List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	The Immune Landscape of Cancer. Immunity, 2018, 48, 812-830.e14.	6.6	3,706
2	Gut microbiome modulates response to anti–PD-1 immunotherapy in melanoma patients. Science, 2018, 359, 97-103.	6.0	3,126
3	An Integrated TCGA Pan-Cancer Clinical Data Resource to Drive High-Quality Survival Outcome Analytics. Cell, 2018, 173, 400-416.e11.	13.5	2,277
4	Oncogenic Signaling Pathways in The Cancer Genome Atlas. Cell, 2018, 173, 321-337.e10.	13.5	2,111
5	Cell-of-Origin Patterns Dominate the Molecular Classification of 10,000 Tumors from 33 Types of Cancer. Cell, 2018, 173, 291-304.e6.	13.5	1,718
6	Comprehensive Characterization of Cancer Driver Genes and Mutations. Cell, 2018, 173, 371-385.e18.	13.5	1,670
7	Melanoma staging: Evidenceâ€based changes in the American Joint Committee on Cancer eighth edition cancer staging manual. Ca-A Cancer Journal for Clinicians, 2017, 67, 472-492.	157.7	1,662
8	B cells and tertiary lymphoid structures promote immunotherapy response. Nature, 2020, 577, 549-555.	13.7	1,421
9	Machine Learning Identifies Stemness Features Associated with Oncogenic Dedifferentiation. Cell, 2018, 173, 338-354.e15.	13.5	1,417
10	Loss of PTEN Promotes Resistance to T Cell–Mediated Immunotherapy. Cancer Discovery, 2016, 6, 202-216.	7.7	1,158
11	B cells are associated with survival and immunotherapy response in sarcoma. Nature, 2020, 577, 556-560.	13.7	1,158
12	The 5th edition of the World Health Organization Classification of Haematolymphoid Tumours: Lymphoid Neoplasms. Leukemia, 2022, 36, 1720-1748.	3.3	1,023
13	Pembrolizumab in advanced soft-tissue sarcoma and bone sarcoma (SARC028): a multicentre, two-cohort, single-arm, open-label, phase 2 trial. Lancet Oncology, The, 2017, 18, 1493-1501.	5.1	921
14	Analysis of Immune Signatures in Longitudinal Tumor Samples Yields Insight into Biomarkers of Response and Mechanisms of Resistance to Immune Checkpoint Blockade. Cancer Discovery, 2016, 6, 827-837.	7.7	785
15	Genomic and Functional Approaches to Understanding Cancer Aneuploidy. Cancer Cell, 2018, 33, 676-689.e3.	7.7	750
16	Comprehensive and Integrated Genomic Characterization of Adult Soft Tissue Sarcomas. Cell, 2017, 171, 950-965.e28.	13.5	738
17	Integrated molecular analysis of tumor biopsies on sequential CTLA-4 and PD-1 blockade reveals markers of response and resistance. Science Translational Medicine, 2017, 9, .	5.8	689
18	Spatial Organization and Molecular Correlation of Tumor-Infiltrating Lymphocytes Using Deep Learning on Pathology Images. Cell Reports, 2018, 23, 181-193.e7.	2.9	683

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19	Circulating Tumor DNA Analysis in Patients With Cancer: American Society of Clinical Oncology and College of American Pathologists Joint Review. Journal of Clinical Oncology, 2018, 36, 1631-1641.	0.8	668
20	Integrative Analysis Identifies Four Molecular and Clinical Subsets in Uveal Melanoma. Cancer Cell, 2017, 32, 204-220.e15.	7.7	642
21	Pathogenic Germline Variants in 10,389 Adult Cancers. Cell, 2018, 173, 355-370.e14.	13.5	620
22	Neoadjuvant immune checkpoint blockade in high-risk resectable melanoma. Nature Medicine, 2018, 24, 1649-1654.	15.2	592
23	<i>NRAS</i> mutation status is an independent prognostic factor in metastatic melanoma. Cancer, 2012, 118, 4014-4023.	2.0	589
24	Fecal microbiota transplantation for refractory immune checkpoint inhibitor-associated colitis. Nature Medicine, 2018, 24, 1804-1808.	15.2	521
25	A Comprehensive Pan-Cancer Molecular Study of Gynecologic and Breast Cancers. Cancer Cell, 2018, 33, 690-705.e9.	7.7	478
26	Specific Mutations in the β-Catenin Gene (CTNNB1) Correlate with Local Recurrence in Sporadic Desmoid Tumors. American Journal of Pathology, 2008, 173, 1518-1527.	1.9	417
27	Driver Fusions and Their Implications in the Development and Treatment of Human Cancers. Cell Reports, 2018, 23, 227-238.e3.	2.9	407
28	Solitary fibrous tumor: a clinicopathological study of 110 cases and proposed risk assessment model. Modern Pathology, 2012, 25, 1298-1306.	2.9	403
29	Comparative Molecular Analysis of Gastrointestinal Adenocarcinomas. Cancer Cell, 2018, 33, 721-735.e8.	7.7	396
30	Dietary fiber and probiotics influence the gut microbiome and melanoma immunotherapy response. Science, 2021, 374, 1632-1640.	6.0	369
31	Imatinib Mesylate in Advanced Dermatofibrosarcoma Protuberans: Pooled Analysis of Two Phase II Clinical Trials. Journal of Clinical Oncology, 2010, 28, 1772-1779.	0.8	351
32	Examination of Mutations in BRAF, NRAS, and PTEN in Primary Cutaneous Melanoma. Journal of Investigative Dermatology, 2006, 126, 154-160.	0.3	334
33	Identification of a Disease-Defining Gene Fusion in Epithelioid Hemangioendothelioma. Science Translational Medicine, 2011, 3, 98ra82.	5.8	328
34	Epithelioid Inflammatory Myofibroblastic Sarcoma. American Journal of Surgical Pathology, 2011, 35, 135-144.	2.1	309
35	NBTXR3, a first-in-class radioenhancer hafnium oxide nanoparticle, plus radiotherapy versus radiotherapy alone in patients with locally advanced soft-tissue sarcoma (Act.In.Sarc): a multicentre, phase 2–3, randomised, controlled trial. Lancet Oncology, The, 2019, 20, 1148-1159.	5.1	288
36	American Joint Committee on Cancer acceptance criteria for inclusion of risk models for individualized prognosis in the practice of precision medicine. Ca-A Cancer Journal for Clinicians, 2016, 66, 370-374.	157.7	280

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37	Histopathologic evaluation of atypical neurofibromatous tumors and their transformation into malignant peripheral nerve sheath tumor in patients with neurofibromatosis 1—a consensus overview. Human Pathology, 2017, 67, 1-10.	1.1	275
38	Perspective on Oncogenic Processes at the End of the Beginning of Cancer Genomics. Cell, 2018, 173, 305-320.e10.	13.5	272
39	Recurrent PTPRB and PLCG1 mutations in angiosarcoma. Nature Genetics, 2014, 46, 376-379.	9.4	269
40	Risk assessment in solitary fibrous tumors: validation and refinement of a risk stratification model. Modern Pathology, 2017, 30, 1433-1442.	2.9	261
41	Clinical, Pathological, and Molecular Variables Predictive of Malignant Peripheral Nerve Sheath Tumor Outcome. Annals of Surgery, 2009, 249, 1014-1022.	2.1	254
42	Genomic, Pathway Network, and Immunologic Features Distinguishing Squamous Carcinomas. Cell Reports, 2018, 23, 194-212.e6.	2.9	245
43	Neoadjuvant plus adjuvant dabrafenib and trametinib versus standard of care in patients with high-risk, surgically resectable melanoma: a single-centre, open-label, randomised, phase 2 trial. Lancet Oncology, The, 2018, 19, 181-193.	5.1	233
44	Molecular Profiling Reveals Unique Immune and Metabolic Features of Melanoma Brain Metastases. Cancer Discovery, 2019, 9, 628-645.	7.7	231
45	MUC4 Is a Sensitive and Extremely Useful Marker for Sclerosing Epithelioid Fibrosarcoma. American Journal of Surgical Pathology, 2012, 36, 1444-1451.	2.1	230
46	Integrated Molecular and Clinical Analysis of AKT Activation in Metastatic Melanoma. Clinical Cancer Research, 2009, 15, 7538-7546.	3.2	221
47	Gut microbiota signatures are associated with toxicity to combined CTLA-4 and PD-1 blockade. Nature Medicine, 2021, 27, 1432-1441.	15.2	216
48	Dermatofibrosarcoma protuberans COL1A1-PDGFB fusion is identified in virtually all dermatofibrosarcoma protuberans cases when investigated by newly developed multiplex reverse transcription polymerase chain reaction and fluorescence in situ hybridization assays. Human Pathology, 2008, 39, 184-193.	1.1	214
49	Activity of temozolomide and bevacizumab in the treatment of locally advanced, recurrent, and metastatic hemangiopericytoma and malignant solitary fibrous tumor. Cancer, 2011, 117, 4939-4947.	2.0	212
50	Modeling Ewing sarcoma tumors in vitro with 3D scaffolds. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 6500-6505.	3.3	210
51	Detection and characterization of EWSR1/ATF1 and EWSR1/CREB1 chimeric transcripts in clear cell sarcoma (melanoma of soft parts). Modern Pathology, 2009, 22, 1201-1209.	2.9	198
52	Correlation between KIT expression and KIT mutation in melanoma: a study of 173 cases with emphasis on the acral-lentiginous/mucosal type. Modern Pathology, 2009, 22, 1446-1456.	2.9	196
53	Frequency and Spectrum of BRAF Mutations in a Retrospective, Single-Institution Study of 1112 Cases of Melanoma. Journal of Molecular Diagnostics, 2013, 15, 220-226.	1.2	195
54	Autophagy inhibition and antimalarials promote cell death in gastrointestinal stromal tumor (GIST). Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 14333-14338.	3.3	194

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55	Telomere dysfunction suppresses spontaneous tumorigenesis in vivo by initiating p53â€dependent cellular senescence. EMBO Reports, 2007, 8, 497-503.	2.0	185
56	Clinical Responses to Vemurafenib in Patients with Metastatic Papillary Thyroid Cancer Harboring BRAF ^{V600E} Mutation. Thyroid, 2013, 23, 1277-1283.	2.4	184
57	Angiosarcoma. Annals of Surgery, 2010, 251, 1098-1106.	2.1	182
58	Activity of dasatinib against <i>L576P KIT</i> mutant melanoma: Molecular, cellular, and clinical correlates. Molecular Cancer Therapeutics, 2009, 8, 2079-2085.	1.9	178
59	Molecular Profiling of Patient-Matched Brain and Extracranial Melanoma Metastases Implicates the PI3K Pathway as a Therapeutic Target. Clinical Cancer Research, 2014, 20, 5537-5546.	3.2	169
60	The SS18-SSX Fusion Oncoprotein Hijacks BAF Complex Targeting and Function to Drive Synovial Sarcoma. Cancer Cell, 2018, 33, 1128-1141.e7.	7.7	169
61	Genetic Alterations in Signaling Pathways in Melanoma. Clinical Cancer Research, 2006, 12, 2301s-2307s.	3.2	168
62	Extensive Survey of STAT6 Expression in a Large Series of Mesenchymal Tumors. American Journal of Clinical Pathology, 2015, 143, 672-682.	0.4	168
63	A Randomized, Phase II Study of Preoperative plus Postoperative Imatinib in GIST: Evidence of Rapid Radiographic Response and Temporal Induction of Tumor Cell Apoptosis. Annals of Surgical Oncology, 2009, 16, 910-919.	0.7	166
64	Vimentin Is a Novel Anti-Cancer Therapeutic Target; Insights from In Vitro and In Vivo Mice Xenograft Studies. PLoS ONE, 2010, 5, e10105.	1.1	166
65	Loss of H3K27 tri-methylation is a diagnostic marker for malignant peripheral nerve sheath tumors and an indicator for an inferior survival. Modern Pathology, 2016, 29, 582-590.	2.9	164
66	Cutaneous Clear Cell Sarcoma: A Clinicopathologic, Immunohistochemical, and Molecular Analysis of 12 Cases Emphasizing its Distinction from Dermal Melanoma. American Journal of Surgical Pathology, 2010, 34, 216-222.	2.1	163
67	CTNNB1 45F mutation is a molecular prognosticator of increased postoperative primary desmoid tumor recurrence. Cancer, 2013, 119, 3696-3702.	2.0	162
68	Adipophilin expression in sebaceous tumors and other cutaneous lesions with clear cell histology: an immunohistochemical study of 117 cases. Modern Pathology, 2010, 23, 567-573.	2.9	156
69	Comprehensive Analysis of Genetic Ancestry and Its Molecular Correlates in Cancer. Cancer Cell, 2020, 37, 639-654.e6.	7.7	151
70	Human sebaceous tumors harbor inactivating mutations in LEF1. Nature Medicine, 2006, 12, 395-397.	15.2	149
71	Complete Loss of PTEN Protein Expression Correlates with Shorter Time to Brain Metastasis and Survival in Stage IIIB/C Melanoma Patients with <i>BRAF</i> V600 Mutations. Clinical Cancer Research, 2014, 20, 5527-5536.	3.2	145
72	PanCancer insights from The Cancer Genome Atlas: the pathologist's perspective. Journal of Pathology, 2018, 244, 512-524.	2.1	144

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73	Immuno-genomic landscape of osteosarcoma. Nature Communications, 2020, 11, 1008.	5.8	143
74	The path to a better biomarker: application of a risk management framework for the implementation of PD‣1 and TILs as immunoâ€oncology biomarkers in breast cancer clinical trials and daily practice. Journal of Pathology, 2020, 250, 667-684.	2.1	142
75	CXCR4/CXCL12 Mediate Autocrine Cell- Cycle Progression in NF1-Associated Malignant Peripheral Nerve Sheath Tumors. Cell, 2013, 152, 1077-1090.	13.5	139
76	Sarcoma Epidemiology and Etiology: Potential Environmental and Genetic Factors. Surgical Clinics of North America, 2008, 88, 451-481.	0.5	138
77	Beyond BRAF V600 : Clinical Mutation Panel Testing by Next-Generation Sequencing in Advanced Melanoma. Journal of Investigative Dermatology, 2015, 135, 508-515.	0.3	138
78	Single-cell dissection of intratumoral heterogeneity and lineage diversity in metastatic gastric adenocarcinoma. Nature Medicine, 2021, 27, 141-151.	15.2	134
79	Primary Cutaneous Apocrine Carcinoma. American Journal of Surgical Pathology, 2008, 32, 682-690.	2.1	133
80	Ewing's Sarcoma: Standard and Experimental Treatment Options. Current Treatment Options in Oncology, 2009, 10, 126-140.	1.3	127
81	New Perspectives for Staging and Prognosis in Soft Tissue Sarcoma. Annals of Surgical Oncology, 2008, 15, 2739-2748.	0.7	126
82	Sebaceous neoplasia and the Muir–Torre syndrome: important connections with clinical implications. Histopathology, 2010, 56, 133-147.	1.6	125
83	The 2021 WHO Classification of Tumors of the Thymus and Mediastinum: What Is New in Thymic Epithelial, Germ Cell, and Mesenchymal Tumors?. Journal of Thoracic Oncology, 2022, 17, 200-213.	0.5	124
84	BCOR–CCNB3 fusions are frequent in undifferentiated sarcomas of male children. Modern Pathology, 2015, 28, 575-586.	2.9	122
85	Use of p63 expression in distinguishing primary and metastatic cutaneous adnexal neoplasms from metastatic adenocarcinoma to skin. Journal of Cutaneous Pathology, 2007, 34, 474-480.	0.7	120
86	Genomic and immune heterogeneity are associated with differential responses to therapy in melanoma. Npj Genomic Medicine, 2017, 2, .	1.7	120
87	Circulating Tumor DNA Analysis in Patients With Cancer: American Society of Clinical Oncology and College of American Pathologists Joint Review. Archives of Pathology and Laboratory Medicine, 2018, 142, 1242-1253.	1.2	120
88	Pilomatrix carcinomas contain mutations in CTNNB1, the gene encoding beta-catenin. Journal of Cutaneous Pathology, 2005, 32, 148-157.	0.7	118
89	Rad51 overexpression contributes to chemoresistance in human soft tissue sarcoma cells: a role for p53/activator protein 2 transcriptional regulation. Molecular Cancer Therapeutics, 2007, 6, 1650-1660.	1.9	116
90	Long-Term Outcomes in Patients with Radiation-Associated Angiosarcomas of the Breast Following Surgery and Radiotherapy for Breast Cancer. Annals of Surgical Oncology, 2013, 20, 1267-1274.	0.7	116

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91	Correlative Analyses of the SARC028 Trial Reveal an Association Between Sarcoma-Associated Immune Infiltrate and Response to Pembrolizumab. Clinical Cancer Research, 2020, 26, 1258-1266.	3.2	115
92	Interleukin-6 blockade abrogates immunotherapy toxicity and promotes tumor immunity. Cancer Cell, 2022, 40, 509-523.e6.	7.7	115
93	Complete response of stage IV anal mucosal melanoma expressing KIT Val560Asp to the multikinase inhibitor sorafenib. Nature Clinical Practice Oncology, 2008, 5, 737-740.	4.3	111
94	Expression of ERG, an Ets family transcription factor, identifies ERG-rearranged Ewing sarcoma. Modern Pathology, 2012, 25, 1378-1383.	2.9	111
95	Primary Vascular Tumors and Tumor-like Lesions of the Kidney: A Clinicopathologic Analysis of 25 Cases. American Journal of Surgical Pathology, 2010, 34, 942-949.	2.1	109
96	Staging Soft Tissue Sarcoma: Evolution and Change. Ca-A Cancer Journal for Clinicians, 2006, 56, 282-291.	157.7	107
97	Can MDM2 analytical tests performed on core needle biopsy be relied upon to diagnose well-differentiated liposarcoma?. Modern Pathology, 2010, 23, 1301-1306.	2.9	107
98	Primitive Nonneural Granular Cell Tumors of Skin. American Journal of Surgical Pathology, 2005, 29, 927-934.	2.1	106
99	Novel algorithmic approach predicts tumor mutation load and correlates with immunotherapy clinical outcomes using a defined gene mutation set. BMC Medicine, 2016, 14, 168.	2.3	106
100	Pitfalls in assessing stromal tumor infiltrating lymphocytes (sTILs) in breast cancer. Npj Breast Cancer, 2020, 6, 17.	2.3	106
101	MiR-155 Is a Liposarcoma Oncogene That Targets Casein Kinase-1α and Enhances β-Catenin Signaling. Cancer Research, 2012, 72, 1751-1762.	0.4	104
102	Wt1 ablation and Igf2 upregulation in mice result in Wilms tumors with elevated ERK1/2 phosphorylation. Journal of Clinical Investigation, 2011, 121, 174-183.	3.9	104
103	Site and Tumor Type Predicts DNA Mismatch Repair Status in Cutaneous Sebaceous Neoplasia. American Journal of Surgical Pathology, 2008, 32, 936-942.	2.1	103
104	NKD2, a negative regulator of Wnt signaling, suppresses tumor growth and metastasis in osteosarcoma. Oncogene, 2015, 34, 5069-5079.	2.6	102
105	Evaluation of response after neoadjuvant treatment in soft tissue sarcomas; the European Organization for Research and Treatment of Cancer–Soft Tissue and Bone Sarcoma Group (EORTC–STBSC) recommendations for pathological examination and reporting. European Journal of Cancer. 2016. 53. 84-95.	1.3	99
106	Gauging NOTCH1 Activation in Cancer Using Immunohistochemistry. PLoS ONE, 2013, 8, e67306.	1.1	98
107	Malignant peripheral nerve sheath tumour (MPNST): the clinical implications of cellular signalling pathways. Expert Reviews in Molecular Medicine, 2009, 11, e30.	1.6	97
108	The role of chemotherapy in advanced solitary fibrous tumors: a retrospective analysis. Clinical Sarcoma Research, 2013, 3, 7.	2.3	96

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109	Locoregional Disease Patterns in Well-Differentiated and Dedifferentiated Retroperitoneal Liposarcoma: Implications for the Extent of Resection?. Annals of Surgical Oncology, 2014, 21, 2136-2143.	0.7	96
110	Noncontact measurement of elasticity for the detection of soft-tissue tumors using phase-sensitive optical coherence tomography combined with a focused air-puff system. Optics Letters, 2012, 37, 5184.	1.7	95
111	Metastatic hidradenocarcinoma with demonstration of Her-2/neu gene amplification by fluorescence in situ hybridization: potential treatment implications. Journal of Cutaneous Pathology, 2007, 34, 49-54.	0.7	94
112	Pleomorphic liposarcoma. Cancer, 2011, 117, 5359-5369.	2.0	92
113	Angiogenesis-Promoting Gene Patterns in Alveolar Soft Part Sarcoma. Clinical Cancer Research, 2007, 13, 7314-7321.	3.2	90
114	Localized and metastatic myxoid/round cell liposarcoma. Cancer, 2013, 119, 1868-1877.	2.0	90
115	Nuclear Î ² -catenin localization and mutation of the CTNNB1 gene: a context-dependent association. Modern Pathology, 2018, 31, 1553-1559.	2.9	90
116	Report on computational assessment of Tumor Infiltrating Lymphocytes from the International Immuno-Oncology Biomarker Working Group. Npj Breast Cancer, 2020, 6, 16.	2.3	90
117	Patterns of recurrence and survival in sporadic, neurofibromatosis Type 1–associated, and radiation-associated malignant peripheral nerve sheath tumors. Journal of Neurosurgery, 2017, 126, 319-329.	0.9	89
118	Cutaneous Digital Papillary Adenocarcinoma. American Journal of Surgical Pathology, 2012, 36, 1883-1891.	2.1	88
119	Clinical characteristics and outcomes with specific <i>BRAF</i> and <i>NRAS</i> mutations in patients with metastatic melanoma. Cancer, 2013, 119, 3821-3829.	2.0	87
120	Activated MET Is a Molecular Prognosticator and Potential Therapeutic Target for Malignant Peripheral Nerve Sheath Tumors. Clinical Cancer Research, 2011, 17, 3943-3955.	3.2	86
121	Molecular Diagnosis of Sarcomas: Chromosomal Translocations in Sarcomas. Archives of Pathology and Laboratory Medicine, 2006, 130, 1199-1207.	1.2	85
122	Galectin-3 Expression Is Associated with Tumor Progression and Pattern of Sun Exposure in Melanoma. Clinical Cancer Research, 2006, 12, 6709-6715.	3.2	84
123	Autophagic Survival in Resistance to Histone Deacetylase Inhibitors: Novel Strategies to Treat Malignant Peripheral Nerve Sheath Tumors. Cancer Research, 2011, 71, 185-196.	0.4	84
124	SAR405838: A Novel and Potent Inhibitor of the MDM2:p53 Axis for the Treatment of Dedifferentiated Liposarcoma. Clinical Cancer Research, 2016, 22, 1150-1160.	3.2	84
125	Dual targeting of AKT and mammalian target of rapamycin: A potential therapeutic approach for malignant peripheral nerve sheath tumor. Molecular Cancer Therapeutics, 2009, 8, 1157-1168.	1.9	83

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127	Involvement of the PI3K/Akt pathway in myxoid/round cell liposarcoma. Modern Pathology, 2012, 25, 212-221.	2.9	81
128	ERG and FLI1 protein expression in epithelioid sarcoma. Modern Pathology, 2014, 27, 496-501.	2.9	81
129	USP6 activation in nodular fasciitis by promoter-swapping gene fusions. Modern Pathology, 2017, 30, 1577-1588.	2.9	79
130	Insights into developmental mechanisms and cancers in the mammalian intestine derived from serial analysis of gene expression and study of the hepatoma-derived growth factor (HDCF). Development (Cambridge), 2005, 132, 415-427.	1.2	77
131	ROR2 is a novel prognostic biomarker and a potential therapeutic target in leiomyosarcoma and gastrointestinal stromal tumour. Journal of Pathology, 2012, 227, 223-233.	2.1	77
132	PRKCI promotes immune suppression in ovarian cancer. Genes and Development, 2017, 31, 1109-1121.	2.7	75
133	Uterine Tumors With Neuroectodermal Differentiation. American Journal of Surgical Pathology, 2008, 32, 219-228.	2.1	74
134	Genetic aberrations of gastrointestinal stromal tumors. Cancer, 2008, 113, 1532-1543.	2.0	72
135	FUS rearrangements are rare in â€~pure' sclerosing epithelioid fibrosarcoma. Modern Pathology, 2012, 25, 846-853.	2.9	72
136	An experimental model for the study of well-differentiated and dedifferentiated liposarcoma; deregulation of targetable tyrosine kinase receptors. Laboratory Investigation, 2011, 91, 392-403.	1.7	71
137	Cross Species Genomic Analysis Identifies a Mouse Model as Undifferentiated Pleomorphic Sarcoma/Malignant Fibrous Histiocytoma. PLoS ONE, 2009, 4, e8075.	1.1	71
138	Clinical, histopathologic, molecular and therapeutic findings in a large kindred with gastrointestinal stromal tumor. International Journal of Cancer, 2008, 122, 711-718.	2.3	70
139	Detection of MDM2 gene amplification or protein expression distinguishes sclerosing mesenteritis and retroperitoneal fibrosis from inflammatory well-differentiated liposarcoma. Modern Pathology, 2009, 22, 66-70.	2.9	70
140	Hotspot Mutation Panel Testing Reveals Clonal Evolution in a Study of 265 Paired Primary and Metastatic Tumors. Clinical Cancer Research, 2015, 21, 2644-2651.	3.2	70
141	A Summary of the Inaugural WHO Classification of Pediatric Tumors: Transitioning from the Optical into the Molecular Era. Cancer Discovery, 2022, 12, 331-355.	7.7	70
142	Synovial Sarcoma: Advances in Diagnosis and Treatment Identification of New Biologic Targets to Improve Multimodal Therapy. Annals of Surgical Oncology, 2017, 24, 2145-2154.	0.7	69
143	EWSR1 fusion proteins mediate PAX7 expression in Ewing sarcoma. Modern Pathology, 2017, 30, 1312-1320.	2.9	69
144	Phase II study of neoadjuvant checkpoint blockade in patients with surgically resectable undifferentiated pleomorphic sarcoma and dedifferentiated liposarcoma. BMC Cancer, 2018, 18, 913.	1.1	69

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145	Multimodality Treatment of Desmoplastic Small Round Cell Tumor: Chemotherapy and Complete Cytoreductive Surgery Improve Patient Survival. Clinical Cancer Research, 2018, 24, 4865-4873.	3.2	68
146	Molecular characterization of epithelioid haemangioendotheliomas identifies novel <i><scp>WWTR</scp>1</i> – <i><scp>CAMTA</scp>1</i> fusion variants. Histopathology, 2015, 67, 699-708.	1.6	67
147	Myxofibrosarcoma. Surgical Oncology Clinics of North America, 2016, 25, 775-788.	0.6	67
148	Detection of myxoid liposarcoma-associated FUS–DDIT3 rearrangement variants including a newly identified breakpoint using an optimized RT-PCR assay. Modern Pathology, 2010, 23, 1307-1315.	2.9	66
149	Comparison of Cancer Prevalence in Patients With Neurofibromatosis Type 1 at an Academic Cancer Center vs in the General Population From 1985 to 2020. JAMA Network Open, 2021, 4, e210945.	2.8	66
150	Liposarcoma in children and young adults: A multiâ€institutional experience. Pediatric Blood and Cancer, 2011, 57, 1142-1146.	0.8	65
151	Outcome of Locally Recurrent and Metastatic Angiosarcoma. Annals of Surgical Oncology, 2009, 16, 2502-2509.	0.7	64
152	Complete Soft Tissue Sarcoma Resection is a Viable Treatment Option for Select Elderly Patients. Annals of Surgical Oncology, 2009, 16, 2579-2586.	0.7	64
153	Targeting the <scp>N</scp> otch pathway: A potential therapeutic approach for desmoid tumors. Cancer, 2015, 121, 4088-4096.	2.0	64
154	NY-ESO-1 (CTAG1B) expression in mesenchymal tumors. Modern Pathology, 2015, 28, 587-595.	2.9	64
155	A Keratinocyte Hypermotility/Growth-Arrest Response Involving Laminin 5 and p16INK4A Activated in Wound Healing and Senescence. American Journal of Pathology, 2006, 168, 1821-1837.	1.9	63
156	Clinical responses to selumetinib (AZD6244; ARRYâ€142886)â€based combination therapy stratified by gene mutations in patients with metastatic melanoma. Cancer, 2013, 119, 799-805.	2.0	63
157	EZH2-miR-30d-KPNB1 pathway regulates malignant peripheral nerve sheath tumour cell survival and tumourigenesis. Journal of Pathology, 2014, 232, 308-318.	2.1	62
158	p53 ls a Master Regulator of Proteostasis in SMARCB1-Deficient Malignant Rhabdoid Tumors. Cancer Cell, 2019, 35, 204-220.e9.	7.7	62
159	New Strategies in Melanoma: Molecular Testing in Advanced Disease. Clinical Cancer Research, 2012, 18, 1195-1200.	3.2	61
160	Uterine Leiomyosarcoma Management, Outcome, and Associated Molecular Biomarkers: A Single Institution's Experience. Annals of Surgical Oncology, 2013, 20, 2364-2372.	0.7	61
161	Overexpressed PRAME is a potential immunotherapy target in sarcoma subtypes. Clinical Sarcoma Research, 2017, 7, 11.	2.3	61
162	R132C IDH1 Mutations Are Found in Spindle Cell Hemangiomas and Not in Other Vascular Tumors or Malformations. American Journal of Pathology, 2013, 182, 1494-1500.	1.9	60

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163	Mechanisms of resistance to imatinib and sunitinib in gastrointestinal stromal tumor. Cancer Chemotherapy and Pharmacology, 2011, 67, 15-24.	1.1	59
164	Diagnosis, Management, and Outcome of Patients with Dedifferentiated Liposarcoma Systemic Metastasis. Annals of Surgical Oncology, 2011, 18, 3762-3770.	0.7	58
165	Phase I Study of the Combination of Sorafenib and Temsirolimus in Patients with Metastatic Melanoma. Clinical Cancer Research, 2012, 18, 1120-1128.	3.2	57
166	Angiosarcoma. American Journal of Dermatopathology, 2013, 35, 432-437.	0.3	57
167	Dual Roles of RNF2 in Melanoma Progression. Cancer Discovery, 2015, 5, 1314-1327.	7.7	57
168	Role of chemotherapy in dedifferentiated liposarcoma of the retroperitoneum: defining the benefit and challenges of the standard. Scientific Reports, 2017, 7, 11836.	1.6	57
169	Integrated Proteomics and Genomics Analysis Reveals a Novel Mesenchymal to Epithelial Reverting Transition in Leiomyosarcoma through Regulation of Slug. Molecular and Cellular Proteomics, 2010, 9, 2405-2413.	2.5	56
170	Expression of ERG, an Ets family transcription factor, distinguishes cutaneous angiosarcoma from histological mimics. Histopathology, 2012, 61, 989-991.	1.6	56
171	A Preexisting Rare <i>PIK3CA</i> E545K Subpopulation Confers Clinical Resistance to MEK plus CDK4/6 Inhibition in <i>NRAS</i> Melanoma and Is Dependent on S6K1 Signaling. Cancer Discovery, 2018, 8, 556-567.	7.7	55
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