

Franck Tirode

List of Publications by Citations

Source: <https://exaly.com/author-pdf/9446670/franck-tirode-publications-by-citations.pdf>

Version: 2024-04-28

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

95
papers

4,677
citations

33
h-index

68
g-index

115
ext. papers

5,787
ext. citations

9.1
avg, IF

5.06
L-index

#	Paper	IF	Citations
95	Mesenchymal stem cell features of Ewing tumors. <i>Cancer Cell</i> , 2007 , 11, 421-9	24.3	390
94	EWS/FLI-1 silencing and gene profiling of Ewing cells reveal downstream oncogenic pathways and a crucial role for repression of insulin-like growth factor binding protein 3. <i>Molecular and Cellular Biology</i> , 2004 , 24, 7275-83	4.8	342
93	A new subtype of bone sarcoma defined by BCOR-CCNB3 gene fusion. <i>Nature Genetics</i> , 2012 , 44, 461-6	36.3	325
92	Genomic landscape of Ewing sarcoma defines an aggressive subtype with co-association of STAG2 and TP53 mutations. <i>Cancer Discovery</i> , 2014 , 4, 1342-53	24.4	310
91	Validated prediction of clinical outcome in sarcomas and multiple types of cancer on the basis of a gene expression signature related to genome complexity. <i>Nature Medicine</i> , 2010 , 16, 781-7	50.5	290
90	Reconstitution of the transcription factor TFIIH: assignment of functions for the three enzymatic subunits, XPB, XPD, and cdk7. <i>Molecular Cell</i> , 1999 , 3, 87-95	17.6	255
89	Cancer-associated SF3B1 mutations affect alternative splicing by promoting alternative branchpoint usage. <i>Nature Communications</i> , 2016 , 7, 10615	17.4	223
88	SMARCA4 inactivation defines a group of undifferentiated thoracic malignancies transcriptionally related to BAF-deficient sarcomas. <i>Nature Genetics</i> , 2015 , 47, 1200-5	36.3	170
87	Translational Activation of HIF1 β by YB-1 Promotes Sarcoma Metastasis. <i>Cancer Cell</i> , 2015 , 27, 682-97	24.3	167
86	Nucleotide excision repair of DNA with recombinant human proteins: definition of the minimal set of factors, active forms of TFIIH, and modulation by CAK. <i>Genes and Development</i> , 2000 , 14, 349-359	12.6	158
85	Transcriptomic definition of molecular subgroups of small round cell sarcomas. <i>Journal of Pathology</i> , 2018 , 245, 29-40	9.4	140
84	Common variants near TARDBP and EGR2 are associated with susceptibility to Ewing sarcoma. <i>Nature Genetics</i> , 2012 , 44, 323-7	36.3	124
83	The oncogenic EWS-FLI1 protein binds in vivo GGAA microsatellite sequences with potential transcriptional activation function. <i>PLoS ONE</i> , 2009 , 4, e4932	3.7	121
82	Chimeric EWSR1-FLI1 regulates the Ewing sarcoma susceptibility gene EGR2 via a GGAA microsatellite. <i>Nature Genetics</i> , 2015 , 47, 1073-8	36.3	103
81	A conditionally expressed third partner stabilizes or prevents the formation of a transcriptional activator in a three-hybrid system. <i>Journal of Biological Chemistry</i> , 1997 , 272, 22995-9	5.4	90
80	A role for the TFIIH XPB DNA helicase in promoter escape by RNA polymerase II. <i>Journal of Biological Chemistry</i> , 1999 , 274, 22127-30	5.4	74
79	ETV4 is a useful marker for the diagnosis of CIC-rearranged undifferentiated round-cell sarcomas: a study of 127 cases including mimicking lesions. <i>Modern Pathology</i> , 2016 , 29, 1523-1531	9.8	71

78	Zoledronic acid as a new adjuvant therapeutic strategy for Ewing's sarcoma patients. <i>Cancer Research</i> , 2010 , 70, 7610-9	10.1	64
77	SMARCA4-deficient Thoracic Sarcomas: Clinicopathologic Study of 30 Cases With an Emphasis on Their Nosology and Differential Diagnoses. <i>American Journal of Surgical Pathology</i> , 2019 , 43, 455-465	6.7	64
76	The orphan nuclear receptor DAX1 is up-regulated by the EWS/FLI1 oncoprotein and is highly expressed in Ewing tumors. <i>International Journal of Cancer</i> , 2006 , 118, 1381-9	7.5	63
75	Strong smooth muscle differentiation is dependent on myocardin gene amplification in most human retroperitoneal leiomyosarcomas. <i>Cancer Research</i> , 2009 , 69, 2269-78	10.1	57
74	Targeting the EWSR1-FLI1 oncogene-induced protein kinase PKC δ abolishes ewing sarcoma growth. <i>Cancer Research</i> , 2012 , 72, 4494-503	10.1	54
73	Combined experience of six independent laboratories attempting to create an Ewing sarcoma mouse model. <i>Oncotarget</i> , 2017 , 8, 34141-34163	3.3	52
72	Preclinical evidence that use of TRAIL in Ewing's sarcoma and osteosarcoma therapy inhibits tumor growth, prevents osteolysis, and increases animal survival. <i>Clinical Cancer Research</i> , 2010 , 16, 2363-74	12.9	51
71	De novo motif identification improves the accuracy of predicting transcription factor binding sites in CHIP-Seq data analysis. <i>Nucleic Acids Research</i> , 2010 , 38, e126	20.1	51
70	Clinicopathologic Features of CIC-NUTM1 Sarcomas, a New Molecular Variant of the Family of CIC-Fused Sarcomas. <i>American Journal of Surgical Pathology</i> , 2019 , 43, 268-276	6.7	44
69	Systems biology of Ewing sarcoma: a network model of EWS-FLI1 effect on proliferation and apoptosis. <i>Nucleic Acids Research</i> , 2013 , 41, 8853-71	20.1	39
68	Transcriptional Programs Define Intratumoral Heterogeneity of Ewing Sarcoma at Single-Cell Resolution. <i>Cell Reports</i> , 2020 , 30, 1767-1779.e6	10.6	39
67	The second European interdisciplinary Ewing sarcoma research summit--A joint effort to deconstructing the multiple layers of a complex disease. <i>Oncotarget</i> , 2016 , 7, 8613-24	3.3	38
66	Alternative PDGFD rearrangements in dermatofibrosarcomas protuberans without PDGFB fusions. <i>Modern Pathology</i> , 2018 , 31, 1683-1693	9.8	37
65	Targeting the epigenetic readers in Ewing sarcoma inhibits the oncogenic transcription factor EWS/FlI1. <i>Oncotarget</i> , 2016 , 7, 24125-40	3.3	35
64	A subset of epithelioid and spindle cell rhabdomyosarcomas is associated with TFCP2 fusions and common ALK upregulation. <i>Modern Pathology</i> , 2020 , 33, 404-419	9.8	35
63	Clinicopathologic and Molecular Features of a Series of 41 Biphenotypic Sinonasal Sarcomas Expanding Their Molecular Spectrum. <i>American Journal of Surgical Pathology</i> , 2019 , 43, 747-754	6.7	33
62	Update on Families of Round Cell Sarcomas Other than Classical Ewing Sarcomas. <i>Surgical Pathology Clinics</i> , 2017 , 10, 587-620	3.9	32
61	deficiency causes a wide tumor spectrum and increases embryonal rhabdomyosarcoma metastasis in zebrafish. <i>ELife</i> , 2018 , 7,	8.9	31

60	Oncostatin M is a growth factor for Ewing sarcoma. <i>American Journal of Pathology</i> , 2012 , 181, 1782-95	5.8	30
59	The first European interdisciplinary ewing sarcoma research summit. <i>Frontiers in Oncology</i> , 2012 , 2, 54	5.3	29
58	Comprehensive Molecular and Pathologic Evaluation of Transitional Mesothelioma Assisted by Deep Learning Approach: A Multi-Institutional Study of the International Mesothelioma Panel from the MESOPATH Reference Center. <i>Journal of Thoracic Oncology</i> , 2020 , 15, 1037-1053	8.9	27
57	Therapeutic Targeting of KDM1A/LSD1 in Ewing Sarcoma with SP-2509 Engages the Endoplasmic Reticulum Stress Response. <i>Molecular Cancer Therapeutics</i> , 2018 , 17, 1902-1916	6.1	27
56	Genome-wide association study identifies multiple new loci associated with Ewing sarcoma susceptibility. <i>Nature Communications</i> , 2018 , 9, 3184	17.4	25
55	BAffling pathologies: Alterations of BAF complexes in cancer. <i>Cancer Letters</i> , 2018 , 419, 266-279	9.9	24
54	SRF-FOXO1 and SRF-NCOA1 Fusion Genes Delineate a Distinctive Subset of Well-differentiated Rhabdomyosarcoma. <i>American Journal of Surgical Pathology</i> , 2020 , 44, 607-616	6.7	22
53	transgenic zebrafish models identify as a mediator of rhabdomyosarcoma tumorigenesis. <i>ELife</i> , 2018 , 7,	8.9	20
52	Bone sarcomas: from biology to targeted therapies. <i>Sarcoma</i> , 2012 , 2012, 301975	3.1	19
51	Loss of connexin43 expression in Ewing sarcoma cells favors the development of the primary tumor and the associated bone osteolysis. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2013 , 1832, 553-64	6.9	17
50	Melanocytic tumors with MAP3K8 fusions: report of 33 cases with morphological-genetic correlations. <i>Modern Pathology</i> , 2020 , 33, 846-857	9.8	17
49	Integrative clinical and biopathology analyses to understand the clinical heterogeneity of infantile rhabdomyosarcoma: A report from the French MMT committee. <i>Cancer Medicine</i> , 2020 , 9, 2698-2709	4.8	16
48	NFATc2-rearranged sarcomas: clinicopathologic, molecular, and cytogenetic study of 7 cases with evidence of AGGRECAN as a novel diagnostic marker. <i>Modern Pathology</i> , 2020 , 33, 1930-1944	9.8	16
47	Genetic analyses of undifferentiated small round cell sarcoma identifies a novel sarcoma subtype with a recurrent CRTCl-SS18 gene fusion. <i>Journal of Pathology</i> , 2018 , 245, 186-196	9.4	15
46	The ENCCA-WP7/EuroSarc/EEC/PROVABES/EURAMOS 3rd European Bone Sarcoma Networking Meeting/Joint Workshop of EU Bone Sarcoma Translational Research Networks; Vienna, Austria, September 24-25, 2015. Workshop Report. <i>Clinical Sarcoma Research</i> , 2016 , 6, 3	2.5	14
45	STAG2 mutations alter CTCF-anchored loop extrusion, reduce cis-regulatory interactions and EWSR1-FLI1 activity in Ewing sarcoma. <i>Cancer Cell</i> , 2021 , 39, 810-826.e9	24.3	13
44	Very long-term survivors among patients with metastatic soft tissue sarcoma. <i>Cancer Medicine</i> , 2019 , 8, 1368-1378	4.8	11
43	Molecular Classification of Endometrial Stromal Sarcomas Using RNA Sequencing Defines Nosological and Prognostic Subgroups with Different Natural History. <i>Cancers</i> , 2020 , 12,	6.6	11

42	Genomic and transcriptomic characterisation of undifferentiated pleomorphic sarcoma of bone. <i>Journal of Pathology</i> , 2019 , 247, 166-176	9.4	11
41	Malignant melanoma with areas of rhabdomyosarcomatous differentiation arising in a giant congenital nevus with RAF1 gene fusion. <i>Pigment Cell and Melanoma Research</i> , 2019 , 32, 708-713	4.5	9
40	A functional, new short isoform of death receptor 4 in Ewing sarcoma cell lines may be involved in TRAIL sensitivity/resistance mechanisms. <i>Molecular Cancer Research</i> , 2012 , 10, 336-46	6.6	9
39	Novel three-way complex rearrangement of TRPM1-PUM1-LCK in a case of agminated Spitz nevi arising in a giant congenital hyperpigmented macule. <i>Pigment Cell and Melanoma Research</i> , 2020 , 33, 767-772	4.5	8
38	CYSLTR2-mutant Cutaneous Melanocytic Neoplasms Frequently Simulate "Pigmented Epithelioid Melanocytoma," Expanding the Morphologic Spectrum of Blue Tumors: A Clinicopathologic Study of 7 Cases. <i>American Journal of Surgical Pathology</i> , 2019 , 43, 1368-1376	6.7	8
37	Brain tumor with an ATXN1-NUTM1 fusion gene expands the histologic spectrum of NUTM1-rearranged neoplasia. <i>Acta Neuropathologica Communications</i> , 2019 , 7, 220	7.3	8
36	Antagonism pattern detection between microRNA and target expression in Ewing sarcoma. <i>PLoS ONE</i> , 2012 , 7, e41770	3.7	7
35	Fusion partners of NTRK3 affect subcellular localization of the fusion kinase and cytomorphology of melanocytes. <i>Modern Pathology</i> , 2021 , 34, 735-747	9.8	7
34	Development of curative therapies for Ewing sarcomas by interdisciplinary cooperative groups in Europe. <i>Klinische Padiatrie</i> , 2015 , 227, 108-15	0.9	6
33	CRTC1-TRIM11 fusion defined melanocytic tumors: A series of four cases. <i>Journal of Cutaneous Pathology</i> , 2019 , 46, 810-818	1.7	6
32	Clear cell tumor with melanocytic differentiation and MITF-CREB translocation: a novel entity similar to clear cell sarcoma. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2021 , 479, 841-846	5.1	6
31	Three-hybrid screens. Inducible third-party systems. <i>Methods in Molecular Biology</i> , 2001 , 177, 271-89	1.4	5
30	SRF Fusions Other Than With RELA Expand the Molecular Definition of SRF-fused Perivascular Tumors. <i>American Journal of Surgical Pathology</i> , 2020 , 44, 1725-1735	6.7	5
29	Neonatal Soft Tissue Sarcoma with YWHAE-NUTM2B Fusion. <i>Case Reports in Oncology</i> , 2019 , 12, 631-638		4
28	Osteoprotegerin inhibits bone resorption and prevents tumor development in a xenogenic model of Ewing sarcoma by inhibiting RANKL. <i>Journal of Bone Oncology</i> , 2013 , 2, 95-104	4.5	4
27	EWS-FLI1 inhibits TNFalpha-induced NFkappaB-dependent transcription in Ewing sarcoma cells. <i>Biochemical and Biophysical Research Communications</i> , 2010 , 399, 705-10	3.4	4
26	Clear Cell Tumor With Melanocytic Differentiation and ACTIN-MITF Translocation: Report of 7 Cases of a Novel Entity. <i>American Journal of Surgical Pathology</i> , 2021 , 45, 962-968	6.7	4
25	Cutaneous Melanocytic Tumors With Concomitant NRASQ61R and IDH1R132C Mutations: A Report of 6 Cases. <i>American Journal of Surgical Pathology</i> , 2020 , 44, 1398-1405	6.7	4

24	Superficial CD34-positive fibroblastic tumor and PRDM10-rearranged soft tissue tumor are overlapping entities: a comprehensive study of 20 cases. <i>Histopathology</i> , 2021 , 79, 810-825	7.3	4
23	Abnormal vascularization of soft-tissue sarcomas on conventional MRI: Diagnostic and prognostic values. <i>European Journal of Radiology</i> , 2019 , 117, 112-119	4.7	3
22	Recurrent novel THBS1-ADGRF5 gene fusion in a new tumor subtype "Acral FibroChondroMyxoid Tumors". <i>Modern Pathology</i> , 2020 , 33, 1360-1368	9.8	3
21	GOPC-ROS1 mosaicism in agminated Spitz naevi: report of two cases. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2021 , 479, 559-564	5.1	3
20	Spitz nevus with a novel TFG-NTRK2 fusion: The first case report of NTRK2-rearranged Spitz/Reed nevus. <i>Journal of Cutaneous Pathology</i> , 2021 , 48, 1193-1196	1.7	3
19	Wholistic approach - transcriptomic analysis and beyond using archival material for molecular diagnosis.. <i>Genes Chromosomes and Cancer</i> , 2022 ,	5	2
18	Morphologic features in a series of 352 Spitz melanocytic proliferations help predict their oncogenic drivers. <i>Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin</i> , 2021 , 480, 369	5.1	2
17	Low-frequency variation near common germline susceptibility loci are associated with risk of Ewing sarcoma. <i>PLoS ONE</i> , 2020 , 15, e0237792	3.7	2
16	Nodular Fasciitis With Malignant Morphology and a Fusion: A Case Report (of a 10-Year-old Boy). <i>International Journal of Surgical Pathology</i> , 2021 , 29, 642-647	1.2	2
15	RASGRF2 gene fusions identified in a variety of melanocytic lesions with distinct morphological features. <i>Pigment Cell and Melanoma Research</i> , 2021 , 34, 1074-1083	4.5	2
14	RASGRF1-rearranged Cutaneous Melanocytic Neoplasms With Spitzoid Cytomorphology: A Clinicopathologic and Genetic Study of 3 Cases. <i>American Journal of Surgical Pathology</i> , 2021 ,	6.7	1
13	CIC-DUX4 expression drives the development of small round cell sarcoma in transgenic zebrafish: a new model revealing a role for ETV4 in CIC-mediated sarcomagenesis		1
12	Tocilizumab for the treatment of paraneoplastic inflammatory syndrome associated with angiomatoid fibrous histiocytoma. <i>ESMO Open</i> , 2020 , 5, e000756	6	1
11	ERG transcription factors have a splicing regulatory function involving RBFOX2 that is altered in the EWS-FLI1 oncogenic fusion. <i>Nucleic Acids Research</i> , 2021 , 49, 5038-5056	20.1	1
10	Aldehyde Dehydrogenase, a Therapeutic Target in Chordoma: Analysis in 3D Cellular Models. <i>Cells</i> , 2021 , 10,	7.9	1
9	Solid papillary mesothelial tumor. <i>Modern Pathology</i> , 2021 ,	9.8	1
8	FNBP1-BRAF fusion in a primary melanoma of the lung. <i>Pathology</i> , 2021 , 53, 785-788	1.6	1
7	Agminated Spitz naevus with an activating HRAS Q61R mutation. <i>Pathology</i> , 2021 ,	1.6	0

6 Biology of Ewing sarcoma **2015**, 245-255

5 Molecular aspects of Ewing sarcomas **2022**, 617-630

4 Low-frequency variation near common germline susceptibility loci are associated with risk of Ewing sarcoma **2020**, 15, e0237792

3 Low-frequency variation near common germline susceptibility loci are associated with risk of Ewing sarcoma **2020**, 15, e0237792

2 Low-frequency variation near common germline susceptibility loci are associated with risk of Ewing sarcoma **2020**, 15, e0237792

1 Low-frequency variation near common germline susceptibility loci are associated with risk of Ewing sarcoma **2020**, 15, e0237792