Yun-Shao Sung

List of Publications by Year
in descending order

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70 papers	6,090 citations	61984 43 h-index	88630 70 g-index
70 all docs	70 docs citations	70 times ranked	4575 citing authors

#	Article	IF	Citations
1	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
2	Epithelioid hemangioma of bone harboring <scp>FOS</scp> and <scp>FOSB</scp> gene rearrangements: A clinicopathologic and molecular study. Genes Chromosomes and Cancer, 2021, 60, 17-25.	2.8	28
3	Novel GATA6-FOXO1 fusions in a subset of epithelioid hemangioma. Modern Pathology, 2021, 34, 934-941.	5.5	27
4	Pediatric Mesothelioma With ALK Fusions. American Journal of Surgical Pathology, 2021, 45, 653-661.	3.7	22
5	Hyalinizing epithelioid tumors with <scp>OGTâ€FOXO</scp> fusions. A case report of a nonâ€sacral soft tissue mass harboring a novel <scp>FOXO4</scp> gene rearrangement. Genes Chromosomes and Cancer, 2021, 60, 498-503.	2.8	7
6	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
7	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
8	A molecular study of synovial chondromatosis. Genes Chromosomes and Cancer, 2020, 59, 144-151.	2.8	31
9	A novel RBMXâ€TFE3 gene fusion in a highly aggressive pediatric renal perivascular epithelioid cell tumor. Genes Chromosomes and Cancer, 2020, 59, 58-63.	2.8	25
10	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
11	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
12	Novel SS18â€NEDD4 gene fusion in a primary renal synovial sarcoma. Genes Chromosomes and Cancer, 2020, 59, 203-208.	2.8	16
13	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
14	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
15	Ewing sarcoma with FEV 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
16	EWSR1/FUSâ€“CREB fusions define a distinctive malignant epithelioid neoplasm with predilection for mesothelial-lined cavities. Modern Pathology, 2020, 33, 2233-2243.	5.5	49
17	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
18	Undifferentiated round cell sarcomas with novelSS18â€POU5F1fusions. Genes Chromosomes and Cancer, 2020, 59, 620-626.	2.8	15

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19	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
20	NTRK3 overexpression in undifferentiated sarcomas with YWHAE and BCOR genetic alterations. Modern Pathology, 2020, 33, 1341-1349.	5.5	53
21	Recurrent YAP1 and MAML2 Gene Rearrangements in Retiform and Composite Hemangioendothelioma. American Journal of Surgical Pathology, 2020, 44, 1677-1684.	3.7	51
22	GLI1-amplifications expand the spectrum of soft tissue neoplasms defined by GLI1 gene fusions. Modern Pathology, 2019, 32, 1617-1626.	5.5	70
23	The histologic spectrum of soft tissue spindle cell tumors with NTRK3 gene rearrangements. Genes Chromosomes and Cancer, 2019, 58, 739-746.	2.8	86
24	PRRXâ€NCOA1/2 rearrangement characterizes a distinctive fibroblastic neoplasm. Genes Chromosomes and Cancer, 2019, 58, 705-712.	2.8	23
25	Novel recurrent PHF1â€TFE3 fusions in ossifying fibromyxoid tumors. Genes Chromosomes and Cancer, 2019, 58, 643-649.	2.8	39
26	Uterine Tumor Resembling Ovarian Sex Cord Tumor. American Journal of Surgical Pathology, 2019, 43, 178-186.	3.7	72
27	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
28	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
29	Expanding the Spectrum of Intraosseous Rhabdomyosarcoma. American Journal of Surgical Pathology, 2019, 43, 695-702.	3.7	93
30	BCOR Overexpression in Renal Malignant Solitary Fibrous Tumors. American Journal of Surgical Pathology, 2019, 43, 773-782.	3.7	24
31	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
32	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
33	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
34	Recurrent BRAF Gene Fusions in a Subset of Pediatric Spindle Cell Sarcomas. American Journal of Surgical Pathology, 2018, 42, 28-38.	3.7	85
35	Recurrent RET Gene Rearrangements in Intraductal Carcinomas of Salivary Gland. American Journal of Surgical Pathology, 2018, 42, 442-452.	3.7	91
36	BCOR-CCNB3 Fusion Positive Sarcomas. American Journal of Surgical Pathology, 2018, 42, 604-615.	3.7	207

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37	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
38	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
39	Dermatofibrosarcoma protuberans with a novel COL6A3â€PDGFD fusion gene and apparent predilection for breast. Genes Chromosomes and Cancer, 2018, 57, 437-445.	2.8	61
40	ETV transcriptional upregulation is more reliable than RNA sequencing algorithms and FISH in diagnosing round cell sarcomas with CIC gene rearrangements. Genes Chromosomes and Cancer, 2017, 56, 501-510.	2.8	52
41	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
42	BCOR upregulation in a poorly differentiated synovial sarcoma with SS18L1â€SSX1 fusionâ€”A pathologic and molecular pitfall. Genes Chromosomes and Cancer, 2017, 56, 296-302.	2.8	30
43	Recurrent SRF-RELA Fusions Define a Novel Subset of Cellular Myofibroma/Myopericytoma. American Journal of Surgical Pathology, 2017, 41, 677-684.	3.7	76
44	TFGâ€MET fusion in an infantile spindle cell sarcoma with neural features. Genes Chromosomes and Cancer, 2017, 56, 663-667.	2.8	57
45	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
46	Expanding the molecular signature of ossifying fibromyxoid tumors with two novel gene fusions: CREBBPâ€BCORL1 and KDM2Aâ€WWTR1. Genes Chromosomes and Cancer, 2017, 56, 42-50.	2.8	51
47	Recurrent CIC Gene Abnormalities in Angiosarcomas. American Journal of Surgical Pathology, 2016, 40, 645-655.	3.7	157
48	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
49	Targeted exome sequencing profiles genetic alterations in leiomyosarcoma. Genes Chromosomes and Cancer, 2016, 55, 124-130.	2.8	38
50	Ewing sarcoma with <scp>ERG</scp> gene rearrangements: A molecular study focusing on the prevalence of <scp>FUSâ€ERG</scp> and common pitfalls in detecting <scp>EWSR1â€ERG</scp> fusions by <scp>FISH</scp>. Genes Chromosomes and Cancer, 2016, 55, 340-349.	2.8	96
51	Secondary EWSR1 gene abnormalities in SMARCB1â€deficient tumors with 22q11â€12 regional deletions: Potential pitfalls in interpreting EWSR1 FISH results. Genes Chromosomes and Cancer, 2016, 55, 767-776.	2.8	44
52	A Molecular Study of Pediatric Spindle and Sclerosing Rhabdomyosarcoma. American Journal of Surgical Pathology, 2016, 40, 224-235.	3.7	208
53	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
54	BCOR Overexpression Is a Highly Sensitive Marker in Round Cell Sarcomas With BCOR Genetic Abnormalities. American Journal of Surgical Pathology, 2016, 40, 1670-1678.	3.7	168

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55	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
56	Frequent HRAS Mutations in Malignant Ectomesenchymoma. American Journal of Surgical Pathology, 2016, 40, 876-885.	3.7	24
57	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
58	Dichotomy of Genetic Abnormalities in PEComas With Therapeutic Implications. American Journal of Surgical Pathology, 2015, 39, 813-825.	3.7	177
59	EWSR1â€PBX3: A novel gene fusion in myoepithelial tumors. Genes Chromosomes and Cancer, 2015, 54, 63-71.	2.8	86
60	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
61	Frequent FOS Gene Rearrangements in Epithelioid Hemangioma. American Journal of Surgical Pathology, 2015, 39, 1313-1321.	3.7	156
62	Novel FUSâ€KLF17 and EWSR1â€KLF17 fusions in myoepithelial tumors. Genes Chromosomes and Cancer, 2015, 54, 267-275.	2.8	82
63	Consistent PLAG1 and HMGA2 abnormalities distinguish carcinoma ex-pleomorphic adenoma from its de novo counterparts. Human Pathology, 2015, 46, 26-33.	2.0	103
64	ZFP36â€FOSB fusion defines a subset of epithelioid hemangioma with atypical features. Genes Chromosomes and Cancer, 2014, 53, 951-959.	2.8	136
65	Distinct transcriptional signature and immunoprofile of CICâ€DUX4 fusionâ€“positive round cell tumors compared to EWSR1â€rearranged ewing sarcomas: Further evidence toward distinct pathologic entities. Genes Chromosomes and Cancer, 2014, 53, 622-633.	2.8	201
66	Novel PRKD gene rearrangements and variant fusions in cribriform adenocarcinoma of salivary gland origin. Genes Chromosomes and Cancer, 2014, 53, 845-856.	2.8	128
67	Novel ZC3H7Bâ€BCOR, MEAF6â€PHF1, and EPC1â€PHF1 fusions in ossifying fibromyxoid tumorsâ€”molecular characterization shows genetic overlap with endometrial stromal sarcoma. Genes Chromosomes and Cancer, 2014, 53, 183-193.	2.8	145
68	Extraskeletal myxoid chondrosarcoma with nonâ€“EWSR1-NR4A3 variant fusions correlate with rhabdoid phenotype and high-grade morphology. Human Pathology, 2014, 45, 1084-1091.	2.0	83
69	Identification of recurrent NAB2-STAT6 gene fusions in solitary fibrous tumor by integrative sequencing. Nature Genetics, 2013, 45, 180-185.	21.4	662
70	Novel MIR143â€NOTCH fusions in benign and malignant glomus tumors. Genes Chromosomes and Cancer, 2013, 52, 1075-1087.	2.8	138