

# David Capper

## List of Publications by Year in descending order

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Version: 2024-02-01

268  
papers

30,590  
citations

5268

83  
h-index

5394

164  
g-index

279  
all docs

279  
docs citations

279  
times ranked

27814  
citing authors

#	ARTICLE	IF	CITATIONS
1	Artificial intelligence and pathology: From principles to practice and future applications in histomorphology and molecular profiling. <i>Seminars in Cancer Biology</i> , 2022, 84, 129-143.	9.6	41
2	Mucosal melanomas of different anatomic sites share a common global <sc>DNA</sc> methylation profile with cutaneous melanoma but show locationâ€dependent patterns of genetic and epigenetic alterations. <i>Journal of Pathology</i> , 2022, 256, 61-70.	4.5	12
3	Comparative investigation of cell cycle and immunomodulatory genes in mucosal and cutaneous melanomas: Preliminary data suggest a potential promising clinical role for p16 and the PD-1/PD-L1 axis. <i>Pathology Research and Practice</i> , 2022, 229, 153689.	2.3	1
4	Machine learning models predict the primary sites of head and neck squamous cell carcinoma metastases based on <sc>DNA</sc> methylation. <i>Journal of Pathology</i> , 2022, 256, 378-387.	4.5	19
5	Pleomorphic xanthoastrocytoma is a heterogeneous entity with pTERT mutations prognosticating shorter survival. <i>Acta Neuropathologica Communications</i> , 2022, 10, 5.	5.2	12
6	Lowâ€grade diffusely infiltrative tumour (LGDIT), SMARCB1â€mutant: A clinical and histopathological distinct entity showing epigenetic similarity with ATRTâ€MYC. <i>Neuropathology and Applied Neurobiology</i> , 2022, 48, .	3.2	5
7	A pan-tissue DNA methylation atlas enables in silico decomposition of human tissue methylomes at cell-type resolution. <i>Nature Methods</i> , 2022, 19, 296-306.	19.0	46
8	DNA methylation reveals distinct cells of origin for pancreatic neuroendocrine carcinomas and pancreatic neuroendocrine tumors. <i>Genome Medicine</i> , 2022, 14, 24.	8.2	12
9	Comprehensive profiling of myxopapillary ependymomas identifies a distinct molecular subtype with relapsing disease. <i>Neuro-Oncology</i> , 2022, 24, 1689-1699.	1.2	11
10	Oligosarcomas, IDH-mutant are distinct and aggressive. <i>Acta Neuropathologica</i> , 2022, 143, 263-281.	7.7	18
11	DNA methylation-based machine learning classification distinguishes pleural mesothelioma from chronic pleuritis, pleural carcinosis, and pleomorphic lung carcinomas. <i>Lung Cancer</i> , 2022, 170, 105-113.	2.0	3
12	Management of pineal region tumors in a pediatric case series. <i>Neurosurgical Review</i> , 2021, 44, 1417-1427.	2.4	25
13	Accurate calling of <i>KIAA1549â€BRAF</i> fusions from DNA of human brain tumours using methylation arrayâ€based copy number and gene panel sequencing data. <i>Neuropathology and Applied Neurobiology</i> , 2021, 47, 406-414.	3.2	12
14	DNA methylation based glioblastoma subclassification is related to tumoral T-cell infiltration and patient survival. <i>Neuro-Oncology</i> , 2021, 23, 240-250.	1.2	31
15	TERT promoter mutation and chromosome 6 loss define a high-risk subtype of ependymoma evolving from posterior fossa subependymoma. <i>Acta Neuropathologica</i> , 2021, 141, 959-970.	7.7	16
16	High-grade astrocytoma with piloid features (HGAP): the CharitÃ© experience with a new central nervous system tumor entity. <i>Journal of Neuro-Oncology</i> , 2021, 153, 109-120.	2.9	35
17	Cross-Species Genomics Reveals Oncogenic Dependencies in ZFTA/C11orf95 Fusionâ€Positive Supratentorial Ependymomas. <i>Cancer Discovery</i> , 2021, 11, 2230-2247.	9.4	39
18	Molecular characterisation of sporadic endolymphatic sac tumours and comparison to von Hippelâ€Lindau diseaseâ€related tumours. <i>Neuropathology and Applied Neurobiology</i> , 2021, 47, 756-767.	3.2	2

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19	Therapeutic implications of improved molecular diagnostics for rare CNS embryonal tumor entities: results of an international, retrospective study. <i>Neuro-Oncology</i> , 2021, 23, 1597-1611.	1.2	22
20	IDH2 R172 Mutations Across Poorly Differentiated Sinonasal Tract Malignancies. <i>American Journal of Surgical Pathology</i> , 2021, 45, 1190-1204.	3.7	26
21	Recurrent fusions in PLAGL1 define a distinct subset of pediatric-type supratentorial neuroepithelial tumors. <i>Acta Neuropathologica</i> , 2021, 142, 827-839.	7.7	33
22	The Pediatric Precision Oncology INFORM Registry: Clinical Outcome and Benefit for Patients with Very High-Evidence Targets. <i>Cancer Discovery</i> , 2021, 11, 2764-2779.	9.4	110
23	GOPC:ROS1 and other ROS1 fusions represent a rare but recurrent drug target in a variety of glioma types. <i>Acta Neuropathologica</i> , 2021, 142, 1065-1069.	7.7	16
24	Molecular profiling of pediatric meningiomas shows tumor characteristics distinct from adult meningiomas. <i>Acta Neuropathologica</i> , 2021, 142, 873-886.	7.7	12
25	Sarcoma classification by DNA methylation profiling. <i>Nature Communications</i> , 2021, 12, 498.	12.8	237
26	Intraoperative DNA methylation classification of brain tumors impacts neurosurgical strategy. <i>Neuro-Oncology Advances</i> , 2021, 3, vdab149.	0.7	23
27	Genetic and epigenetic characterization of posterior pituitary tumors. <i>Acta Neuropathologica</i> , 2021, 142, 1025-1043.	7.7	7
28	DNA methylation profiling identifies two distinct subgroups in breast cancers with low hormone receptor expression, mainly associated with HER2 amplification status. <i>Clinical Epigenetics</i> , 2021, 13, 184.	4.1	2
29	FGFR3 overexpression is a useful detection tool for FGFR3 fusions and sequence variations in glioma. <i>Neuro-Oncology Practice</i> , 2021, 8, 209-221.	1.6	7
30	Isomorphic diffuse glioma is a morphologically and molecularly distinct tumour entity with recurrent gene fusions of MYBL1 or MYB and a benign disease course. <i>Acta Neuropathologica</i> , 2020, 139, 193-209.	7.7	83
31	Reflection of neuroblastoma intratumor heterogeneity in the new OHC&NB1 disease model. <i>International Journal of Cancer</i> , 2020, 146, 1031-1041.	5.1	9
32	Machine learning workflows to estimate class probabilities for precision cancer diagnostics on DNA methylation microarray data. <i>Nature Protocols</i> , 2020, 15, 479-512.	12.0	89
33	DNA methylation-based profiling for paediatric CNS tumour diagnosis and treatment: a population-based study. <i>The Lancet Child and Adolescent Health</i> , 2020, 4, 121-130.	5.6	55
34	High density DNA methylation array is a reliable alternative for PCR-based analysis of the MGMT promoter methylation status in glioblastoma. <i>Pathology Research and Practice</i> , 2020, 216, 152728.	2.3	8
35	Response to trametinib treatment in progressive pediatric low-grade glioma patients. <i>Journal of Neuro-Oncology</i> , 2020, 149, 499-510.	2.9	68
36	Infratentorial IDH-mutant astrocytoma is a distinct subtype. <i>Acta Neuropathologica</i> , 2020, 140, 569-581.	7.7	45

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37	Molecular characterization of CNS paragangliomas identifies cauda equina paragangliomas as a distinct tumor entity. <i>Acta Neuropathologica</i> , 2020, 140, 893-906.	7.7	19
38	Malignant transformation and genetic alterations are uncoupled in early colorectal cancer progression. <i>BMC Biology</i> , 2020, 18, 116.	3.8	16
39	Acquired resistance to DZNep-mediated apoptosis is associated with copy number gains of AHCY in a B-cell lymphoma model. <i>BMC Cancer</i> , 2020, 20, 427.	2.6	3
40	An 8-Year-Old Girl with Posterior Fossa Mass. <i>Brain Pathology</i> , 2020, 30, 713-714.	4.1	1
41	cIMPACT-NOW update 7: advancing the molecular classification of ependymal tumors. <i>Brain Pathology</i> , 2020, 30, 863-866.	4.1	168
42	Genome-wide DNA methylation profiles distinguish silent from non-silent ACTH adenomas. <i>Acta Neuropathologica</i> , 2020, 140, 95-97.	7.7	7
43	Phase 1b/2a study of galunisertib, a small molecule inhibitor of transforming growth factor-beta receptor I, in combination with standard temozolomide-based radiochemotherapy in patients with newly diagnosed malignant glioma. <i>Investigational New Drugs</i> , 2020, 38, 1570-1579.	2.6	70
44	Invited Review: DNA methylation-based classification of paediatric brain tumours. <i>Neuropathology and Applied Neurobiology</i> , 2020, 46, 28-47.	3.2	33
45	cIMPACT-NOW update 6: new entity and diagnostic principle recommendations of the cIMPACT-Utrecht meeting on future CNS tumor classification and grading. <i>Brain Pathology</i> , 2020, 30, 844-856.	4.1	363
46	Infant High-Grade Gliomas Comprise Multiple Subgroups Characterized by Novel Targetable Gene Fusions and Favorable Outcomes. <i>Cancer Discovery</i> , 2020, 10, 942-963.	9.4	157
47	Review: Challenges in the histopathological classification of ganglioglioma and DNT: microscopic agreement studies and a preliminary genotype-phenotype analysis. <i>Neuropathology and Applied Neurobiology</i> , 2019, 45, 95-107.	3.2	46
48	DNA methylation signature is prognostic of choroid plexus tumor aggressiveness. <i>Clinical Epigenetics</i> , 2019, 11, 117.	4.1	21
49	New therapeutic target for pediatric anaplastic ependymoma control: study of anti-tumor activity by a Kunitz-type molecule, Amblyomin-X. <i>Scientific Reports</i> , 2019, 9, 9973.	3.3	6
50	Extent of Resection, MGMT Promoter Methylation Status and Tumor Location Independently Predict Progression-Free Survival in Adult Sporadic Pilocytic Astrocytoma. <i>Cancers</i> , 2019, 11, 1072.	3.7	16
51	Tumors diagnosed as cerebellar glioblastoma comprise distinct molecular entities. <i>Acta Neuropathologica Communications</i> , 2019, 7, 163.	5.2	37
52	YAP1 subgroup supratentorial ependymoma requires TEAD and nuclear factor I-mediated transcriptional programmes for tumorigenesis. <i>Nature Communications</i> , 2019, 10, 3914.	12.8	65
53	Machine learning analysis of DNA methylation profiles distinguishes primary lung squamous cell carcinomas from head and neck metastases. <i>Science Translational Medicine</i> , 2019, 11, .	12.4	100
54	Location-Dependent Patient Outcome and Recurrence Patterns in IDH1-Wildtype Glioblastoma. <i>Cancers</i> , 2019, 11, 122.	3.7	25

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55	DNA methylation profiling reliably distinguishes pulmonary enteric adenocarcinoma from metastatic colorectal cancer. <i>Modern Pathology</i> , 2019, 32, 855-865.	5.5	36
56	Rosette-forming glioneuronal tumors share a distinct DNA methylation profile and mutations in FGFR1, with recurrent co-mutation of PIK3CA and NF1. <i>Acta Neuropathologica</i> , 2019, 138, 497-504.	7.7	57
57	Lack of H3K27 trimethylation is associated with 1p/19q codeletion in diffuse gliomas. <i>Acta Neuropathologica</i> , 2019, 138, 331-334.	7.7	22
58	Methylation array profiling of adult brain tumours: diagnostic outcomes in a large, single centre. <i>Acta Neuropathologica Communications</i> , 2019, 7, 24.	5.2	101
59	Papillary glioneuronal tumor (PGNT) exhibits a characteristic methylation profile and fusions involving PRKCA. <i>Acta Neuropathologica</i> , 2019, 137, 837-846.	7.7	43
60	RBTT-06. TESSA JOWELL BRAIN MATRIX STUDY: A BRITISH FEASIBILITY STUDY OF MOLECULAR STRATIFICATION AND TARGETED THERAPY TO OPTIMIZE THE CLINICAL MANAGEMENT OF PATIENTS WITH GLIOMA. <i>Neuro-Oncology</i> , 2019, 21, vi219-vi220.	1.2	1
61	MRI Features of Histologically Diagnosed Supratentorial Primitive Neuroectodermal Tumors and Pineoblastomas in Correlation with Molecular Diagnoses and Outcomes: A Report from the Children's Oncology Group ACNS0332 Trial. <i>American Journal of Neuroradiology</i> , 2019, 40, 1796-1803.	2.4	11
62	Actively personalized vaccination trial for newly diagnosed glioblastoma. <i>Nature</i> , 2019, 565, 240-245.	27.8	637
63	Diagnostics of pediatric supratentorial RELA ependymomas: integration of information from histopathology, genetics, DNA methylation and imaging. <i>Brain Pathology</i> , 2019, 29, 325-335.	4.1	55
64	Integrated molecular characterization of IDH-mutant glioblastomas. <i>Neuropathology and Applied Neurobiology</i> , 2019, 45, 108-118.	3.2	68
65	The Senescence-associated Secretory Phenotype Mediates Oncogene-induced Senescence in Pediatric Pilocytic Astrocytoma. <i>Clinical Cancer Research</i> , 2019, 25, 1851-1866.	7.0	55
66	Multicentric malignant glioma with striking morphologic heterogeneity and early and extensive metastatic spread to the bone. , 2019, 38, 261-268.		0
67	The miR-139-5p regulates proliferation of supratentorial paediatric low-grade gliomas by targeting the PI3K/AKT/mTORC1 signalling. <i>Neuropathology and Applied Neurobiology</i> , 2018, 44, 687-706.	3.2	31
68	cIMPACT-NOW update 2: diagnostic clarifications for diffuse midline glioma, H3 K27M-mutant and diffuse astrocytoma/anaplastic astrocytoma, IDH-mutant. <i>Acta Neuropathologica</i> , 2018, 135, 639-642.	7.7	281
69	Novel, improved grading system(s) for IDH-mutant astrocytic gliomas. <i>Acta Neuropathologica</i> , 2018, 136, 153-166.	7.7	298
70	Preoperative assessment of haemostasis in patients undergoing stereotactic brain biopsy. <i>Journal of Clinical Neuroscience</i> , 2018, 53, 112-116.	1.5	13
71	Bevacizumab plus hypofractionated radiotherapy versus radiotherapy alone in elderly patients with glioblastoma: the randomized, open-label, phase II ARTE trial. <i>Annals of Oncology</i> , 2018, 29, 1423-1430.	1.2	65
72	cIMPACT-NOW update 1: Not Otherwise Specified (NOS) and Not Elsewhere Classified (NEC). <i>Acta Neuropathologica</i> , 2018, 135, 481-484.	7.7	145

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73	Feasibility of real-time molecular profiling for patients with newly diagnosed glioblastoma without MGMT promoter hypermethylation in the NCT Neuro Master Match (N2M2) pilot study. <i>Neuro-Oncology</i> , 2018, 20, 826-837.	1.2	32
74	Sellar Region Atypical Teratoid/Rhabdoid Tumors (ATRT) in Adults Display DNA Methylation Profiles of the ATRT-MYC Subgroup. <i>American Journal of Surgical Pathology</i> , 2018, 42, 506-511.	3.7	43
75	DNA methylation-based reclassification of olfactory neuroblastoma. <i>Acta Neuropathologica</i> , 2018, 136, 255-271.	7.7	59
76	DNA methylation-based classification of central nervous system tumours. <i>Nature</i> , 2018, 555, 469-474.	27.8	1,872
77	Anaplastic astrocytoma with piloid features, a novel molecular class of IDH wildtype glioma with recurrent MAPK pathway, CDKN2A/B and ATRX alterations. <i>Acta Neuropathologica</i> , 2018, 136, 273-291.	7.7	190
78	CASP9 germline mutation in a family with multiple brain tumors. <i>Brain Pathology</i> , 2018, 28, 94-102.	4.1	11
79	Glial papillary tumour of the spinal cord with SMARCB1/INI1 loss and favourable long-term outcome. <i>Neuropathology and Applied Neurobiology</i> , 2018, 44, 229-232.	3.2	3
80	Radiomic subtyping improves disease stratification beyond key molecular, clinical, and standard imaging characteristics in patients with glioblastoma. <i>Neuro-Oncology</i> , 2018, 20, 848-857.	1.2	170
81	The case for DNA methylation based molecular profiling to improve diagnostic accuracy for central nervous system embryonal tumors (not otherwise specified) in adults. <i>Journal of Clinical Neuroscience</i> , 2018, 47, 163-167.	1.5	8
82	Epithelioid glioblastomas stratify into established diagnostic subsets upon integrated molecular analysis. <i>Brain Pathology</i> , 2018, 28, 656-662.	4.1	89
83	WHO 2016 Classification of gliomas. <i>Neuropathology and Applied Neurobiology</i> , 2018, 44, 139-150.	3.2	612
84	Extensive Molecular and Clinical Heterogeneity in Patients With Histologically Diagnosed CNS-PNET Treated as a Single Entity: A Report From the Children's Oncology Group Randomized ACNS0332 Trial. <i>Journal of Clinical Oncology</i> , 2018, 36, 3388-3395.	1.6	58
85	Modern Principles of CNS Tumor Classification. , 2018, , 117-129.		0
86	Distribution of EGFR amplification, combined chromosome 7 gain and chromosome 10 loss, and TERT promoter mutation in brain tumors and their potential for the reclassification of IDHwt astrocytoma to glioblastoma. <i>Acta Neuropathologica</i> , 2018, 136, 793-803.	7.7	195
87	EMBR-01. MOLECULAR AND CLINICAL HETEROGENEITY IN HISTOLOGICALLY-DIAGNOSED CNS-PNET PATIENTS PROSPECTIVELY TREATED AS A SINGLE ENTITY: A REPORT FROM THE CHILDREN'S ONCOLOGY GROUP ACNS0332 TRIAL. <i>Neuro-Oncology</i> , 2018, 20, i68-i69.	1.2	0
88	Satisfying your neuro-oncologist: a fast approach to routine molecular glioma diagnostics. <i>Neuro-Oncology</i> , 2018, 20, 1682-1683.	1.2	8
89	Molecularly defined diffuse leptomeningeal glioneuronal tumor (DLGNT) comprises two subgroups with distinct clinical and genetic features. <i>Acta Neuropathologica</i> , 2018, 136, 239-253.	7.7	118
90	Molecular, Pathological, Radiological, and Immune Profiling of Non-brainstem Pediatric High-Grade Glioma from the HERBY Phase II Randomized Trial. <i>Cancer Cell</i> , 2018, 33, 829-842.e5.	16.8	140

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91	Practical implementation of DNA methylation and copy-number-based CNS tumor diagnostics: the Heidelberg experience. <i>Acta Neuropathologica</i> , 2018, 136, 181-210.	7.7	308
92	DNA methylation-based classification of ependymomas in adulthood: implications for diagnosis and treatment. <i>Neuro-Oncology</i> , 2018, 20, 1616-1624.	1.2	65
93	NFM-11. PEDIATRIC MENINGIOMAS ARE MOLECULARLY DISTINCT FROM ADULT COUNTERPARTS. <i>Neuro-Oncology</i> , 2018, 20, i144-i145.	1.2	1
94	Concurrent IDH1 and SMARCB1 Mutations in Pediatric Medulloblastoma: A Case Report. <i>Frontiers in Neurology</i> , 2018, 9, 398.	2.4	10
95	Desmoplastic Infantile Ganglioglioma/Astrocytoma (DIG/DIA) Are Distinct Entities with Frequent BRAFV600 Mutations. <i>Molecular Cancer Research</i> , 2018, 16, 1491-1498.	3.4	39
96	Recurrent homozygous deletion of DROSHA and microduplication of PDE4DIP in pineoblastoma. <i>Nature Communications</i> , 2018, 9, 2868.	12.8	54
97	Voxel-wise radiogenomic mapping of tumor location with key molecular alterations in patients with glioma. <i>Neuro-Oncology</i> , 2018, 20, 1517-1524.	1.2	36
98	Pericytes/vessel-associated mural cells (VAMCs) are the major source of key epithelial-mesenchymal transition (EMT) factors SLUG and TWIST in human glioma. <i>Oncotarget</i> , 2018, 9, 24041-24053.	1.8	8
99	Pan-mutant IDH1 inhibitor BAY 1436032 for effective treatment of IDH1 mutant astrocytoma in vivo. <i>Acta Neuropathologica</i> , 2017, 133, 629-644.	7.7	146
100	A 49-year old female with multiple extra-axial tumors. <i>Brain Pathology</i> , 2017, 27, 235-236.	4.1	0
101	H3-/IDH-wild type pediatric glioblastoma is comprised of molecularly and prognostically distinct subtypes with associated oncogenic drivers. <i>Acta Neuropathologica</i> , 2017, 134, 507-516.	7.7	144
102	DNA methylation-based classification and grading system for meningioma: a multicentre, retrospective analysis. <i>Lancet Oncology</i> , The, 2017, 18, 682-694.	10.7	586
103	Gain of 12p encompassing CCND2 is associated with gemistocytic histology in IDH mutant astrocytomas. <i>Acta Neuropathologica</i> , 2017, 133, 325-327.	7.7	12
104	Announcing cIMPACT-NOW: the Consortium to Inform Molecular and Practical Approaches to CNS Tumor Taxonomy. <i>Acta Neuropathologica</i> , 2017, 133, 1-3.	7.7	120
105	Oncolytic H-1 Parvovirus Shows Safety and Signs of Immunogenic Activity in a First Phase I/IIa Glioblastoma Trial. <i>Molecular Therapy</i> , 2017, 25, 2620-2634.	8.2	199
106	The whole-genome landscape of medulloblastoma subtypes. <i>Nature</i> , 2017, 547, 311-317.	27.8	787
107	Molecular Transition of an Adult Low-Grade Brain Tumor to an Atypical Teratoid/Rhabdoid Tumor Over a Time-Course of 14 Years. <i>Journal of Neuropathology and Experimental Neurology</i> , 2017, 76, 655-664.	1.7	13
108	Histone 3.3 hotspot mutations in conventional osteosarcomas: a comprehensive clinical and molecular characterization of six H3F3A mutated cases. <i>Clinical Sarcoma Research</i> , 2017, 7, 9.	2.3	51

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109	Polymorphous low-grade neuroepithelial tumor of the young (PLNTY): an epileptogenic neoplasm with oligodendroglioma-like components, aberrant CD34 expression, and genetic alterations involving the MAP kinase pathway. <i>Acta Neuropathologica</i> , 2017, 133, 417-429.	7.7	172
110	cIMPACTâ€NOW (the consortium to inform molecular and practical approaches to CNS tumor) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 70 27, 851-852.	4.1	63
111	Cribriform neuroepithelial tumor: molecular characterization of a SMARCB1â€deficient nonâ€rhabdoid tumor with favorable longâ€term outcome. <i>Brain Pathology</i> , 2017, 27, 411-418.	4.1	58
112	Biomarker and Histopathology Evaluation of Patients with Recurrent Glioblastoma Treated with Galunisertib, Lomustine, or the Combination of Galunisertib and Lomustine. <i>International Journal of Molecular Sciences</i> , 2017, 18, 995.	4.1	32
113	Effects of soluble CPE on glioma cell migration are associated with mTOR activation and enhanced glucose flux. <i>Oncotarget</i> , 2017, 8, 67567-67591.	1.8	11
114	Establishment and application of a novel patient-derived KIAA1549:BRAF-driven pediatric pilocytic astrocytoma model for preclinical drug testing. <i>Oncotarget</i> , 2017, 8, 11460-11479.	1.8	43
115	Dabrafenib in patients with recurrent, BRAF V600E mutated malignant glioma and leptomeningeal disease. <i>Oncology Reports</i> , 2017, 38, 3291-3296.	2.6	46
116	Pediatric Targeted Therapy: Clinical Feasibility of Personalized Diagnostics in Children with Relapsed and Progressive Tumors. <i>Brain Pathology</i> , 2016, 26, 506-516.	4.1	14
117	<i>LOC283731</i> promoter hypermethylation prognosticates survival after radiochemotherapy in IDH1 wildâ€type glioblastoma patients. <i>International Journal of Cancer</i> , 2016, 139, 424-432.	5.1	18
118	Papillary Tumor of the Pineal Region: A Distinct Molecular Entity. <i>Brain Pathology</i> , 2016, 26, 199-205.	4.1	39
119	Somatic mutations of <i>DICER1</i> and <i>KMT2D</i> are frequent in intraocular medulloepitheliomas. <i>Genes Chromosomes and Cancer</i> , 2016, 55, 418-427.	2.8	34
120	Genetic Alterations in Gliosarcoma and Giant Cell Glioblastoma. <i>Brain Pathology</i> , 2016, 26, 517-522.	4.1	63
121	Spatial transcriptome analysis reveals Notch pathway-associated prognostic markers in IDH1 wild-type glioblastoma involving the subventricular zone. <i>BMC Medicine</i> , 2016, 14, 170.	5.5	31
122	Poorly differentiated chordoma with SMARCB1/INI1 loss: a distinct molecular entity with dismal prognosis. <i>Acta Neuropathologica</i> , 2016, 132, 149-151.	7.7	127
123	Temozolomide chemotherapy versus radiotherapy in high-risk low-grade glioma (EORTC 22033-26033): a randomised, open-label, phase 3 intergroup study. <i>Lancet Oncology</i> , The, 2016, 17, 1521-1532.	10.7	396
124	HG-68COMBINED ALTERATIONS IN MAPK PATHWAY GENES, CDKN2A/B AND ATRX CHARACTERIZE ANAPLASTIC PILOCYTIC ASTROCYTOMA. <i>Neuro-Oncology</i> , 2016, 18, iii63.2-iii63.	1.2	0
125	Next-generation personalised medicine for high-risk paediatric cancer patients â€ The INFORM pilot study. <i>European Journal of Cancer</i> , 2016, 65, 91-101.	2.8	262
126	SHH desmoplastic/nodular medulloblastoma and Gorlin syndrome in the setting of Down syndrome: case report, molecular profiling, and review of the literature. <i>Child's Nervous System</i> , 2016, 32, 2439-2446.	1.1	15



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127	Evidence of H3 K27M mutations in posterior fossa ependymomas. <i>Acta Neuropathologica</i> , 2016, 132, 635-637.	7.7	73
128	Radiogenomics of Glioblastoma: Machine Learning-based Classification of Molecular Characteristics by Using Multiparametric and Multiregional MR Imaging Features. <i>Radiology</i> , 2016, 281, 907-918.	7.3	236
129	Recurrent MET fusion genes represent a drug target in pediatric glioblastoma. <i>Nature Medicine</i> , 2016, 22, 1314-1320.	30.7	183
130	Low-grade epilepsy-associated neuroepithelial tumours – the 2016 WHO classification. <i>Nature Reviews Neurology</i> , 2016, 12, 732-740.	10.1	113
131	P08.48 Combined alterations in MAPK pathway genes, CDKN2A/B and ATRX characterize anaplastic pilocytic astrocytoma. <i>Neuro-Oncology</i> , 2016, 18, iv52-iv52.	1.2	0
132	Chemotherapy with BCNU in recurrent glioma: Analysis of clinical outcome and side effects in chemotherapy-naïve patients. <i>BMC Cancer</i> , 2016, 16, 81.	2.6	51
133	Adamantinomatous and papillary craniopharyngiomas are characterized by distinct epigenomic as well as mutational and transcriptomic profiles. <i>Acta Neuropathologica Communications</i> , 2016, 4, 20.	5.2	136
134	Prognostic value of the extent of resection in supratentorial WHO grade II astrocytomas stratified for IDH1 mutation status: a single-center volumetric analysis. <i>Journal of Neuro-Oncology</i> , 2016, 129, 319-328.	2.9	25
135	Next-generation sequencing in routine brain tumor diagnostics enables an integrated diagnosis and identifies actionable targets. <i>Acta Neuropathologica</i> , 2016, 131, 903-910.	7.7	203
136	Malignant Transformation of a Dysembryoplastic Neuroepithelial Tumor (DNET) Characterized by Genome-Wide Methylation Analysis. <i>Journal of Neuropathology and Experimental Neurology</i> , 2016, 75, 358-365.	1.7	27
137	Methylation-based classification of benign and malignant peripheral nerve sheath tumors. <i>Acta Neuropathologica</i> , 2016, 131, 877-887.	7.7	151
138	Methylation profiling of choroid plexus tumors reveals 3 clinically distinct subgroups. <i>Neuro-Oncology</i> , 2016, 18, 790-796.	1.2	67
139	Atypical Teratoid/Rhabdoid Tumors Are Comprised of Three Epigenetic Subgroups with Distinct Enhancer Landscapes. <i>Cancer Cell</i> , 2016, 29, 379-393.	16.8	438
140	New Brain Tumor Entities Emerge from Molecular Classification of CNS-PNETs. <i>Cell</i> , 2016, 164, 1060-1072.	28.9	702
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