List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Identification of recurrent NAB2-STAT6 gene fusions in solitary fibrous tumor by integrative sequencing. Nature Genetics, 2013, 45, 180-185.	21.4	662
2	Molecular Characterization of Inflammatory Myofibroblastic Tumors With Frequent ALK and ROS1 Gene Fusions and Rare Novel RET Rearrangement. American Journal of Surgical Pathology, 2015, 39, 957-967.	3.7	281
3	A Molecular Study of Pediatric Spindle and Sclerosing Rhabdomyosarcoma. American Journal of Surgical Pathology, 2016, 40, 224-235.	3.7	208
4	BCOR-CCNB3 Fusion Positive Sarcomas. American Journal of Surgical Pathology, 2018, 42, 604-615.	3.7	207
5	Distinct transcriptional signature and immunoprofile of <i>CICâ€DUX4</i> fusion–positive round cell tumors compared to <i>EWSR1</i> â€rearranged ewing sarcomas: Further evidence toward distinct pathologic entities. Genes Chromosomes and Cancer, 2014, 53, 622-633.	2.8	201
6	Dichotomy of Genetic Abnormalities in PEComas With Therapeutic Implications. American Journal of Surgical Pathology, 2015, 39, 813-825.	3.7	177
7	Recurrent NTRK1 Gene Fusions Define a Novel Subset of Locally Aggressive Lipofibromatosis-like Neural Tumors. American Journal of Surgical Pathology, 2016, 40, 1407-1416.	3.7	177
8	BCOR Overexpression ls a Highly Sensitive Marker in Round Cell Sarcomas With BCOR Genetic Abnormalities. American Journal of Surgical Pathology, 2016, 40, 1670-1678.	3.7	168
9	Recurrent CIC Gene Abnormalities in Angiosarcomas. American Journal of Surgical Pathology, 2016, 40, 645-655.	3.7	157
10	Frequent FOS Gene Rearrangements in Epithelioid Hemangioma. American Journal of Surgical Pathology, 2015, 39, 1313-1321.	3.7	156
11	Recurrent BCOR Internal Tandem Duplication and YWHAE-NUTM2B Fusions in Soft Tissue Undifferentiated Round Cell Sarcoma of Infancy. American Journal of Surgical Pathology, 2016, 40, 1009-1020.	3.7	155
12	Novel <i>ZC3H7Bâ€BCOR</i> , <i>MEAF6â€PHF1</i> , and <i>EPC1â€PHF1</i> fusions in ossifying fibromyxoid tumors—molecular characterization shows genetic overlap with endometrial stromal sarcoma. Genes Chromosomes and Cancer, 2014, 53, 183-193.	2.8	145
13	Novel BCOR-MAML3 and ZC3H7B-BCOR Gene Fusions in Undifferentiated Small Blue Round Cell Sarcomas. American Journal of Surgical Pathology, 2016, 40, 433-442.	3.7	145
14	A novel group of spindle cell tumors defined by S100 and CD34 coâ€expression shows recurrent fusions involving RAF1, BRAF, and NTRK1/2 genes. Genes Chromosomes and Cancer, 2018, 57, 611-621.	2.8	144
15	Novel MIR143â€NOTCH fusions in benign and malignant glomus tumors. Genes Chromosomes and Cancer, 2013, 52, 1075-1087.	2.8	138
16	<i>ZFP36â€FOSB</i> fusion defines a subset of epithelioid hemangioma with atypical features. Genes Chromosomes and Cancer, 2014, 53, 951-959.	2.8	136
17	Novel <i>PRKD</i> gene rearrangements and variant fusions in cribriform adenocarcinoma of salivary gland origin. Genes Chromosomes and Cancer, 2014, 53, 845-856.	2.8	128
18	EWSR1 Fusions With CREB Family Transcription Factors Define a Novel Myxoid Mesenchymal Tumor With Predilection for Intracranial Location. American Journal of Surgical Pathology, 2017, 41, 482-490.	3.7	112

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19	A Distinct Malignant Epithelioid Neoplasm With GLI1 Gene Rearrangements, Frequent S100 Protein Expression, and Metastatic Potential. American Journal of Surgical Pathology, 2018, 42, 553-560.	3.7	109
20	Consistent PLAG1 and HMGA2 abnormalities distinguish carcinoma ex-pleomorphic adenoma from its de novo counterparts. Human Pathology, 2015, 46, 26-33.	2.0	103
21	NUTM1 Gene Fusions Characterize a Subset of Undifferentiated Soft Tissue and Visceral Tumors. American Journal of Surgical Pathology, 2018, 42, 636-645.	3.7	97
22	Ewing sarcoma with <scp> <i>ERG </i> </scp> gene rearrangements: A molecular study focusing on the prevalence of <scp> <i>FUSâ€ERG </i> </scp> and common pitfalls in detecting <scp> <i>EWSR1â€ERG </i> </scp> fusions by <scp>FISH </scp> . Genes Chromosomes and Cancer, 2016, 55, 340-349.	2.8	96
23	A genetic dichotomy between pure sclerosing epithelioid fibrosarcoma (SEF) and hybrid SEF/lowâ€grade fibromyxoid sarcoma: A pathologic and molecular study of 18 cases. Genes Chromosomes and Cancer, 2015, 54, 28-38.	2.8	95
24	Expanding the Spectrum of Intraosseous Rhabdomyosarcoma. American Journal of Surgical Pathology, 2019, 43, 695-702.	3.7	93
25	Recurrent RET Gene Rearrangements in Intraductal Carcinomas of Salivary Gland. American Journal of Surgical Pathology, 2018, 42, 442-452.	3.7	91
26	<i>EWSR1â€₽BX3</i> : A novel gene fusion in myoepithelial tumors. Genes Chromosomes and Cancer, 2015, 54, 63-71.	2.8	86
27	The histologic spectrum of soft tissue spindle cell tumors with <i>NTRK3</i> gene rearrangements. Genes Chromosomes and Cancer, 2019, 58, 739-746.	2.8	86
28	Recurrent BRAF Gene Fusions in a Subset of Pediatric Spindle Cell Sarcomas. American Journal of Surgical Pathology, 2018, 42, 28-38.	3.7	85
29	Extraskeletal myxoid chondrosarcoma with non–EWSR1-NR4A3 variant fusions correlate with rhabdoid phenotype and high-grade morphology. Human Pathology, 2014, 45, 1084-1091.	2.0	83
30	Novel <i>FUSâ€KLF17</i> and <i>EWSR1â€KLF17</i> fusions in myoepithelial tumors. Genes Chromosomes and Cancer, 2015, 54, 267-275.	2.8	82
31	Spindle Cell Tumors With RET Gene Fusions Exhibit a Morphologic Spectrum Akin to Tumors With NTRK Gene Fusions. American Journal of Surgical Pathology, 2019, 43, 1384-1391.	3.7	78
32	Recurrent SRF-RELA Fusions Define a Novel Subset of Cellular Myofibroma/Myopericytoma. American Journal of Surgical Pathology, 2017, 41, 677-684.	3.7	76
33	Uterine Tumor Resembling Ovarian Sex Cord Tumor. American Journal of Surgical Pathology, 2019, 43, 178-186.	3.7	72
34	GLI1-amplifications expand the spectrum of soft tissue neoplasms defined by GLI1 gene fusions. Modern Pathology, 2019, 32, 1617-1626.	5.5	70
35	Dermatofibrosarcoma protuberans with a novel <i>COL6A3â€PDGFD</i> fusion gene and apparent predilection for breast. Genes Chromosomes and Cancer, 2018, 57, 437-445.	2.8	61
36	Recurrent YAP1 and KMT2A Gene Rearrangements in a Subset of MUC4-negative Sclerosing Epithelioid Fibrosarcoma. American Journal of Surgical Pathology, 2020, 44, 368-377.	3.7	61

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37	<i>TFGâ€MET</i> fusion in an infantile spindle cell sarcoma with neural features. Genes Chromosomes and Cancer, 2017, 56, 663-667.	2.8	57
38	Novel EWSR1-SMAD3 Gene Fusions in a Group of Acral Fibroblastic Spindle Cell Neoplasms. American Journal of Surgical Pathology, 2018, 42, 522-528.	3.7	57
39	Soft tissue tumors characterized by a wide spectrum of kinase fusions share a lipofibromatosisâ€like neural tumor pattern. Genes Chromosomes and Cancer, 2020, 59, 575-583.	2.8	56
40	NTRK3 overexpression in undifferentiated sarcomas with YWHAE and BCOR genetic alterations. Modern Pathology, 2020, 33, 1341-1349.	5.5	53
41	ETV transcriptional upregulation is more reliable than RNA sequencing algorithms and FISH in diagnosing round cell sarcomas with <i>CIC</i> gene rearrangements. Genes Chromosomes and Cancer, 2017, 56, 501-510.	2.8	52
42	Expanding the molecular signature of ossifying fibromyxoid tumors with two novel gene fusions: <i>CREBBPâ€BCORL1</i> and <i>KDM2Aâ€WWTR1</i> . Genes Chromosomes and Cancer, 2017, 56, 42-50.	2.8	51
43	Recurrent YAP1 and MAML2 Gene Rearrangements in Retiform and Composite Hemangioendothelioma. American Journal of Surgical Pathology, 2020, 44, 1677-1684.	3.7	51
44	EWSR1/FUS–CREB fusions define a distinctive malignant epithelioid neoplasm with predilection for mesothelial-lined cavities. Modern Pathology, 2020, 33, 2233-2243.	5.5	49
45	Secondary <i>EWSR1</i> gene abnormalities in <i>SMARCB1</i> â€deficient tumors with 22q11â€12 regional deletions: Potential pitfalls in interpreting <i>EWSR1</i> FISH results. Genes Chromosomes and Cancer, 2016, 55, 767-776.	2.8	44
46	A morphologic and molecular reappraisal of myoepithelial tumors of soft tissue, bone, and viscera with EWSR1 and FUS gene rearrangements. Genes Chromosomes and Cancer, 2020, 59, 348-356.	2.8	44
47	Recurrent BRAF Gene Rearrangements in Myxoinflammatory Fibroblastic Sarcomas, but Not Hemosiderotic Fibrolipomatous Tumors. American Journal of Surgical Pathology, 2017, 41, 1456-1465.	3.7	40
48	Novel recurrent <i>PHF1â€TFE3</i> fusions in ossifying fibromyxoid tumors. Genes Chromosomes and Cancer, 2019, 58, 643-649.	2.8	39
49	Targeted exome sequencing profiles genetic alterations in leiomyosarcoma. Genes Chromosomes and Cancer, 2016, 55, 124-130.	2.8	38
50	Novel MEIS1-NCOA2 Gene Fusions Define a Distinct Primitive Spindle Cell Sarcoma of the Kidney. American Journal of Surgical Pathology, 2018, 42, 1562-1570.	3.7	35
51	A molecular study of synovial chondromatosis. Genes Chromosomes and Cancer, 2020, 59, 144-151.	2.8	31
52	BCOR upregulation in a poorly differentiated synovial sarcoma with <i>SS18L1 SX1</i> fusion—A pathologic and molecular pitfall. Genes Chromosomes and Cancer, 2017, 56, 296-302.	2.8	30
53	PGR Gene Fusions Identify a Molecular Subset of Uterine Epithelioid Leiomyosarcoma With Rhabdoid Features. American Journal of Surgical Pathology, 2019, 43, 810-818.	3.7	28
54	Epithelioid hemangioma of bone harboring <scp><i>FOS</i></scp> and <scp><i>FOSB</i></scp> gene rearrangements: A clinicopathologic and molecular study. Genes Chromosomes and Cancer, 2021, 60, 17-25.	2.8	28

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55	Novel GATA6-FOXO1 fusions in a subset of epithelioid hemangioma. Modern Pathology, 2021, 34, 934-941.	5.5	27
56	Clinical and molecular characterization of primary sclerosing epithelioid fibrosarcoma of bone and review of the literature. Genes Chromosomes and Cancer, 2020, 59, 217-224.	2.8	26
57	A novel <i>RBMXâ€TFE3</i> gene fusion in a highly aggressive pediatric renal perivascular epithelioid cell tumor. Genes Chromosomes and Cancer, 2020, 59, 58-63.	2.8	25
58	Frequent HRAS Mutations in Malignant Ectomesenchymoma. American Journal of Surgical Pathology, 2016, 40, 876-885.	3.7	24
59	BCOR Overexpression in Renal Malignant Solitary Fibrous Tumors. American Journal of Surgical Pathology, 2019, 43, 773-782.	3.7	24
60	<i>PRRXâ€NCOA1/2</i> rearrangement characterizes a distinctive fibroblastic neoplasm. Genes Chromosomes and Cancer, 2019, 58, 705-712.	2.8	23
61	Pediatric Mesothelioma With ALK Fusions. American Journal of Surgical Pathology, 2021, 45, 653-661.	3.7	22
62	Genetic diversity in alveolar soft part sarcoma: A subset contain variant fusion genes, highlighting broader molecular kinship with other MiT family tumors. Genes Chromosomes and Cancer, 2020, 59, 23-29.	2.8	19
63	Ewing sarcoma with <i>FEV</i> gene rearrangements is a rare subset with predilection for extraskeletal locations and aggressive behavior. Genes Chromosomes and Cancer, 2020, 59, 286-294.	2.8	18
64	Novel <i>SS18â€NEDD4</i> gene fusion in a primary renal synovial sarcoma. Genes Chromosomes and Cancer, 2020, 59, 203-208.	2.8	16
65	Novel SRF-ICA1L Fusions in Cellular Myoid Neoplasms With Potential For Malignant Behavior. American Journal of Surgical Pathology, 2020, 44, 55-60.	3.7	15
66	Undifferentiated round cell sarcomas with novelSS18â€₽OU5F1fusions. Genes Chromosomes and Cancer, 2020, 59, 620-626.	2.8	15
67	Comprehensive genomic profiling of EWSR1/FUS::CREB translocation-associated tumors uncovers prognostically significant recurrent genetic alterations and methylation-transcriptional correlates. Modern Pathology, 2022, 35, 1055-1065.	5.5	13
68	Hyalinizing epithelioid tumors with <scp><i>OGTâ€FOXO</i></scp> fusions. A case report of a nonâ€acral soft tissue mass harboring a novel <scp><i>FOXO4</i></scp> gene rearrangement. Genes Chromosomes and Cancer, 2021, 60, 498-503.	2.8	7
69	Generation of human embryonic stem cell models to exploit the EWSR1-CREB fusion promiscuity as a common pathway of transformation in human tumors. Oncogene, 2021, 40, 5095-5104.	5.9	7
70	Unclassified low grade spindle cell sarcoma with storiform pattern characterized by recurrent novel EWSR1/FUS-NACC1 fusions. Modern Pathology, 2021, 34, 1541-1546.	5.5	5