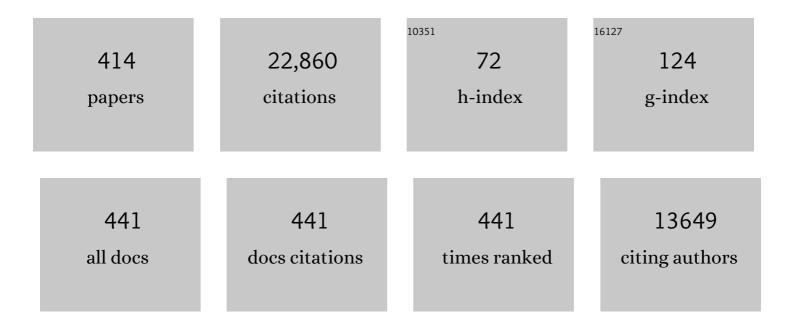
List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Transcriptomics paving the way for improved diagnostics and precision medicine of acute leukemia. Seminars in Cancer Biology, 2022, 84, 40-49.	4.3	11
2	HUGO Gene Nomenclature Committee (HGNC) recommendations for the designation of gene fusions. Leukemia, 2021, 35, 3040-3043.	3.3	42
3	Farewell message from the Editorâ€inâ€Chief of <i>Genes</i> , <i>Chromosomes</i> & <i>Cancer</i> . Genes Chromosomes and Cancer, 2020, 59, 3-4.	1.5	0
4	Analysis of fusion transcripts indicates widespread deregulation of snoRNAs and their host genes in breast cancer. International Journal of Cancer, 2020, 146, 3343-3353.	2.3	8
5	Most gene fusions in cancer are stochastic events. Genes Chromosomes and Cancer, 2019, 58, 607-611.	1.5	33
6	Cancer chromosome breakpoints cluster in geneâ€rich genomic regions. Genes Chromosomes and Cancer, 2019, 58, 149-154.	1.5	10
7	Frequent miRNA-convergent fusion gene events in breast cancer. Nature Communications, 2017, 8, 788.	5.8	24
8	ldentification of ETV6-RUNX1-like and DUX4-rearranged subtypes in paediatric B-cell precursor acute lymphoblastic leukaemia. Nature Communications, 2016, 7, 11790.	5.8	225
9	Gene fusions in soft tissue tumors: Recurrent and overlapping pathogenetic themes. Genes Chromosomes and Cancer, 2016, 55, 291-310.	1.5	107
10	The emerging complexity of gene fusions in cancer. Nature Reviews Cancer, 2015, 15, 371-381.	12.8	544
11	Whole-exome sequencing of pediatric acute lymphoblastic leukemia. Leukemia, 2012, 26, 1602-1607.	3.3	34
12	Diseaseâ€associated patterns of disomic chromosomes in hyperhaploid neoplasms. Genes Chromosomes and Cancer, 2012, 51, 536-544.	1.5	26
13	The correlation pattern of acquired copy number changes in 164 ETV6/RUNX1-positive childhood acute lymphoblastic leukemias. Human Molecular Genetics, 2010, 19, 3150-3158.	1.4	46
14	Generation of trisomies in cancer cells by multipolar mitosis and incomplete cytokinesis. Proceedings of the United States of America, 2010, 107, 20489-20493.	3.3	67
15	Identification by fluorescence of the G chromosome lost in human meningomas. Hereditas, 2009, 71, 163-168.	0.5	108
16	Isozymes of experimentally induced primary sarcomas in the rat. Hereditas, 2009, 73, 318-321.	0.5	1
17	The specificity of chromosome A2 involvement in DMBA-induced rat sarcomas. Hereditas, 2009, 77, 263-280.	0.5	54
18	Clustering of aberrations to specific chromosomes in human neoplasms:. Hereditas, 2009, 82, 167-174.	0.5	92

#	Article	IF	CITATIONS
19	Chromosomes and cancer. Hereditas, 2009, 86, 15-29.	0.5	129
20	Lysosomal enzymes in rat sarcomas induced by 7,12-dimethylbenz(α)anthracene and Rous sarcoma virus. Hereditas, 2009, 86, 103-106.	0.5	1
21	Prophase analysis of ring chromosome 13-an attempt at phenotypekaryotype correlation. Hereditas, 2009, 93, 231-233.	0.5	5
22	Cytogenetic study of pesticides in agricultural work. Hereditas, 2009, 92, 177-178.	0.5	20
23	Clustering of aberrations to specific chromosomes in human neoplasms. Hereditas, 2009, 95, 79-139.	0.5	253
24	The interrelations of micronuclei, chromosomal instability, and mutational activity in acute non-lymphocytic leukemia-A hypothesis. Hereditas, 2009, 95, 165-167.	0.5	9
25	Molecular screening for new fusion genes in cancer. Nature Genetics, 2008, 40, 685-686.	9.4	34
26	The prognostic implication of cytogenetic findings in non-Hodgkin lymphomas. Clinical Genetics, 2008, 29, 464-464.	1.0	0
27	Chromosome abnormalities in mesenchymal neoplasms. Clinical Genetics, 2008, 29, 466-467.	1.0	0
28	Chromosome aberrations and micronuclei in bone marrow cells and peripheral blood lymphocytes in humans exposed to ethylene oxide. Hereditas, 2008, 98, 105-113.	0.5	78
29	Comparison between five Nordic laboratories on scoring of human lymphocyte chromosome aberrations. Hereditas, 2008, 100, 209-218.	0.5	33
30	Increased sister chromatid exchange frequency in chronic alcoholic users. Hereditas, 2008, 101, 265-266.	0.5	10
31	Nonrandom de novo chromosome aberrations in human lymphocytes and amniotic cells. Hereditas, 2008, 102, 33-38.	0.5	21
32	Cytogenetic studies of childhood non-Hodgkin lymphomas. Hereditas, 2008, 103, 77-84.	0.5	30
33	C-band polymorphism in non-Hodgkin lymphoma. Hereditas, 2008, 103, 85-87.	0.5	5
34	Proliferation-specific and differentiation-associated chromosomal breakpoints in human neoplasia-a unifying model. Hereditas, 2008, 104, 307-312.	0.5	24
35	Breakprone chromosome bands in fibroblasts from patients with non-Hodgkin's lymphoma do not coincide with bands involved in primary rearrangements in non-Hodgkin's lymphomas. Hereditas, 2008, 109, 131-137.	0.5	4
36	Constitutional C-band polymorphism in lymphocytes from patients with chronic myeloid leukemia. Hereditas, 2008, 110, 145-148.	0.5	4

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37	Telomeric location of retroviral oncogenes in humans. Hereditas, 2008, 114, 207-211.	0.5	14
38	Wolf-Hirschhorn syndrome and balanced (4;10) translocation in the father. Clinical Genetics, 2008, 12, 101-103.	1.0	10
39	Chromosome Aberration in Human Neoplasm: Occurrence and Significance. Clinical Genetics, 2008, 14, 300-300.	1.0	Ο
40	ISCN (2005) is not acceptable for describing clonal evolution in cancer. Genes Chromosomes and Cancer, 2007, 46, 213-214.	1.5	9
41	Response to letter by the ISCN standing committee. Genes Chromosomes and Cancer, 2007, 46, 516-516.	1.5	1
42	The impact of translocations and gene fusions on cancer causation. Nature Reviews Cancer, 2007, 7, 233-245.	12.8	1,191
43	A novel and cytogenetically cryptic t(7;21)(p22;q22) in acute myeloid leukemia results in fusion of RUNX1 with the ubiquitin-specific protease gene USP42. Leukemia, 2006, 20, 224-229.	3.3	62
44	A gene fusion network in human neoplasia. Oncogene, 2006, 25, 2674-2678.	2.6	29
45	Gene expression profiling of leukemic cell lines reveals conserved molecular signatures among subtypes with specific genetic aberrations. Leukemia, 2005, 19, 1042-1050.	3.3	65
46	Formation of der(19)t(1;19)(q23;p13) in acute lymphoblastic leukemia. Genes Chromosomes and Cancer, 2005, 42, 144-148.	1.5	17
47	Statistical behavior of complex cancer karyotypes. Genes Chromosomes and Cancer, 2005, 42, 327-341.	1.5	53
48	Prevalence estimates of recurrent balanced cytogenetic aberrations and gene fusions in unselected patients with neoplastic disorders. Genes Chromosomes and Cancer, 2005, 43, 350-366.	1.5	56
49	Ewing tumours and synovial sarcomas have critical features of karyotype evolution in common with epithelial tumours. International Journal of Cancer, 2005, 116, 401-406.	2.3	11
50	Molecular signatures in childhood acute leukemia and their correlations to expression patterns in normal hematopoietic subpopulations. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 19069-19074.	3.3	99
51	Structural and numerical chromosome changes in colon cancer develop through telomere-mediated anaphase bridges, not through mitotic multipolarity. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 5541-5546.	3.3	161
52	ldentification of a commonly amplified 4.3 Mb region with overexpression of C8FW, but not MYC in MYC-containing double minutes in myeloid malignancies. Human Molecular Genetics, 2004, 13, 1479-1485.	1.4	56
53	High frequencies of chromosomal aberrations in multiple myeloma and monoclonal gammopathy of undetermined significance in direct chromosome preparation. British Journal of Haematology, 2004, 126, 487-494.	1.2	10
54	Fusion genes and rearranged genes as a linear function of chromosome aberrations in cancer. Nature Genetics, 2004, 36, 331-334.	9.4	286

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55	Clinical impact of internal tandem duplications and activating point mutations in FLT3 in acute myeloid leukemia in elderly patients. European Journal of Haematology, 2004, 72, 307-313.	1.1	52
56	Statistical analyses of karyotypic complexity in head and neck squamous cell carcinoma. Cancer Genetics and Cytogenetics, 2004, 150, 1-8.	1.0	18
57	Wilms tumors develop through two distinct karyotypic pathways. Cancer Genetics and Cytogenetics, 2004, 150, 9-15.	1.0	28
58	Dissecting karyotypic patterns in renal cell carcinoma: an analysis of the accumulated cytogenetic data. Cancer Genetics and Cytogenetics, 2004, 153, 1-9.	1.0	41
59	Statistical dissection of cytogenetic patterns in lung cancer reveals multiple modes of karyotypic evolution independent of histological classification. Cancer Genetics and Cytogenetics, 2004, 154, 99-109.	1.0	21
60	Identification of cytogenetic subgroups and karyotypic pathways of clonal evolution in follicular lymphomas. Genes Chromosomes and Cancer, 2004, 39, 195-204.	1.5	114
61	A model for karyotypic evolution in testicular germ cell tumors. Genes Chromosomes and Cancer, 2004, 40, 172-178.	1.5	17
62	MDS/AML-associated cytogenetic abnormalities in multiple myeloma and monoclonal gammopathy of undetermined significance: Evidence for frequent de novo occurrence and multipotent stem cell involvement of del(20q). Genes Chromosomes and Cancer, 2004, 41, 223-231.	1.5	26
63	MLL/GRAFfusion in an infant acute monocytic leukemia (AML M5b) with a cytogenetically cryptic ins(5;11)(q31;q23q23). Genes Chromosomes and Cancer, 2004, 41, 400-404.	1.5	17
64	Dissecting karyotypic patterns in malignant melanomas: Temporal clustering of losses and gains in melanoma karyotypic evolution. International Journal of Cancer, 2004, 108, 57-65.	2.3	122
65	Clinical and biological importance of cytogenetic abnormalities in childhood and adult acute lymphoblastic leukemia. Annals of Medicine, 2004, 36, 492-503.	1.5	47
66	The Breakpoint Region of the Most Common Isochromosome, i(17q), in Human Neoplasia Is Characterized by a Complex Genomic Architecture with Large, Palindromic, Low-Copy Repeats. American Journal of Human Genetics, 2004, 74, 1-10.	2.6	122
67	Evidence for a Single Step Mechanism in the Origin of Hyperdiploid Childhood Acute Lymphoblastic Leukemia Blood, 2004, 104, 1966-1966.	0.6	1
68	Gene Expression Profiling of Leukemic Cell Lines and Primary Leukemias Reveals Conserved Molecular Signatures among Subtypes with Specific Genetic Aberrations: Identification of Fusion Gene-Specific Transcriptional Profiles and Expression Pattern of Tyrosine Kinase-Encoding Genes Blood, 2004, 104, 2044-2044.	0.6	5
69	Genomic characterization ofMOZ/CBP andCBP/MOZ chimeras in acute myeloid leukemia suggests the involvement of a damage-repair mechanism in the origin of the t(8;16)(p11;p13). Genes Chromosomes and Cancer, 2003, 36, 90-98.	1.5	29
70	Fusion of theNUP98gene and the homeobox geneHOXC13in acute myeloid leukemia with t(11;12)(p15;q13). Genes Chromosomes and Cancer, 2003, 36, 107-112.	1.5	51
71	Introduction to special issue for Alfred G. Knudson. Genes Chromosomes and Cancer, 2003, 38, 285-285.	1.5	0
72	A pooled analysis of karyotypic patterns, breakpoints and imbalances in 783 cytogenetically abnormal multiple myelomas reveals frequently involved chromosome segments as well as significant age- and sex-related differences. British Journal of Haematology, 2003, 120, 960-969.	1.2	64

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73	Clinical and genetic studies of ETV6/ABL1-positive chronic myeloid leukaemia in blast crisis treated with imatinib mesylate. British Journal of Haematology, 2003, 122, 85-93.	1.2	38
74	Formation of trisomies and their parental origin in hyperdiploid childhood acute lymphoblastic leukemia. Blood, 2003, 102, 3010-3015.	0.6	47
75	A novel gene, MSI2, encoding a putative RNA-binding protein is recurrently rearranged at disease progression of chronic myeloid leukemia and forms a fusion gene with HOXA9 as a result of the cryptic t(7;17)(p15;q23). Cancer Research, 2003, 63, 1202-6.	0.4	69
76	Ovarian carcinoma develops through multiple modes of chromosomal evolution. Cancer Research, 2003, 63, 3378-85.	0.4	43
77	Power law distribution of chromosome aberrations in cancer. Cancer Research, 2003, 63, 7094-7.	0.4	20
78	Cytogenetic and Molecular Genetic Evolution of Chronic Myeloid Leukemia. Acta Haematologica, 2002, 107, 76-94.	0.7	405
79	Coping with complexity. Cancer Genetics and Cytogenetics, 2002, 135, 103-109.	1.0	62
80	Cytogenetic features of multiple myeloma: impact of gender, age, disease phase, culture time, and cytokine stimulation. European Journal of Haematology, 2002, 68, 345-353.	1.1	19
81	Acute myeloid leukemia with inv(16)(p13q22): Involvement of cervical lymph nodes and tonsils is common and may be a negative prognostic sign. American Journal of Hematology, 2002, 71, 15-19.	2.0	19
82	Expression ofNUP98/TOP1, but not ofTOP1/NUP98, in a treatment-related myelodysplastic syndrome with t(10;20;11)(q24;q11;p15). Genes Chromosomes and Cancer, 2002, 34, 249-254.	1.5	15
83	Multicolor COBRA-FISH analysis of chronic myeloid leukemia reveals novel cryptic balanced translocations during disease progression. Genes Chromosomes and Cancer, 2002, 35, 127-137.	1.5	23
84	RT-PCR analysis of acute myeloid leukemia with t(8;16)(p11;p13): Identification of a novelMOZ/CBP transcript and absence ofCBP/MOZ expression. Genes Chromosomes and Cancer, 2002, 35, 372-374.	1.5	19
85	Pooled analysis of clinical and cytogenetic features in treatment-related and de novo adult acute myeloid leukemia and myelodysplastic syndromes based on a consecutive series of 761 patients analyzed 1976–1993 and on 5098 unselected cases reported in the literature 1974–2001. Leukemia, 2002, 16, 2366-2378.	3.3	239
86	Multivariate analysis of chromosomal imbalances in breast cancer delineates cytogenetic pathways and reveals complex relationships among imbalances. Cancer Research, 2002, 62, 2675-80.	0.4	38
87	Prognostically important chromosomal aberrations in soft tissue sarcomas: a report of the Chromosomes and Morphology (CHAMP) Study Group. Cancer Research, 2002, 62, 3980-4.	0.4	20
88	Dissecting karyotypic patterns in colorectal tumors: two distinct but overlapping pathways in the adenoma-carcinoma transition. Cancer Research, 2002, 62, 5939-46.	0.4	48
89	The prognostic impact of karyotypic subgroups in myelodysplastic syndromes is strongly modified by sex. British Journal of Haematology, 2001, 113, 347-356.	1.2	28
90	Multivariate analyses of genomic imbalances in solid tumors reveal distinct and converging pathways of karyotypic evolution. Genes Chromosomes and Cancer, 2001, 31, 156-171.	1.5	85

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91	Fusion of theBCRand the fibroblast growth factor receptor-1 (FGFR1) genes as a result of t(8;22)(p11;q11) in a myeloproliferative disorder: The first fusion gene involvingBCRbut notABL. Genes Chromosomes and Cancer, 2001, 32, 302-310.	1.5	90
92	Characterization of chromosome aberrations in salivary gland tumors by FISH, including multicolor COBRA-FISH. Genes Chromosomes and Cancer, 2001, 30, 161-167.	1.5	57
93	Isodicentric 7p, idic(7)(q11.2), in acute myeloid leukemia associated with older age and favorable response to induction chemotherapy: A new clinical entity?. Genes Chromosomes and Cancer, 2001, 30, 261-266.	1.5	5
94	Paired multiplex reverse-transcriptase polymerase chain reaction (PMRT-PCR) analysis as a rapid and accurate diagnostic tool for the detection of MLL fusion genes in hematologic malignancies. Leukemia, 2001, 15, 1293-1300.	3.3	38
95	The incidence of trisomy 8 as a sole chromosomal aberration in myeloid malignancies varies in relation to gender, age, prior iatrogenic genotoxic exposure, and morphology. Cancer Genetics and Cytogenetics, 2001, 130, 160-165.	1.0	59
96	Telomere dysfunction triggers extensive DNA fragmentation and evolution of complex chromosome abnormalities in human malignant tumors. Proceedings of the National Academy of Sciences of the United States of America, 2001, 98, 12683-12688.	3.3	381
97	Fusion of the MORF and CBP genes in acute myeloid leukemia with the t(10;16)(q22;p13). Human Molecular Genetics, 2001, 10, 395-404.	1.4	190
98	Are occupational, hobby, or lifestyle exposures associated with Philadelphia chromosome positive chronic myeloid leukaemia?. Occupational and Environmental Medicine, 2001, 58, 722-727.	1.3	39
99	Cytogenetic characterization of peripheral nerve sheath tumours: a report of the CHAMP study group. , 2000, 190, 31-38.		141
100	Granulocytic sarcomas in body cavities in childhood acute myeloid leukemias with 11q23/MLL rearrangements. Genes Chromosomes and Cancer, 2000, 27, 136-142.	1.5	28
101	Karyotypic characterization of urinary bladder transitional cell carcinomas. Genes Chromosomes and Cancer, 2000, 29, 256-265.	1.5	52
102	RT-PCR analysis of theMOZ-CBP andCBP-MOZ chimeric transcripts in acute myeloid leukemias with t(8;16)(p11;p13). Genes Chromosomes and Cancer, 2000, 28, 415-424.	1.5	39
103	Cytogenetic evidence of clonality in cutaneous benign fibrous histiocytomas: a report of the CHAMP Study Group. Histopathology, 2000, 37, 212-217.	1.6	60
104	Survival time in a population-based consecutive series of adult acute myeloid leukemia – the prognostic impact of karyotype during the time period 1976–1993. Leukemia, 2000, 14, 1039-1043.	3.3	15
105	Cytogenetic and FISH studies of a single center consecutive series of 152 childhood acute lymphoblastic leukemias. European Journal of Haematology, 2000, 65, 40-51.	1.1	72
106	Recurrent chromosome aberrations in fibrous dysplasia of the bone. Cancer Genetics and Cytogenetics, 2000, 122, 30-32.	1.0	44
107	Radiation-Associated Sarcomas are Characterized by Complex Karyotypes with Frequent Rearrangements of Chromosome Arm 3p. Cancer Genetics and Cytogenetics, 2000, 116, 89-96.	1.0	55
108	Comparative Cytogenetic Study of Spindle Cell and Pleomorphic Leiomyosarcomas of Soft Tissues. Cancer Genetics and Cytogenetics, 2000, 116, 66-73.	1.0	60

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109	Cytogenetic-Morphologic Correlations in Aneurysmal Bone Cyst, Giant Cell Tumor of Bone and Combined Lesions. A Report from the CHAMP Study Group. Modern Pathology, 2000, 13, 1206-1210.	2.9	101
110	Chromosomal breakage-fusion-bridge events cause genetic intratumor heterogeneity. Proceedings of the United States of America, 2000, 97, 5357-5362.	3.3	368
111	Cytogenetic, Clinical, and Morphologic Correlations in 78 Cases of Fibromatosis: A Report from the CHAMP Study Group. Modern Pathology, 2000, 13, 1080-1085.	2.9	99
112	Recurrent chromosome aberrations in cancer. Mutation Research - Reviews in Mutation Research, 2000, 462, 247-253.	2.4	291
113	Acute Myeloid Leukemia with inv(8)(p11q13). Leukemia and Lymphoma, 2000, 39, 651-656.	0.6	19
114	Cytogenetic monoclonality in multifocal uroepithelial carcinomas: evidence of intraluminal tumour seeding. British Journal of Cancer, 1999, 81, 6-12.	2.9	50
115	The structure and dynamics of ring chromosomes in human neoplastic and non-neoplastic cells. Human Genetics, 1999, 104, 315-325.	1.8	108
116	Cytogenetic polyclonality in hematologic malignancies. , 1999, 24, 222-229.		27
117	Cytogenetic analysis of 363 consecutively ascertained diffuse large B-cell lymphomas. , 1999, 25, 123-133.		144
118	Different patterns of chromosomal imbalances in metastasising and non-metastasising primary breast carcinomas. , 1999, 84, 370-375.		21
119	Correlation between Clinicopathological Features and Karyotype in Spindle Cell Sarcomas. American Journal of Pathology, 1999, 154, 1841-1847.	1.9	109
120	Genetic changes in bone and soft tissue tumors. Acta Orthopaedica, 1999, 70, 30-40.	1.4	3
121	A singleâ€center populationâ€based consecutive series of 1500 cytogenetically investigated adult hematological malignancies: karyotypic features in relation to morphology, age and gender. European Journal of Haematology, 1999, 62, 95-102.	1.1	52
122	Prognostic implications of cytogenetic aberrations in diffuse large B ell lymphomas. European Journal of Haematology, 1999, 62, 184-190.	1.1	21
123	Cytogenetic analysis of 363 consecutively ascertained diffuse large B-cell lymphomas. , 1999, 25, 123.		4
124	Fluorescence in situ hybridization analyses of hematologic malignancies reveal frequent cytogenetically unrecognized 12p rearrangements. Leukemia, 1998, 12, 390-400.	3.3	48
125	ETV6/ABL fusion is rare in Ph-negative chronic myeloid disorders. Leukemia, 1998, 12, 1167-1168.	3.3	11
126	Molecular characterization of jumping translocations reveals spatial and temporal breakpoint heterogeneity. Leukemia, 1998, 12, 1411-1416.	3.3	19

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127	Cytogenetic and molecular genetic demonstration of polyclonality in an acinic cell carcinoma. British Journal of Cancer, 1998, 78, 292-295.	2.9	10
128	Cytogenetic analysis of four angiosarcomas from deep and superficial soft tissue. Cancer Genetics and Cytogenetics, 1998, 100, 52-56.	1.0	46
129	Nonrandom Pattern of Telomeric Associations in Atypical Lipomatous Tumors with Ring and Giant Marker Chromosomes. Cancer Genetics and Cytogenetics, 1998, 103, 25-34.	1.0	38
130	Genomic Amplification of CCND2 Is Rare in Non-Hodgkin Lymphomas. Cancer Genetics and Cytogenetics, 1998, 102, 81-82.	1.0	8
131	Comparison of Chromosomal Patterns with Clinical Features in 165 Lipomas: A Report of the CHAMP Study Group. Cancer Genetics and Cytogenetics, 1998, 102, 46-49.	1.0	60
132	Nonrandom Numerical Chromosome Abnormalities in Basal Cell Carcinomas. Cancer Genetics and Cytogenetics, 1998, 103, 35-42.	1.0	23
133	Chromosomal Abnormalities in Two Bladder Carcinomas with Secondary Squamous Cell Differentiation. Cancer Genetics and Cytogenetics, 1998, 102, 125-130.	1.0	24
134	Cytogenetic findings in invasive breast carcinomas with prognostically favourable histology: A less complex karyotypic pattern?. , 1998, 79, 361-364.		17
135	Cytogenetic and fluorescence in situ hybridization analyses of chromosome 19 aberrations in pancreatic carcinomas: Frequent loss of 19p13.3 and gain of 19q13.1-13.2. Genes Chromosomes and Cancer, 1998, 21, 8-16.	1.5	49
136	Cytogenetic analysis of 46 pleomorphic soft tissue sarcomas and correlation with morphologic and clinical features: A report of the CHAMP study group. , 1998, 22, 16-25.		161
137	Cytogenetic comparison of primary tumors and lymph node metastases in breast cancer patients. , 1998, 22, 122-129.		50
138	FISH characterization of head and neck carcinomas reveals that amplification of band 11q13 is associated with deletion of distal 11q. Genes Chromosomes and Cancer, 1998, 22, 312-320.	1.5	60
139	Frequent rearrangements of chromosomes 1, 7, and 8 in primary liver cancer. Genes Chromosomes and Cancer, 1998, 23, 26-35.	1.5	66
140	Cytogenetic analysis of pancreatic carcinomas: Intratumor heterogeneity and nonrandom pattern of chromosome aberrations. , 1998, 23, 81-99.		98
141	Chromosomal organization of amplified chromosome 12 sequences in mesenchymal tumors detected by fluorescence in situ hybridization. , 1998, 23, 203-212.		71
142	Cytogenetic comparison of primary tumors and lymph node metastases in breast cancer patients. , 1998, 22, 122.		1
143	Cytogenetic comparison of primary tumors and lymph node metastases in breast cancer patients. Genes Chromosomes and Cancer, 1998, 22, 122-129.	1.5	1
144	Frequent rearrangements of chromosomes 1, 7, and 8 in primary liver cancer. Genes Chromosomes and Cancer, 1998, 23, 26-35.	1.5	30

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#	Article	IF	CITATIONS
145	Cytogenetic comparisons of synchronous carcinomas and polyps in patients with colorectal cancer. British Journal of Cancer, 1997, 76, 765-769.	2.9	37
146	Characterization and Chromosomal Mapping of the HumanTFGGene Involved in Thyroid Carcinoma. Genomics, 1997, 41, 327-331.	1.3	42
147	Deletion of chromosome arm 3p in hematologic malignancies. Leukemia, 1997, 11, 1207-1213.	3.3	54
148	Characteristic sequence motifs at the breakpoints of the hybrid genes FUS/CHOP, EWS/CHOP and FUS/ERG in myxoid liposarcoma and acute myeloid leukemia. Oncogene, 1997, 15, 1357-1362.	2.6	65
149	A breakpoint map of recurrent chromosomal rearrangements in human neoplasia. Nature Genetics, 1997, 15, 417-474.	9.4	674
150	Clinical significance of cytogenetic findings in solid tumors. Cancer Genetics and Cytogenetics, 1997, 95, 1-8.	1.0	131
151	A subgroup of breast carcinomas is cytogenetically characterized by trisomy 12. Cancer Genetics and Cytogenetics, 1997, 97, 119-121.	1.0	10
152	Translocations between the Long Arms of Chromosomes 1 and 5 in Hematologic Malignancies Are Strongly Associated with Neoplasms of the Myeloid Lineages. Cancer Genetics and Cytogenetics, 1997, 99, 97-101.	1.0	17
153	Clonal CD5-positive B lymphocytes in myelodysplastic syndrome with systemic vasculitis and trisomy 8. Annals of Hematology, 1997, 74, 37-40.	0.8	16
154	Childhood acute lymphoblastic leukaemia with ider(21)(q10)t(12;21)(p12;q22): a new recurrent abnormality showing ETV6/CBFA2 fusion. British Journal of Haematology, 1997, 98, 216-218.	1.2	13
155	Deletion of the short arm of chromosome 3 in breast tumors. Genes Chromosomes and Cancer, 1997, 18, 241-245.	1.5	41
156	Deletions of CDKN1B and ETV6 in acute myeloid leukemia and myelodysplastic syndromes without cytogenetic evidence of 12p abnormalities. , 1997, 19, 77-83.		34
157	VariableFHIT transcripts in non-neoplastic tissues. , 1997, 19, 215-219.		38
158	BCR/ABL-negative chronic myeloid leukemia withETV6/ABL fusion. Genes Chromosomes and Cancer, 1997, 20, 299-304.	1.5	88
159	Long-term survival of patients with acute myeloid leukemia. Cancer, 1997, 80, 2191-2198.	2.0	110
160	Karyotypic abnormalities in fibroadenomas of the breast. , 1997, 70, 282-286.		27
161	Cytogenetic analysis of inverted nasal papillomas and demonstration of genetic convergence during in vitro passaging. International Journal of Cancer, 1997, 70, 668-673.	2.3	6
162	Cytogenetic findings in metastases from colorectal cancer. , 1997, 72, 604-607.		19

#	Article	IF	CITATIONS
163	Poor survival in t(8;21)(q22;q22)â€associated acute myeloid leukaemia with leukocytosis. European Journal of Haematology, 1997, 59, 47-52.	1.1	39
164	Longâ€ŧerm survival of patients with acute myeloid leukemia. Cancer, 1997, 80, 2191-2198.	2.0	6
165	Deletion of the short arm of chromosome 3 in breast tumors. , 1997, 18, 241.		1
166	BCR/ABL-negative chronic myeloid leukemia with ETV6/ABL fusion. , 1997, 20, 299.		3
167	Cytogenetic Analysis of Subcutaneous Angiolipoma: Further Evidence Supporting Its Difference from Ordinary Pure Lipomas. American Journal of Surgical Pathology, 1997, 21, 441-444.	2.1	86
168	Cytogenetic Abnormalities and Clonal Evolution in an Adult Hepatoblastoma. American Journal of Surgical Pathology, 1997, 21, 1381-1386.	2.1	38
169	Expression Patterns of the Human Sarcoma-Associated GenesFUSandEWSand the Genomic Structure ofFUS. Genomics, 1996, 37, 1-8.	1.3	144
170	Chromosome rearrangements in synovial chondromatous lesions. British Journal of Cancer, 1996, 74, 251-254.	2.9	61
171	Cytogenetic findings in four malignant mixed mesodermal tumors of the ovary. Cancer Genetics and Cytogenetics, 1996, 88, 53-56.	1.0	10
172	No FISH evidence for trisomy 7 in normal or leukemic bone Marrow. Cancer Genetics and Cytogenetics, 1996, 88, 133-135.	1.0	3
173	Ring marker containing 17q and chromosome 22 in a case of dermatofibrosarcoma protuberans. Cancer Genetics and Cytogenetics, 1996, 89, 88-91.	1.0	28
174	Evidence of somatic mutations in osteoarthritis. Human Genetics, 1996, 98, 651-656.	1.8	30
175	Cytogenetic abnormalities in anin situ ductal carcinoma and five prophylactically removed breasts from members of a family with hereditary breast cancer. Breast Cancer Research and Treatment, 1996, 38, 177-182.	1.1	29
176	Dichotomy of hyperdiploid acute lymphoblastic leukemia on the basis of the distribution of gained chromosomes. Cancer Genetics and Cytogenetics, 1996, 92, 8-10.	1.0	31
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