# 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 | 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 |
| 2 | 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 |
| 3 | Molecular pathogenesis and prognostication of "lowâ€grade" and "highâ€grade" endometrial stromal sarcoma. Genes Chromosomes and Cancer, 2021, 60, 160-167. | 1.5 | 25 |
| 4 | 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 |
| 5 | 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 |
| 6 | Interstitial Deletions Generating Fusion Genes. Cancer Genomics and Proteomics, 2021, 18, 167-196. | 1.0 | 10 |
| 7 | RareKMT2A-ELLand NovelZNF56-KMT2AFusion Genes in Pediatric T-cell Acute Lymphoblastic Leukemia. Cancer Genomics and Proteomics, 2021, 18, 121-131. | 1.0 | 3 |
| 8 | 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 |
| 9 | 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 |
| 10 | Monosomy 13 in Mammary Myofibroblastoma. Anticancer Research, 2021, 41, 3747-3751. | 0.5 | 0 |
| 11 | 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 |
| 12 | Chronic Expanding Hematoma with a t(11;19)(q13;q13) Chromosomal Translocation. Anticancer Research, 2020, 40, 97-100. | 0.5 | 0 |
| 13 | MGMT promoter methylation is a rare epigenetic change in malignant effusions. Cytopathology, 2020, 31, 12-15. | 0.4 | 2 |
| 14 | 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 |
| 15 | 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 |
| 16 | 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 |
| 17 | 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 |
| 18 | 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 |

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|---|---|---|---|
| 19 | An Unbalanced Chromosome Translocation Between 7p22 and 12q13 Leads to ACTB-GLI1 Fusion in Pericytoma. Anticancer Research, 2020, 40, 1239-1245. | 0.5 | 12 |
| 20 | A Small-Molecule Tankyrase Inhibitor Reduces Glioma Stem Cell Proliferation and Sphere Formation. Cancers, 2020, 12, 1630. | 1.7 | 18 |
| 21 | FOS-ANKH and FOS-RUNX2 Fusion Genes in Osteoblastoma. Cancer Genomics and Proteomics, 2020, 17, 161-168. | 1.0 | 8 |
| 22 | NDRG1-PLAG1 and TRPS1-PLAG1 Fusion Genes in Chondroid Syringoma. Cancer Genomics and Proteomics, 2020, 17, 237-248. | 1.0 | 29 |
| 23 | 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 |
| 24 | 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 |
| 25 | Fusion of the Genes WWTR1 and FOSB in Pseudomyogenic Hemangioendothelioma. Cancer Genomics and Proteomics, 2019, 16, 293-298. | 1.0 | 41 |
| 26 | Genetic Characterization of Myoid Hamartoma of the Breast. Cancer Genomics and Proteomics, 2019, 16, 563-568. | 1.0 | 16 |
| 27 | Fusion of the Genes PHF1 and TFE3 in Malignant Chondroid Syringoma. Cancer Genomics and Proteomics, 2019, 16, 345-351. | 1.0 | 21 |
| 28 | Fusion of the COL1A1 and FYN Genes in Epithelioid Osteoblastoma. Cancer Genomics and Proteomics, 2019, 16, 361-368. | 1.0 | 5 |
| 29 | Novel GTF2Iâ€“PDGFRB and IKZF1â€“TYW1 fusions in pediatric leukemia with normal karyotype. Experimental Hematology and Oncology, 2019, 8, 12. | 2.0 | 12 |
| 30 | 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 |
| 31 | 18F-Fluciclovine PET/CT in Suspected Residual or Recurrent High-Grade Glioma. Clinical Nuclear Medicine, 2019, 44, 605-611. | 0.7 | 30 |
| 32 | 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 |
| 33 | Identification of novel cyclin gene fusion transcripts in endometrioid ovarian carcinomas. International Journal of Cancer, 2018, 143, 1379-1387. | 2.3 | 8 |
| 34 | 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 |
| 35 | 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 |
| 36 | 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 |

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|---|---|---|---|
| 37 | 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 |
| 38 | Pyrosequencing Analysis of MGMT Promoter Methylation in Meningioma. Cancer Genomics and Proteomics, 2018, 15, 379-385. | 1.0 | 11 |
| 39 | K27/G34 versus K28/G35 in histone H3-mutant gliomas: A note of caution. Acta Neuropathologica, 2018, 136, 175-176. | 3.9 | 12 |
| 40 | The microRNA miR-192/215 family is upregulated in mucinous ovarian carcinomas. Scientific Reports, 2018, 8, 11069. | 1.6 | 18 |
| 41 | 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 |
| 42 | Identification of anEPC2-PHF1fusion transcript in low-grade endometrial stromal sarcoma. Oncotarget, 2018, 9, 19203-19208. | 0.8 | 32 |
| 43 | Consistent Involvement of Chromosome 13 in Angiolipoma. Cancer Genomics and Proteomics, 2018, 15, 61-65. | 1.0 | 7 |
| 44 | Cytogenetics of Spindle Cell/Pleomorphic Lipomas: Karyotyping and FISH Analysis of 31 Tumors. Cancer Genomics and Proteomics, 2018, 15, 193-200. | 1.0 | 12 |
| 45 | FAM53B truncation caused by t(10;19)(q26;q13) chromosome translocation in acute lymphoblastic leukemia. Oncology Letters, 2017, 13, 2216-2220. | 0.8 | 14 |
| 46 | 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 |
| 47 | Loss of chromosome 13 material in cellular angiofibromas indicates pathogenetic similarity with spindle cell lipomas. Diagnostic Pathology, 2017, 12, 17. | 0.9 | 16 |
| 48 | Fusion of the genes BRD8 and PHF1 in endometrial stromal sarcoma. Genes Chromosomes and Cancer, 2017, 56, 841-845. | 1.5 | 45 |
| 49 | Karyotyping and analysis of GNAS locus in intramuscular myxomas. Oncotarget, 2017, 8, 22086-22094. | 0.8 | 7 |
| 50 | Recurrent fusion transcripts in squamous cell carcinomas of the vulva. Oncotarget, 2017, 8, 16843-16850. | 0.8 | 4 |
| 51 | 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 |
| 52 | Genetic heterogeneity in leiomyomas of deep soft tissue. Oncotarget, 2017, 8, 48769-48781. | 0.8 | 17 |
| 53 | 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 |
| 54 | 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 |

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|---|---|---|---|
| 55 | Cytogenetic Analysis of a Pseudoangiomatous Pleomorphic/Spindle Cell Lipoma. Anticancer Research, 2017, 37, 2119-2123. | 0.5 | 7 |
| 56 | 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 |
| 57 | Cytogenetic and molecular profile of endometrial stromal sarcoma. Genes Chromosomes and Cancer, 2016, 55, 834-846. | 1.5 | 54 |
| 58 | 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 |
| 59 | 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 |
| 60 | 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 |
| 61 | Recurrent fusion of the genes FN1 and ALK in gastrointestinal leiomyomas. Modern Pathology, 2016, 29, 1415-1423. | 2.9 | 28 |
| 62 | 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 |
| 63 | A novel truncated form of HMGA2 in tumors of the ovaries. Oncology Letters, 2016, 12, 1559-1563. | 0.8 | 7 |
| 64 | Novel fusion genes and chimeric transcripts in ependymal tumors. Genes Chromosomes and Cancer, 2016, 55, 944-953. | 1.5 | 10 |
| 65 | Molecular characterization of the t(4;12)(q27~28;q14~15) chromosomal rearrangement in lipoma. Oncology Letters, 2016, 12, 1701-1704. | 0.8 | 8 |
| 66 | Fusion of the HMGA2 and C9orf92 genes in myolipoma with t(9;12)(p22;q14). Diagnostic Pathology, 2016, 11, 22. | 0.9 | 18 |
| 67 | 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 |
| 68 | HMGA2 expression pattern and TERT mutations in tumors of the vulva. Oncology Reports, 2015, 33, 2675-2680. | 1.2 | 17 |
| 69 | 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 |
| 70 | 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 |
| 71 | 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 |
| 72 | 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 |

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|---|---|---|---|
| 73 | 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 |
| 74 | Fusion genes withALKas recurrent partner in ependymoma-like gliomas: a new brain tumor entity?. Neuro-Oncology, 2015, 17, 1365-1373. | 0.6 | 44 |
| 75 | 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 |
| 76 | 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 |
| 77 | 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 |
| 78 | Fusion of the Genes EWSR1 and PBX3 in Retroperitoneal Leiomyoma with t(9;22)(q33;q12). PLoS ONE, 2015, 10, e0124288. | 1.1 | 15 |
| 79 | 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 |
| 80 | 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 |
| 81 | Low Frequency of ESRRAâ€“C11orf20 Fusion Gene in Ovarian Carcinomas. PLoS Biology, 2014, 12, e1001784. | 2.6 | 13 |
| 82 | Chromosome aberrations and HEY1-NCOA2 fusion gene in a mesenchymal chondrosarcoma. Oncology Reports, 2014, 32, 40-44. | 1.2 | 43 |
| 83 | MEAF6/PHF1 is a recurrent gene fusion in endometrial stromal sarcoma. Cancer Letters, 2014, 347, 75-78. | 3.2 | 79 |
| 84 | 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 |
| 85 | 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 |
| 86 | 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 |
| 87 | Cryptic FUS-ERG fusion identified by RNA-sequencing in childhood acute myeloid leukemia. Oncology Reports, 2013, 30, 2587-2592. | 1.2 | 17 |
| 88 | 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 |
| 89 | 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 |
| 90 | Myeloid leukemia with t(7;21)(p22;q22) and 5q deletion. Oncology Reports, 2013, 30, 1549-1552. | 1.2 | 16 |

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| 91 | Fusion of ZMYND8 and RELA Genes in Acute Erythroid Leukemia. PLoS ONE, 2013, 8, e63663. | 1.1 | 40 |
| 92 | 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 |
| 93 | 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 |
| 94 | Novel Fusion of MYST/Esa1-Associated Factor 6 and PHF1 in Endometrial Stromal Sarcoma. PLoS ONE, 2012, 7, e39354. | 1.1 | 94 |
| 95 | Characterization of an alternative transcript of the human CREB3L2 gene. Oncology Reports, 2010, 24, 1133-9. | 1.2 | 0 |
| 96 | 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 |
| 97 | 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 |
| 98 | Expression levels of HMGA2 in adipocytic tumors correlate with morphologic and cytogenetic subgroups. Molecular Cancer, 2009, 8, 36. | 7.9 | 35 |
| 99 | Characterization of the human CREB3L2 gene promoter. Oncology Reports, 2009, 21, 615-24. | 1.2 | 3 |
| 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 | 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 |
| 102 | Heterogeneous genetic profiles in soft tissue myoepitheliomas. Modern Pathology, 2008, 21, 1311-1319. | 2.9 | 44 |
| 103 | Fusion of the COL1A1 and USP6 genes in a benign bone tumor. Cancer Genetics and Cytogenetics, 2008, 180, 70-73. | 1.0 | 24 |
| 104 | An endometrial stromal sarcoma cell line with the JAZF1/PHF1 chimera. Cancer Genetics and Cytogenetics, 2008, 185, 74-77. | 1.0 | 37 |
| 105 | The POU5F1P1 pseudogene encodes a putative protein similar to POU5F1 isoform 1. Oncology Reports, 2008, 20, 1029-33. | 1.2 | 50 |
| 106 | Characterization of the native CREB3L2 transcription factor and the FUS/CREB3L2 chimera. Genes Chromosomes and Cancer, 2007, 46, 181-191. | 1.5 | 54 |
| 107 | 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 |
| 108 | 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 |

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|---|---|---|---|
| 109 | 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 |
| 110 | Confirmation of the high frequency of theTMPRSS2/ERG fusion gene in prostate cancer. Genes Chromosomes and Cancer, 2006, 45, 717-719. | 1.5 | 182 |
| 111 | Fusion of the SEC31L1 and ALK genes in an inflammatory myofibroblastic tumor. International Journal of Cancer, 2006, 118, 1181-1186. | 2.3 | 113 |
| 112 | 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 |
| 113 | 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 |
| 114 | Absence of mutations of the BRAF gene in malignant melanoma of soft parts (clear cell sarcoma of) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5 | 1.0 | 82 |
| 115 | The chimericFUS/CREB3l2gene is specific for low-grade fibromyxoid sarcoma. Genes Chromosomes and Cancer, 2004, 40, 218-228. | 1.5 | 188 |
| 116 | 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 |
| 117 | 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 |
| 118 | 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 |
| 119 | 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 |
| 120 | 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 |
| 121 | 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 |
| 122 | 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 |
| 123 | Fusion, Disruption, and Expression of HMGA2 in Bone and Soft Tissue Chondromas. Modern Pathology, 2003, 16, 1132-1140. | 2.9 | 82 |
| 124 | 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 |
| 125 | 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 |
| 126 | Molecular genetic characterization of theEWS/CHNandRBP56/CHNfusion genes in extraskeletal myxoid chondrosarcoma. Genes Chromosomes and Cancer, 2002, 35, 340-352. | 1.5 | 104 |

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|---|---|---|---|
| 127 | 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 |
| 128 | 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 |
| 129 | NoEWS/FLI1 fusion transcripts in giant-cell tumors of bone. International Journal of Cancer, 2001, 93, 769-772. | 2.3 | 35 |
| 130 | Clinical impact of molecular and cytogenetic findings in synovial sarcoma. Genes Chromosomes and Cancer, 2001, 31, 362-372. | 1.5 | 108 |
| 131 | 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 |
| 132 | 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 |
| 133 | Lentivirus Vector Gene Expression during ES Cell-Derived Hematopoietic Development In Vitro. Journal of Virology, 2000, 74, 10778-10784. | 1.5 | 100 |
| 134 | Acute Myeloid Leukemia with inv(8)(p11q13). Leukemia and Lymphoma, 2000, 39, 651-656. | 0.6 | 19 |
| 135 | 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 |
| 136 | A novel PCR-based approach for the detection of the Huntington disease associated trinucleotide repeat expansion. , 1999, 13, 232-236. | | 8 |
| 137 | A methylation PCR approach for detection of fragile X syndrome. , 1999, 14, 71-79. | | 39 |
| 138 | Identification of genes differentially expressed in TLS-CHOP carrying myxoid liposarcomas. , 1999, 83, 30-33. | | 64 |
| 139 | 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 |
| 140 | VariableFHIT transcripts in non-neoplastic tissues. , 1997, 19, 215-219. | | 38 |
| 141 | Genomic PCR detects tumor cells in peripheral blood from patients with myxoid liposarcoma. , 1996, 17, 102-107. | | 25 |
| 142 | Cloning and sequencing of a cDNA encoding ratd-dopachrome tautomerase. FEBS Letters, 1995, 373, 203-206. | 1.3 | 52 |
| 143 | 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 |