Ioannis Panagopoulos

List of Publications by Year
in descending order

143	4,991	76196	110170
papers	citations	40 h-index	64 g-index

144	144	144	4881
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Consistent Rearrangement of Chromosomal Band 6p21 with Generation of Fusion Genes JAZF1/PHF1 and EPC1/PHF1 in Endometrial Stromal Sarcoma. Cancer Research, 2006, 66, 107-112.	0.4	225
2	The chimericFUS/CREB3l2gene is specific for low-grade fibromyxoid sarcoma. Genes Chromosomes and Cancer, 2004, 40, 218-228.	1.5	188
3	Confirmation of the high frequency of theTMPRSS2/ERG fusion gene in prostate cancer. Genes Chromosomes and Cancer, 2006, 45, 717-719.	1.5	182
4	Molecular genetic characterization of theEWS/ATF1fusion gene in clear cell sarcoma of tendons and aponeuroses. International Journal of Cancer, 2002, 99, 560-567.	2.3	135
5	Activation of the GLI Oncogene through Fusion with the Î²-Actin Gene (ACTB) in a Group of Distinctive Pericytic Neoplasms. American Journal of Pathology, 2004, 164, 1645-1653.	1.9	130
6	ZC3H7B-BCOR high-grade endometrial stromal sarcomas: a report of 17 cases of a newly defined entity. Modern Pathology, 2018, 31, 674-684.	2.9	130
7	t(19;22)(q13;q12) Translocation leading to the novel fusion gene EWSR1â€ZNF444 in soft tissue myoepithelial carcinoma. Genes Chromosomes and Cancer, 2009, 48, 1051-1056.	1.5	125
8	Detection of a t(1;22)(q23;q12) translocation leading to an EWSR1â€PBX1 fusion gene in a myoepithelioma. Genes Chromosomes and Cancer, 2008, 47, 558-564.	1.5	123
9	Fusion of theFUS gene withERG in acute myeloid leukemia with t(16;21)(p11;q22). Genes Chromosomes and Cancer, 1994, 11, 256-262.	1.5	116
10	Fusion of the ZC3H7B and BCOR genes in endometrial stromal sarcomas carrying an X;22â€translocation. Genes Chromosomes and Cancer, 2013, 52, 610-618.	1.5	114
11	Fusion of the SEC31L1 and ALK genes in an inflammatory myofibroblastic tumor. International Journal of Cancer, 2006, 118, 1181-1186.	2.3	113
12	Clinical impact of molecular and cytogenetic findings in synovial sarcoma. Genes Chromosomes and Cancer, 2001, 31, 362-372.	1.5	108
13	Molecular genetic characterization of theEWS/CHNandRBP56/CHNfusion genes in extraskeletal myxoid chondrosarcoma. Genes Chromosomes and Cancer, 2002, 35, 340-352.	1.5	104
14	Lentivirus Vector Gene Expression during ES Cell-Derived Hematopoietic Development In Vitro. Journal of Virology, 2000, 74, 10778-10784.	1.5	100
15	Novel Fusion of MYST/Esa1-Associated Factor 6 and PHF1 in Endometrial Stromal Sarcoma. PLoS ONE, 2012, 7, e39354.	1.1	94
16	Cytogenetic and molecular genetic analyses of endometrial stromal sarcoma: nonrandom involvement of chromosome arms 6p and 7p and confirmation of JAZF1/JJAZ1 gene fusion in t(7;17). Cancer Genetics and Cytogenetics, 2003, 144, 119-124.	1.0	92
17	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
18	Fusion, Disruption, and Expression of HMGA2 in Bone and Soft Tissue Chondromas. Modern Pathology, 2003, 16, 1132-1140.	2.9	82

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19	Absence of mutations of the BRAF gene in malignant melanoma of soft parts (clear cell sarcoma of) Tj ETQq1 1 0.784314 rgBT /Overlock	1.0	82
20	MEAF6/PHF1 is a recurrent gene fusion in endometrial stromal sarcoma. Cancer Letters, 2014, 347, 75-78.	3.2	79
21	Whole-Transcriptome Sequencing Identifies Novel IRF2BP2-CDX1 Fusion Gene Brought about by Translocation t(1;5)(q42;q32) in Mesenchymal Chondrosarcoma. PLoS ONE, 2012, 7, e49705.	1.1	77
22	Assessment of the clinical and molecular impact of different cytogenetic subgroups in a series of 272 lipomas with abnormal karyotype. Genes Chromosomes and Cancer, 2007, 46, 594-606.	1.5	75
23	Fusion of the RBP56 and CHN genes in extraskeletal myxoid chondrosarcomas with translocation t(9;17)(q22;q11). Oncogene, 1999, 18, 7594-7598.	2.6	69
24	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
25	Identification of genes differentially expressed in TLS-CHOP carrying myxoid liposarcomas. , 1999, 83, 30-33.		64
26	RNA sequencing identifies fusion of the EWSR1 and YY1 genes in mesothelioma with t(14;22)(q32;q12). Genes Chromosomes and Cancer, 2013, 52, 733-740.	1.5	64
27	Characterization of the native CREB3L2 transcription factor and the FUS/CREB3L2 chimera. Genes Chromosomes and Cancer, 2007, 46, 181-191.	1.5	54
28	Cytogenetic and molecular profile of endometrial stromal sarcoma. Genes Chromosomes and Cancer, 2016, 55, 834-846.	1.5	54
29	Rearrangement of theCOL12A1andCOL4A5genes in subungual exostosis: molecular cytogenetic delineation of the tumor-specific translocationt(X;6)(q13-14;q22). International Journal of Cancer, 2006, 118, 1972-1976.	2.3	53
30	The â€œGrepâ€•Command But Not FusionMap, FusionFinder or ChimeraScan Captures the CIC-DUX4 Fusion Gene from Whole Transcriptome Sequencing Data on a Small Round Cell Tumor with t(4;19)(q35;q13). PLoS ONE, 2014, 9, e99439.	1.1	53
31	Cloning and sequencing of a cDNA encoding ratd-dopachrome tautomerase. FEBS Letters, 1995, 373, 203-206.	1.3	52
32	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
33	The POU5F1P1 pseudogene encodes a putative protein similar to POU5F1 isoform 1. Oncology Reports, 2008, 20, 1029-33.	1.2	50
34	Fusion of NUP98 and the SET binding protein 1 (SETBP1) gene in a paediatric acute T cell lymphoblastic leukaemia with t(11;18)(p15;q12). British Journal of Haematology, 2007, 136, 294-296.	1.2	49
35	Molecular genetic characterization of the genomic ACTB-GLI fusion in pericytoma with t(7;12). Biochemical and Biophysical Research Communications, 2004, 325, 1318-1323.	1.0	46
36	Fusion of the genes BRD8 and PHF1 in endometrial stromal sarcoma. Genes Chromosomes and Cancer, 2017, 56, 841-845.	1.5	45

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37	Heterogeneous genetic profiles in soft tissue myoepitheliomas. Modern Pathology, 2008, 21, 1311-1319.	2.9	44
38	Fusion genes withALKas recurrent partner in ependymoma-like gliomas: a new brain tumor entity?. Neuro-Oncology, 2015, 17, 1365-1373.	0.6	44
39	MGMT Gene Promoter Methylation Status â€“ Assessment of Two Pyrosequencing Kits and Three Methylation-specific PCR Methods for their Predictive Capacity in Glioblastomas. Cancer Genomics and Proteomics, 2018, 15, 437-446.	1.0	44
40	Chromosome aberrations and HEY1-NCOA2 fusion gene in a mesenchymal chondrosarcoma. Oncology Reports, 2014, 32, 40-44.	1.2	43
41	Novel CSF1-S100A10 fusion gene and CSF1 transcript identified by RNA sequencing in tenosynovial giant cell tumors. International Journal of Oncology, 2014, 44, 1425-1432.	1.4	42
42	Fusion of the Genes WWTR1 and FOSB in Pseudomyogenic Hemangioendothelioma. Cancer Genomics and Proteomics, 2019, 16, 293-298.	1.0	41
43	Fusion of ZMYND8 and RELA Genes in Acute Erythroid Leukemia. PLoS ONE, 2013, 8, e63663.	1.1	40
44	A methylation PCR approach for detection of fragile X syndrome. , 1999, 14, 71-79.		39
45	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
46	VariableFHIT transcripts in non-neoplastic tissues. , 1997, 19, 215-219.		38
47	An endometrial stromal sarcoma cell line with the JAZF1/PHF1 chimera. Cancer Genetics and Cytogenetics, 2008, 185, 74-77.	1.0	37
48	NoEWS/FLI1 fusion transcripts in giant-cell tumors of bone. International Journal of Cancer, 2001, 93, 769-772.	2.3	35
49	Expression levels of HMGA2 in adipocytic tumors correlate with morphologic and cytogenetic subgroups. Molecular Cancer, 2009, 8, 36.	7.9	35
50	Novel KAT6B-KANSL1 Fusion Gene Identified by RNA Sequencing in Retroperitoneal Leiomyoma with t(10;17)(q22;q21). PLoS ONE, 2015, 10, e0117010.	1.1	34
51	Identification of anEPC2-PHF1fusion transcript in low-grade endometrial stromal sarcoma. Oncotarget, 2018, 9, 19203-19208.	0.8	32
52	18F-Fluciclovine PET/CT in Suspected Residual or Recurrent High-Grade Glioma. Clinical Nuclear Medicine, 2019, 44, 605-611.	0.7	30
53	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
54	NDRG1-PLAG1 and TRPS1-PLAG1 Fusion Genes in Chondroid Syringoma. Cancer Genomics and Proteomics, 2020, 17, 237-248.	1.0	29

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55	Recurrent fusion of the genes FN1 and ALK in gastrointestinal leiomyomas. Modern Pathology, 2016, 29, 1415-1423.	2.9	28
56	Genomic PCR detects tumor cells in peripheral blood from patients with myxoid liposarcoma. , 1996, 17, 102-107.		25
57	Rearrangement of chromosome bands 12q14–15 causing HMGA2-SOX5 gene fusion and HMGA2 expression in extraskeletal osteochondroma. Oncology Reports, 2015, 34, 577-584.	1.2	25
58	Molecular pathogenesis and prognostication of "lowâ€grade'' and "highâ€grade" endometrial stromal sarcoma. Genes Chromosomes and Cancer, 2021, 60, 160-167.	1.5	25
59	Fusion of ETV6 with an intronic sequence of the BAZ2A gene in a paediatric pre-B acute lymphoblastic leukaemia with a cryptic chromosome 12 rearrangement. British Journal of Haematology, 2006, 133, 270-275.	1.2	24
60	Fusion of the COL1A1 and USP6 genes in a benign bone tumor. Cancer Genetics and Cytogenetics, 2008, 180, 70-73.	1.0	24
61	RNAâ€sequencing identifies novel GREB1â€NCOA2 fusion gene in a uterine sarcoma with the chromosomal translocation t(2;8)(p25;q13). Genes Chromosomes and Cancer, 2018, 57, 176-181.	1.5	24
62	Fusion of the Genes PHF1 and TFE3 in Malignant Chondroid Syringoma. Cancer Genomics and Proteomics, 2019, 16, 345-351.	1.0	21
63	LAMTOR1-PRKCD and NUMA1-SFMBT1 fusion genes identified by RNA sequencing in aneurysmal benign fibrous histiocytoma with t(3;11)(p21;q13). Cancer Genetics, 2015, 208, 545-551.	0.2	20
64	Gene fusions AHRR-NCOA2, NCOA2-ETV4, ETV4-AHRR, P4HA2-TBCK, and TBCK-P4HA2 resulting from the translocations t(5;8;17)(p15;q13;q21) and t(4;5)(q24;q31) in a soft tissue angiofibroma. Oncology Reports, 2016, 36, 2455-2462.	1.2	20
65	Acute Myeloid Leukemia with inv(8)(p11q13). Leukemia and Lymphoma, 2000, 39, 651-656.	0.6	19
66	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
67	Fusion of the HMGA2 and C9orf92 genes in myolipoma with t(9;12)(p22;q14). Diagnostic Pathology, 2016, 11, 22.	0.9	18
68	The microRNA miR-192/215 family is upregulated in mucinous ovarian carcinomas. Scientific Reports, 2018, 8, 11069.	1.6	18
69	A Small-Molecule Tankyrase Inhibitor Reduces Glioma Stem Cell Proliferation and Sphere Formation. Cancers, 2020, 12, 1630.	1.7	18
70	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
71	A novel TCF3-HLF fusion transcript in acute lymphoblastic leukemia with a t(17;19)(q22;p13). Cancer Genetics, 2012, 205, 669-672.	0.2	17
72	Cryptic FUS-ERG fusion identified by RNA-sequencing in childhood acute myeloid leukemia. Oncology Reports, 2013, 30, 2587-2592.	1.2	17

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73	HMGA2 expression pattern and TERT mutations in tumors of the vulva. Oncology Reports, 2015, 33, 2675-2680.	1.2	17
74	Novel TNS3-MAP3K3 and ZFPM2-ELF5 fusion genes identified by RNA sequencing in multicystic mesothelioma with t(7;17)(p12;q23) and t(8;11)(q23;p13). Cancer Letters, 2015, 357, 502-509.	3.2	17
75	Genetic heterogeneity in leiomyomas of deep soft tissue. Oncotarget, 2017, 8, 48769-48781.	0.8	17
76	Myeloid leukemia with t(7;21)(p22;q22) and 5q deletion. Oncology Reports, 2013, 30, 1549-1552.	1.2	16
77	Sequential combination of karyotyping and RNA-sequencing in the search for cancer-specific fusion genes. International Journal of Biochemistry and Cell Biology, 2014, 53, 462-465.	1.2	16
78	Loss of chromosome 13 material in cellular angiofibromas indicates pathogenetic similarity with spindle cell lipomas. Diagnostic Pathology, 2017, 12, 17.	0.9	16
79	Genetic Characterization of Myoid Hamartoma of the Breast. Cancer Genomics and Proteomics, 2019, 16, 563-568.	1.0	16
80	Expressions of miR-30c and let-7a are inversely correlated with HMGA2 expression in squamous cell carcinoma of the vulva. Oncotarget, 2016, 7, 85058-85062.	0.8	16
81	Fusion of the genes ataxin 2 like, ATXN2L, and Janus kinase 2, JAK2, in cutaneous CD4 positive T-cell lymphoma. Oncotarget, 2017, 8, 103775-103784.	0.8	16
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	t(3;21)(q22;q22) leading to truncation of the RYK gene in atypical chronic myeloid leukemia. Cancer Letters, 2009, 277, 205-211.	3.2	15
84	Comparison between Karyotyping-FISH-Reverse Transcription PCR and RNA- Sequencing-Fusion Gene Identification Programs in the Detection of KAT6A-CREBBP in Acute Myeloid Leukemia. PLoS ONE, 2014, 9, e96570.	1.1	15
85	Novel ZEB2-BCL11B Fusion Gene Identified by RNA-Sequencing in Acute Myeloid Leukemia with t(2;14)(q22;q32). PLoS ONE, 2015, 10, e0132736.	1.1	15
86	Fusion of the Genes EWSR1 and PBX3 in Retroperitoneal Leiomyoma with t(9;22)(q33;q12). PLoS ONE, 2015, 10, e0124288.	1.1	15
87	Genomic imbalances are involved in miR-30c and let-7a deregulation in ovarian tumors: implications for HMGA2 expression. Oncotarget, 2017, 8, 21554-21560.	0.8	15
88	FAM53B truncation caused by t(10;19)(q26;q13) chromosome translocation in acute lymphoblastic leukemia. Oncology Letters, 2017, 13, 2216-2220.	0.8	14
89	Expression and clinical role of the dipeptidyl peptidases DPP8 and DPP9 in ovarian carcinoma. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 2019, 474, 177-185.	1.4	14
90	Fusion of RDC1 with HMGA2 in lipomas as the result of chromosome aberrations involving 2q35-37 and 12q13-15. International Journal of Oncology, 2002, 21, 321-6.	1.4	14

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91	Low Frequency of ESRRAâ€“C11orf20 Fusion Gene in Ovarian Carcinomas. PLoS Biology, 2014, 12, e1001784.	2.6	13
92	The recurrent chromosomal translocation t(12;18) (q14~15;q12~21) causes the fusion gene HMGA2-SETBP1 and HMGA2 expression in lipoma and osteochondrolipoma. International Journal of Oncology, 2015, 47, 884-890.	1.4	13
93	Recurrent Fusion of the Genes for High-mobility Group AT-hook 2 (HMGA2) and Nuclear Receptor Co-repressor 2 (NCOR2) in Osteoclastic Giant Cell-rich Tumors of Bone. Cancer Genomics and Proteomics, 2022, 19, 163-177.	1.0	13
94	Several fusion genes identified by whole transcriptome sequencing in a spindle cell sarcoma with rearrangements of chromosome arm 12q and MDM2 amplification. International Journal of Oncology, 2014, 45, 1829-1836.	1.4	12
95	K27/G34 versus K28/G35 in histone H3-mutant gliomas: A note of caution. Acta Neuropathologica, 2018, 136, 175-176.	3.9	12
96	Novel GTF2Iâ€“PDGFRB and IKZF1â€“TYW1 fusions in pediatric leukemia with normal karyotype. Experimental Hematology and Oncology, 2019, 8, 12.	2.0	12
97	Fusion of the Lumican (LUM) Gene With the Ubiquitin Specific Peptidase 6 (USP6) Gene in an Aneurysmal Bone Cyst Carrying a t(12;17)(q21;p13) Chromosome Translocation. Cancer Genomics and Proteomics, 2020, 17, 555-561.	1.0	12
98	An Unbalanced Chromosome Translocation Between 7p22 and 12q13 Leads to ACTB-GLI1 Fusion in Pericytoma. Anticancer Research, 2020, 40, 1239-1245.	0.5	12
99	Cytogenetics of Spindle Cell/Pleomorphic Lipomas: Karyotyping and FISH Analysis of 31 Tumors. Cancer Genomics and Proteomics, 2018, 15, 193-200.	1.0	12
100	A PCR/restriction digestion assay for the detection of the transcript variants 1 and 2 of POU5F1. Genes Chromosomes and Cancer, 2008, 47, 521-529.	1.5	11
101	Pyrosequencing Analysis of MGMT Promoter Methylation in Meningioma. Cancer Genomics and Proteomics, 2018, 15, 379-385.	1.0	11
102	Novel fusion genes and chimeric transcripts in ependymal tumors. Genes Chromosomes and Cancer, 2016, 55, 944-953.	1.5	10
103	Interstitial Deletions Generating Fusion Genes. Cancer Genomics and Proteomics, 2021, 18, 167-196.	1.0	10
104	Rearrangement of the Chromatin Organizer Special AT-rich Binding Protein 1 Gene, SATB1, Resulting from a t(3;5)(p24;q14) Chromosomal Translocation in Acute Myeloid Leukemia. Anticancer Research, 2017, 37, 693-698.	0.5	10
105	A novel PCR-based approach for the detection of the Huntington disease associated trinucleotide repeat expansion. , 1999, 13, 232-236.		8
106	Upregulation of INS-IGF2 read-through expression and identification of a novel INS-IGF2 splice variant in insulinomas. Oncology Reports, 2016, 36, 2653-2662.	1.2	8
107	Molecular characterization of the t(4;12)(q27~28;q14~15) chromosomal rearrangement in lipoma. Oncology Letters, 2016, 12, 1701-1704.	0.8	8
108	Identification of novel cyclin gene fusion transcripts in endometrioid ovarian carcinomas. International Journal of Cancer, 2018, 143, 1379-1387.	2.3	8

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109	PAN3â€“PSMA2 fusion resulting from a novel t(7;13)(p14;q12) chromosome translocation in a myelodysplastic syndrome that evolved into acute myeloid leukemia. Experimental Hematology and Oncology, 2018, 7, 7.	2.0	8
110	FOS-ANKH and FOS-RUNX2 Fusion Genes in Osteoblastoma. Cancer Genomics and Proteomics, 2020, 17, 161-168.	1.0	8
111	RUNX1 truncation resulting from a cryptic and novel t(6;21)(q25;q22) chromosome translocation in acute myeloid leukemia: A case report. Oncology Reports, 2016, 36, 2481-2488.	1.2	7
112	A novel truncated form of HMGA2 in tumors of the ovaries. Oncology Letters, 2016, 12, 1559-1563.	0.8	7
113	Karyotyping and analysis of GNAS locus in intramuscular myxomas. Oncotarget, 2017, 8, 22086-22094.	0.8	7
114	Therapy-induced Deletion in 11q23 Leading to Fusion ofKMT2AWithARHGEF12and Development of B Lineage Acute Lymphoplastic Leukemia in a Child Treated for Acute Myeloid Leukemia Caused by t(9;11)(p21;q23)/KMT2A-MLLT3. Cancer Genomics and Proteomics, 2021, 18, 67-81.	1.0	7
115	Cytogenetic Analysis of a Pseudoangiomatous Pleomorphic/Spindle Cell Lipoma. Anticancer Research, 2017, 37, 2119-2123.	0.5	7
116	Consistent Involvement of Chromosome 13 in Angiolipoma. Cancer Genomics and Proteomics, 2018, 15, 61-65.	1.0	7
117	Recurrent 12q13-15 chromosomal aberrations, high frequency of isocitrate dehydrogenase 1 mutations, and absence of high mobility group AT-hook 2 expression in periosteal chondromas. Oncology Letters, 2015, 10, 163-167.	0.8	5
118	Rare MLL-ELL fusion transcripts in childhood acute myeloid leukemiaâ€”association with young age and myeloid sarcomas?. Experimental Hematology and Oncology, 2015, 5, 8.	2.0	5
119	Fusion of the COL1A1 and FYN Genes in Epithelioid Osteoblastoma. Cancer Genomics and Proteomics, 2019, 16, 361-368.	1.0	5
120	Recurrent Fusion of the GRB2 Associated Binding Protein 1 (GAB1) Gene With ABL Proto-oncogene 1 (ABL1) in Benign Pediatric Soft Tissue Tumors. Cancer Genomics and Proteomics, 2020, 17, 499-508.	1.0	5
121	Cytogenetic and molecular analyses of 291 gastrointestinal stromal tumors: site-specific cytogenetic evolution as evidence of pathogenetic heterogeneity. Oncotarget, 2022, 13, 508-517.	0.8	5
122	Therapy-related Myeloid Leukemia With the Translocation t(8;19)(p11;q13) Leading to a KAT6A-LEUTX Fusion Gene. Anticancer Research, 2021, 41, 1753-1760.	0.5	4
123	Recurrent fusion transcripts in squamous cell carcinomas of the vulva. Oncotarget, 2017, 8, 16843-16850.	0.8	4
124	DEK-NUP214-Fusion Identified by RNA-Sequencing of an Acute Myeloid Leukemia with t(9;12)(q34;q15). Cancer Genomics and Proteomics, 2017, 14, 437-443.	1.0	4
125	Fusion of the TBL1XR1 and HMGA1 genes in splenic hemangioma with t(3;6)(q26;p21). International Journal of Oncology, 2016, 48, 1242-1250.	1.4	3
126	Chromosome Translocation t(14;21)(q11;q22) Activates Both OLIG1 and OLIG2 in Pediatric T-cell Lymphoblastic Malignancies and May Signify Adverse Prognosis. Cancer Genomics and Proteomics, 2020, 17, 41-48.	1.0	3

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127	Cytogenetic and Molecular Study of an Adult Sclerosing Rhabdomyosarcoma of the Extremity: MYOD1-mutation and Clonal Evolution. Cancer Genomics and Proteomics, 2020, 17, 563-569.	1.0	3
128	RareKMT2A-ELLand NovelZNF56-KMT2AFusion Genes in Pediatric T-cell Acute Lymphoblastic Leukemia. Cancer Genomics and Proteomics, 2021, 18, 121-131.	1.0	3
129	Mutation analysis and genomic imbalances of cells found in effusion fluids from patients with ovarian cancer. Oncology Letters, 2020, 20, 2273-2279.	0.8	3
130	Characterization of the human CREB3L2 gene promoter. Oncology Reports, 2009, 21, 615-24.	1.2	3
131	Identification of SETD2-NF1 fusion gene in a pediatric spindle cell tumor with the chromosomal translocation t(3;17)(p21;q12). Oncology Reports, 2017, 37, 3181-3188.	1.2	2
132	RUNX1-PDCD6 fusion resulting from a novel t(5;21)(p15;q22) chromosome translocation in myelodysplastic syndrome secondary to chronic lymphocytic leukemia. PLoS ONE, 2018, 13, e0196181.	1.1	2
133	MGMT promoter methylation is a rare epigenetic change in malignant effusions. Cytopathology, 2020, 31, 12-15.	0.4	2
134	Fusion of the COL4A5 Gene With NR2F2-AS1 in a Hemangioma Carrying a t(X;15)(q22;q26) Chromosomal Translocation. Cancer Genomics and Proteomics, 2020, 17, 383-390.	1.0	2
135	TYRO3 Truncation Resulting From a t(10;15)(p11;q15) Chromosomal Translocation in Pediatric Acute Myeloid Leukemia. Anticancer Research, 2020, 40, 6115-6121.	0.5	2
136	Expression of DOL54 is not restricted to myxoid liposarcomas with the FUS-DDIT3 chimera but is found in various sarcomas. Oncology Reports, 2004, 12, 107-10.	1.2	2
137	Molecular Genetic Characterization of Acute Myeloid Leukemia With Trisomy 4 as the Sole Chromosome Abnormality. Cancer Genomics and Proteomics, 2019, 16, 175-178.	1.0	1
138	Several Fusion Genes Identified in a Spermatic Cord Leiomyoma With Rearrangements of Chromosome Arms 3p and 21q. Cancer Genomics and Proteomics, 2021, 18, 531-542.	1.0	1
139	Characterization of an alternative transcript of the human CREB3L2 gene. Oncology Reports, 2010, 24, 1133-9.	1.2	0
140	Chronic Expanding Hematoma with a t(11;19)(q13;q13) Chromosomal Translocation. Anticancer Research, 2020, 40, 97-100.	0.5	0
141	Chromosomal complexity as a biomarker to de-escalate adjuvant imatinib treatment in high-risk gastrointestinal stromal tumor.. Journal of Clinical Oncology, 2021, 39, 11535-11535.	0.8	0
142	Monosomy 13 in Mammary Myofibroblastoma. Anticancer Research, 2021, 41, 3747-3751.	0.5	0
143	Fusion of the Paired Box 3 (PAX3) and Myocardin (MYOCD) Genes in Pediatric Rhabdomyosarcoma. Cancer Genomics and Proteomics, 2021, 18, 723-734.	1.0	0