63-90.	0.6	44
25	Ovarian cancer therapeutic potential of glutamine depletion based on GS expression. Carcinogenesis, 2018, 39, 758-766.	1.3	29
26	Regional evaluation of childhood acute lymphoblastic leukemia genetic susceptibility loci among Japanese. Scientific Reports, 2018, 8, 789.	1.6	23
27	miR-3140 suppresses tumor cell growth by targeting BRD4 via its coding sequence and downregulates the BRD4-NUT fusion oncoprotein. Scientific Reports, 2018, 8, 4482.	1.6	25
28	Genomeâ€wide association study identifies gastric cancer susceptibility loci at 12q24.11â€12 and 20q11.21. Cancer Science, 2018, 109, 4015-4024.	1.7	39
29	Identification and characterization of transforming growth factor betaâ€induced in circulating tumor cell subline from pancreatic cancer cell line. Cancer Science, 2018, 109, 3623-3633.	1.7	11
30	Receptor tyrosine kinase amplification is predictive of distant metastasis in patients with oral squamous cell carcinoma. Cancer Science, 2017, 108, 256-266.	1.7	17
31	miR-432 Induces NRF2 Stabilization by Directly Targeting KEAP1. Molecular Cancer Research, 2017, 15, 1570-1578.	1.5	53
32	Genome-wide screening of DNA methylation associated with lymph node metastasis in esophageal squamous cell carcinoma. Oncotarget, 2017, 8, 37740-37750.	0.8	27
33	Comprehensive investigation of CASK mutations and other genetic etiologies in 41 patients with intellectual disability and microcephaly with pontine and cerebellar hypoplasia (MICPCH). PLoS ONE, 2017, 12, e0181791.	1.1	44
34	Subcloning and characterization of highly metastatic cells derived from human esophageal squamous cell carcinoma KYSE150 cells by in vivo selection. Oncotarget, 2017, 8, 34670-34677.	0.8	10
35	Cytogenetic analysis of spontaneously discharged products of conception by array-based comparative genomic hybridization. SpringerPlus, 2016, 5, 874.	1.2	8
36	Significant impact of miRNAâ€“target gene networks on genetics of human complex traits. Scientific Reports, 2016, 6, 22223.	1.6	44

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37	Exosomal microRNA miR-1246 induces cell motility and invasion through the regulation of DENND2D in oral squamous cell carcinoma. Scientific Reports, 2016, 6, 38750.	1.6	147
38	Association between maternal education and malocclusion in Mongolian adolescents: a cross-sectional study. BMJ Open, 2016, 6, e012283.	0.8	18
39	Association of variations in HLA class II and other loci with susceptibility to EGFR-mutated lung adenocarcinoma. Nature Communications, 2016, 7, 12451.	5.8	49
40	SNP array screening of cryptic genomic imbalances in 450 Japanese subjects with intellectual disability and multiple congenital anomalies previously negative for large rearrangements. Journal of Human Genetics, 2016, 61, 335-343.	1.1	7
41	Chromothripsis-like chromosomal rearrangements induced by ionizing radiation using proton microbeam irradiation system. Oncotarget, 2016, 7, 10182-10192.	0.8	44
42	Down-regulation of LAPTM5 in human cancer cells. Oncotarget, 2016, 7, 28320-28328.	0.8	32
43	Simultaneous Detection of Both Single Nucleotide Variations and Copy Number Alterations by Next-Generation Sequencing in Gorlin Syndrome. PLoS ONE, 2015, 10, e0140480.	1.1	16
44	Identical deletion at 14q13.3 including PAX9 and NKX2-1 in siblings from mosaicism of unaffected parent. Journal of Human Genetics, 2015, 60, 203-206.	1.1	13
45	miR-634 Activates the Mitochondrial Apoptosis Pathway and Enhances Chemotherapy-Induced Cytotoxicity. Cancer Research, 2015, 75, 3890-3901.	0.4	50
46	The oncogenic role of GASC1 in chemically induced mouse skin cancer. Mammalian Genome, 2015, 26, 591-597.	1.0	10
47	miR-544ainduces epithelialâ€“mesenchymal transition through the activation of WNT signaling pathway in gastric cancer. Carcinogenesis, 2015, 36, 1363-1371.	1.3	76
48	High Expression of p62 Protein Is Associated with Poor Prognosis and Aggressive Phenotypes in Endometrial Cancer. American Journal of Pathology, 2015, 185, 2523-2533.	1.9	42
49	Autophagy Inhibition Sensitizes Acute Lymphoblastic Leukemia Cells to L-Asparaginase. Blood, 2015, 126, 3772-3772.	0.6	3
50	NF-Î²B Inducing Kinase, a Central Signaling Component of the Non-Canonical Pathway of NF-Î²B, Contributes to Ovarian Cancer Progression. PLoS ONE, 2014, 9, e88347.	1.1	36
51	The Impact of miRNA-Based Molecular Diagnostics and Treatment of NRF2-Stabilized Tumors. Molecular Cancer Research, 2014, 12, 58-68.	1.5	64
52	Chromosome 9q33q34 Microdeletion With Early Infantile Epileptic Encephalopathy, Severe Dystonia, Abnormal Eye Movements, and Nephroureteral Malformations. Pediatric Neurology, 2014, 51, 170-175.	1.0	14
53	Î³-Glutamylcyclotransferase as a novel immunohistochemical biomarker for the malignancy of esophageal squamous tumors. Human Pathology, 2014, 45, 331-341.	1.1	26
54	High Expression of SQSTM1/p62 Protein Is Associated with Poor Prognosis in Epithelial Ovarian Cancer. Acta Histochemica Et Cytochemica, 2014, 47, 295-301.	0.8	44

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55	The Tumor-Suppressive miR-497-195 Cluster Targets Multiple Cell-Cycle Regulators in Hepatocellular Carcinoma. PLoS ONE, 2013, 8, e60155.	1.1	132
56	A Rare Genetic Polymorphism In C5 Confers Poor Response To The Anti-C5 Monoclonal Antibody Eculizumab In 11 Japanese Patients With PNH. Blood, 2013, 122, 3709-3709.	0.6	0
57	Tumor-suppressive microRNA silenced by tumor-specific DNA hypermethylation in cancer cells. Cancer Science, 2012, 103, 837-845.	1.7	43
58	A Rare Genetic Polymorphism in C5 Confers Poor Response to the Anti-C5 Monoclonal Antibody Eculizumab by Nine Japanese Patients with PNH. Blood, 2012, 120, 3197-3197.	0.6	3
59	Reproducibility, performance, and clinical utility of a genetic risk prediction model for prostate cancer in Japanese patients.. Journal of Clinical Oncology, 2012, 30, 10520-10520.	0.8	0
60	Clinical application of array-based comparative genomic hybridization by two-stage screening for 536 patients with mental retardation and multiple congenital anomalies. Journal of Human Genetics, 2011, 56, 110-124.	1.1	22
61	HECT-type Ubiquitin Ligase ITCH Targets Lysosomal-associated Protein Multispanning Transmembrane 5 (LAPTM5) and Prevents LAPTM5-mediated Cell Death. Journal of Biological Chemistry, 2011, 286, 44086-44094.	1.6	20
62	Frequent silencing of protocadherin 17 , a candidate tumour suppressor for esophageal squamous cell carcinoma. Carcinogenesis, 2010, 31, 1027-1036.	1.3	86
63	SKI and MEL1 Cooperate to Inhibit Transforming Growth Factor-Î² Signal in Gastric Cancer Cells. Journal of Biological Chemistry, 2009, 284, 3334-3344.	1.6	74
64	Overexpression of SMYD2 relates to tumor cell proliferation and malignant outcome of esophageal squamous cell carcinoma. Carcinogenesis, 2009, 30, 1139-1146.	1.3	154
65	PH Domain-Only Protein PHLDA3 Is a p53-Regulated Repressor of Akt. Cell, 2009, 136, 535-550.	13.5	201
66	Identification of SMURF1 as a possible target for 7q21.3â€22.1 amplification detected in a pancreatic cancer cell line by inâ€house arrayâ€based comparative genomic hybridization. Cancer Science, 2008, 99, 986-994.	1.7	35
67	Construction of a high-density and high-resolution human chromosome X array for comparative genomic hybridization analysis. Journal of Human Genetics, 2007, 52, 397-405.	1.1	20
68	Redefining the disease locus of 16q22.1-linked autosomal dominant cerebellar ataxia. Journal of Human Genetics, 2007, 52, 643-649.	1.1	28
69	Nuclear expression of cIAP-1, an apoptosis inhibiting protein, predicts lymph node metastasis and poor patient prognosis in head and neck squamous cell carcinomas. Cancer Letters, 2005, 224, 141-151.	3.2	65
70	Frequent Silencing of Low Density Lipoprotein Receptor-Related Protein 1B (LRP1B) Expression by Genetic and Epigenetic Mechanisms in Esophageal Squamous Cell Carcinoma. Cancer Research, 2004, 64, 3741-3747.	0.4	132
71	Comparative genomic hybridization (CGH)-arrays pave the way for identification of novel cancer-related genes. Cancer Science, 2004, 95, 559-563.	1.7	130
72	Translocation (1;22)(p36;q11.2) with concurrent del(22)(q11.2) resulted in homozygous deletion of SNF5/INI1 in a newly established cell line derived from extrarenal rhabdoid tumor. Journal of Human Genetics, 2004, 49, 586-589.	1.1	26

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73	GPC5 is a possible target for the 13q31-q32 amplification detected in lymphoma cell lines. Journal of Human Genetics, 2003, 48, 331-335.	1.1	32
74	Association of over-expressed TFDP1 with progression of hepatocellular carcinomas. Journal of Human Genetics, 2003, 48, 609-613.	1.1	47
75	PPM1D is a potential target for 17q gain in neuroblastoma. Cancer Research, 2003, 63, 1876-83.	0.4	197
76	The Xq22 Inversion Breakpoint Interrupted a Novel Ras-Like GTPase Gene in a Patient with Duchenne Muscular Dystrophy and Profound Mental Retardation. American Journal of Human Genetics, 2002, 71, 637-645.	2.6	37
77	Expression of cIAP1, a target for 11q22 amplification, correlates with resistance of cervical cancers to radiotherapy. Cancer Research, 2002, 62, 4860-6.	0.4	124
78	Familial nasal NK/T-cell lymphoma and pesticide use. American Journal of Hematology, 2001, 66, 145-147.	2.0	20
79	Chromosomal aberrations in colorectal cancers and liver metastases analyzed by comparative genomic hybridization. International Journal of Cancer, 2001, 94, 623-629.	2.3	72
80	Homozygous deletion in a neuroblastoma cell line defined by a high-density STS map spanning human chromosome band 1p36. Genes Chromosomes and Cancer, 2001, 31, 326-332.	1.5	21
81	Identification of target genes within an amplicon at 14q12-q13 in esophageal squamous cell carcinoma. Genes Chromosomes and Cancer, 2001, 32, 112-118.	1.5	29
82	An 8-cM interstitial deletion on 4q21-q22 in DNA from an infant with hepatoblastoma overlaps with a commonly deleted region in adult liver cancers. American Journal of Medical Genetics Part A, 2001, 103, 176-180.	2.4	15
83	A Novel Amplicon at 9p23-24 in Squamous Cell Carcinoma of the Esophagus That Lies Proximal toGASC1and HarborsNFIB. Japanese Journal of Cancer Research, 2001, 92, 423-428.	1.7	44
84	Identification of the homozygously deleted region at chromosome 1p36.2 in human neuroblastoma. Medical and Pediatric Oncology, 2000, 35, 516-521.	1.0	7
85	Detection of chromosomal aneusomy by fluorescence in situ hybridization in fine-needle aspirates from breast tumors. Cancer, 2000, 90, 373-378.	2.0	10
86	Amplification and over-expression of theAIB1 nuclear receptor co-activator gene in primary gastric cancers. International Journal of Cancer, 2000, 89, 217-223.	2.3	146
87	CD44 is a potential target of amplification within the 11p13 amplicon detected in gastric cancer cell lines. Genes Chromosomes and Cancer, 2000, 29, 315-324.	1.5	44
88	Cloning and expression of human B cell-specific transcription factor BACH2 mapped to chromosome 6q15. Oncogene, 2000, 19, 3739-3749.	2.6	61
89	Identification and characterization of a 500-kb homozygously deleted region at 1p36.2-p36.3 in a neuroblastoma cell line. Oncogene, 2000, 19, 4302-4307.	2.6	82
90	Molecular Cytogenetic Analysis of 17 Renal Cancer Cell Lines: Increased Copy Number at 5q31-33 in Cell Lines from Nonpapillary Carcinomas. Japanese Journal of Cancer Research, 2000, 91, 156-163.	1.7	29

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91	Detection of chromosomal aneusomy by fluorescence in situ hybridization in fine-needle aspirates from breast tumors. , 2000, 90, 373.		2
92	Amplification and overâ€expression of the AIB1 nuclear receptor coâ€activator gene in primary gastric cancers. International Journal of Cancer, 2000, 89, 217-223.	2.3	2
93	Repeat-directed isolation of a novel gene preferentially expressed from the maternal allele in human placenta. Journal of Human Genetics, 1999, 44, 1-9.	1.1	3
94	Gains, losses, and amplifications of genomic materials in primary gastric cancers analyzed by comparative genomic hybridization. , 1999, 24, 299-305.		153
95	Combined spectral karyotyping and DAPI banding analysis of chromosome abnormalities in myelodysplastic syndrome. Genes Chromosomes and Cancer, 1999, 26, 336-345.	1.5	93
96	Isolation of novel genes within the amplified regions in tumors detected by comparative genomic hybridization (CGH).. Seibutsu Butsuri Kagaku, 1999, 43, 91-95.	0.1	0
97	Loss of material from chromosome arm 1p during malignant progression of meningioma revealed by fluorescent in Situ hybridization. Cancer, 1998, 83, 360-366.	2.0	69
98	Grb10/GrbIR as anin vivosubstrate of Tec tyrosine kinase. Genes To Cells, 1998, 3, 431-441.	0.5	33
99	Identification of an Efs isoform that lacks the SH3 domain and chromosomal mapping of human Efs. Oncogene, 1997, 15, 1741-1745.	2.6	12
100	The Whole Nucleotide Sequence and Chromosomal Localization of the Gene for Human Metabotropic Glutamate Receptor Subtype 6. European Journal of Neuroscience, 1997, 9, 1226-1235.	1.2	32
101	Detection of numerical alterations of chromosomes 3, 7, 17 and X in low-grade intracystic papillary tumors of the breast by multi-color fluorescencein situ hybridization. Breast Cancer, 1997, 4, 247-252.	1.3	14
102	A complete Not I restriction map covering the entire long arm of human chromosome 11. Genes To Cells, 1997, 2, 345-357.	0.5	12
103	The brain finger protein gene (ZNF179), a member of the RING finger family, maps within the Smith-Magenis syndrome region at 17p11.2. American Journal of Medical Genetics Part A, 1997, 69, 320-324.	2.4	14
104	The brain finger protein gene (ZNF179), a member of the RING finger family, maps within the Smith-Magenis syndrome region at 17p11.2. , 1997, 69, 320.		1
105	Assignment of the human ST2 gene to chromosome 2 at q11.2. Human Genetics, 1996, 97, 561-563.	1.8	27
106	Molecular cloning of the breakpoint of t(11;22)(q23;q11) chromosome translocation in an adult acute myelomonocytic leukaemia. British Journal of Haematology, 1996, 92, 687-691.	1.2	10
107	p53 gene mutation is not directly related to tumoricidal effects of preoperative radiochemohyperthermia therapy for rectal cancers. , 1996, 63, 87-90.		4
108	Analysis of numerical aberrations in specific chromosomes by fluorescent in situ hybridization as a diagnostic tool in breast cancer. , 1996, 77, 2064-2069.		73

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109	Multiple Primary Cancers with Microsatellite Instability: Report of a Case. Japanese Journal of Cancer Research, 1996, 87, 105-108.	1.7	3
110	Analysis of numerical aberrations in specific chromosomes by fluorescent in situ hybridization as a diagnostic tool in breast cancer. , 1996, 77, 2064.		6
111	Assignment of the human ST2 gene to chromosome 2 at q11.2. Human Genetics, 1996, 97, 561-563.	1.8	3
112	Induction of mcl1/EAT, Bcl-2 Related Gene, by Retinoic Acid or Heat Shock in the Human Embryonal Carcinoma Cells, NCR-G3.. Cell Structure and Function, 1996, 21, 143-150.	0.5	21
113	Multi-color FISH: Application to the Simultaneous Detection of Chromosome Aberrations in Neuro-epithelial Tumor Cell Lines.. Acta Histochemica Et Cytochemica, 1995, 28, 569-573.	0.8	0
114	Frequent deletions of material from chromosome arm 1p in oligodendroglial tumors revealed by double-target fluorescence in situ hybridization and microsatellite analysis. Genes Chromosomes and Cancer, 1995, 14, 295-300.	1.5	35
115	Reexamination of chromosomal loci of human muscle actin genes by fluorescencein situ hybridization. Japanese Journal of Human Genetics, 1995, 40, 145-148.	0.8	4
116	Fluorescencein situ hybridization analysis of chromosomal localization of three human cytochrome P450 2C genes (CYP2C8, 2C9, and 2C10) at 10q24.1. Japanese Journal of Human Genetics, 1994, 39, 337-343.	0.8	17
117	Myelomonocytic crisis with t(5;17) and a p53 mutation in a patient with chronic myelogenous leukemia. American Journal of Hematology, 1994, 45, 335-340.	2.0	5
118	Isolation, Characterization and Structural Organization of the Gene and Pseudogene for the Dihydrolipoamide Succinyltransferase Component of the Human 2-Oxoglutarate Dehydrogenase Complex. FEBS Journal, 1994, 224, 179-189.	0.2	33
119	Chromosomal localization of the protein tyrosine phosphatase G1 gene and characterization of the aberrant transcripts in human colon cancer cells. FEBS Letters, 1994, 339, 222-228.	1.3	12
120	Expression and chromosomal assignment of PTPH1 gene encoding a cytosolic protein tyrosine phosphatase homologous to cytoskeletal-associated proteins. International Journal of Cancer, 1993, 55, 947-951.	2.3	12
121	Chromosomal Assignment of the Gene for Protein Tyrosine Phosphatase HPTPÎ´. Japanese Journal of Cancer Research, 1993, 84, 1219-1222.	1.7	1
122	A Simple G-Banding Technique Adaptable for Fluorescent in situ Hybridization (FISH) and Physical Ordering of Human Renin (REN) and Catepsin E (CTSE) Genes by Multi-Color FISH.. Acta Histochemica Et Cytochemica, 1993, 26, 319-324.	0.8	10
123	Physical Ordering of Three Polymorphic DNA Markers Spanning the Regions Containing a Tumor Suppressor Gene of Renal Cell Carcinoma by Three-color Fluorescentin situHybridization. Japanese Journal of Cancer Research, 1992, 83, 1248-1252.	1.7	46
124	Assignment of the human cytochrome P-450 nifedipine oxidase gene (CYP3A4) to chromosome 7 at band q22.1 by fluorescencein situ hybridization. Japanese Journal of Human Genetics, 1992, 37, 133-138.	0.8	53
125	Structure of the human ferrochelatase gene. Exon/intron gene organization and location of the gene to chromosome 18. FEBS Journal, 1992, 205, 217-222.	0.2	150
126	Different drug sensitivity in two neuroblastoma cell lines established from the same patient before and after chemotherapy. International Journal of Cancer, 1991, 47, 732-737.	2.3	39

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127	Alpha-smooth-muscle actin and desmin expressions in human neuroblastoma cell lines. International Journal of Cancer, 1991, 48, 277-283.	2.3	40
128	Carcinoembryonic antigen(CEA) family genes are located on human chromosome 19 at band q13.2.. Acta Histochemica Et Cytochemica, 1990, 23, 411-422.	0.8	0
129	Characterization of an embryonal rhabdomyosarcoma cell line showing amplification and over-expression of the N-myc oncogene. International Journal of Cancer, 1990, 45, 705-711.	2.3	25
130	The unbalanced 1;7 translocation inde novo myelodysplastic syndrome and its clinical implication. Cancer, 1990, 65, 1350-1354.	2.0	42
131	A case of incomplete DiGeorge syndrome associated with partial monosomy 22q11.1 due to maternal 14;22 translocation. Japanese Journal of Human Genetics, 1989, 34, 235-241.	0.8	3
132	Diverse responses to retinoid in morphological differentiation, tumorigenesis and n-myc expression in human neuroblastoma sublines. International Journal of Cancer, 1989, 44, 286-291.	2.3	9
133	Distribution of breakpoint within the breakpoint cluster region (bcr) in chronic myelogenous leukemia with a complex philadelphia chromosome translocation. American Journal of Hematology, 1989, 32, 194-199.	2.0	7
134	Rejoining between 9q+ and Philadelphia chromosomes results in normal-looking chromosomes 9 and 22 in Ph1-negative chronic myelocytic leukemia. Human Genetics, 1989, 83, 115-118.	1.8	39
135	DNA analysis using long-term preserved fixed cytogenetic preparations. Japanese Journal of Human Genetics, 1988, 33, 417-421.	0.8	5