Matija Snuderl

List of Publications by Year in descending order

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61984 30922 11,824 177 43 102 citations h-index g-index papers 189 189 189 17479 docs citations times ranked citing authors all docs

#	Article	IF	Citations
1	TSC2-mutant uterine sarcomas with JAZF1-SUZ12 fusions demonstrate hybrid features of endometrial stromal sarcoma and PEComa and are responsive to mTOR inhibition. Modern Pathology, 2022, 35, 117-127.	5.5	16
2	Deep Learning and Pathomics Analyses Reveal Cell Nuclei as Important Features for Mutation Prediction of BRAF-Mutated Melanomas. Journal of Investigative Dermatology, 2022, 142, 1650-1658.e6.	0.7	22
3	Molecular neuropathology: The times they are a-changin'. Neuro-Oncology, 2022, , .	1.2	O
4	Integrated Analysis of Ovarian Juvenile Granulosa Cell Tumors Reveals Distinct Epigenetic Signatures and Recurrent <i>TERT</i> Rearrangements. Clinical Cancer Research, 2022, 28, 1724-1733.	7.0	8
5	Utility of multimodality molecular profiling for pediatric patients with central nervous system tumors. Neuro-Oncology Advances, 2022, 4, vdac031.	0.7	1
6	Proteomic differences in hippocampus and cortex of sudden unexplained death in childhood. Acta Neuropathologica, 2022, 143, 585-599.	7.7	7
7	Comprehensive profiling of myxopapillary ependymomas identifies a distinct molecular subtype with relapsing disease. Neuro-Oncology, 2022, 24, 1689-1699.	1.2	11
8	Epigenetic and genomic profiling of chordoid meningioma: implications for clinical management. Acta Neuropathologica Communications, 2022, 10, 56.	5.2	6
9	Thoracic low grade glial neoplasm with concurrent H3 K27M and PTPN11 mutations. Acta Neuropathologica Communications, 2022, 10, 64.	5.2	1
10	Global DNA Methylation Profiles in Peripheral Blood of WTC-Exposed Community Members with Breast Cancer. International Journal of Environmental Research and Public Health, 2022, 19, 5104.	2.6	4
11	DNA methylation as a diagnostic tool. Acta Neuropathologica Communications, 2022, 10, 71.	5.2	24
12	EPEN-18. Oncogenic 3D genome conformations identify novel therapeutic targets in ependymoma. Neuro-Oncology, 2022, 24, i42-i42.	1.2	0
13	ETMR-06. Molecular and clinical characteristics of CNS tumors with <i>BCOR(L1 </i>) fusion/internal tandem duplication. Neuro-Oncology, 2022, 24, i50-i50.	1.2	2
14	MEDB-14. Clinical outcome of pediatric medulloblastoma patients with Li-Fraumeni syndrome. Neuro-Oncology, 2022, 24, i107-i107.	1.2	1
15	MEDB-83. A novel epigenetic nanotherapeutic strategy to induce medulloblastoma differentiation. Neuro-Oncology, 2022, 24, i126-i126.	1,2	O
16	HGG-60. Structural variants shape driver combinations and outcomes in pediatric high-grade glioma. Neuro-Oncology, 2022, 24, i75-i76.	1.2	0
17	EPEN-27. Epigenetic dissection of spinal ependymomas (SP-EPN) separates tumors with and without <i>NF2</i> mutation. Neuro-Oncology, 2022, 24, i44-i45.	1.2	O
18	RARE-15. Astroblastoma, <i>MN1</i> altered comprises two molecularly and clinically distinct subgroups defined by the fusion partners <i>BEND2</i> and <i>CXXC5</i> Neuro-Oncology, 2022, 24, i12-i13.	1,2	1

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19	DNA Methylation Profiling Identifies Subgroups of Lung Adenocarcinoma with Distinct Immune Cell Composition, DNA Methylation Age, and Clinical Outcome. Clinical Cancer Research, 2022, 28, 3824-3835.	7.0	6
20	A Phase I Trial of TB-403 in Relapsed Medulloblastoma, Neuroblastoma, Ewing Sarcoma, and Alveolar Rhabdomyosarcoma. Clinical Cancer Research, 2022, 28, 3950-3957.	7.0	5
21	LMNAâ€NTRK1 rearranged mesenchymal tumor (lipofibromatosisâ€like neural tumor) mimicking pigmented dermatofibrosarcoma protuberans. Journal of Cutaneous Pathology, 2021, 48, 290-294.	1.3	9
22	Analytical performance of lateral flow immunoassay for SARS-CoV-2 exposure screening on venous and capillary blood samples. Journal of Immunological Methods, 2021, 489, 112909.	1.4	32
23	Primary mismatch repair deficient IDH-mutant astrocytoma (PMMRDIA) is a distinct type with a poor prognosis. Acta Neuropathologica, 2021, 141, 85-100.	7.7	52
24	Somatic Focal Copy Number Gains of Noncoding Regions of Receptor Tyrosine Kinase Genes in Treatment-Resistant Epilepsy. Journal of Neuropathology and Experimental Neurology, 2021, 80, 160-168.	1.7	7
25	A subset of pediatric-type thalamic gliomas share a distinct DNA methylation profile, H3K27me3 loss and frequent alteration of <i>EGFR</i> . Neuro-Oncology, 2021, 23, 34-43.	1.2	75
26	Molecular classification and deconvolution of the immune microenvironment in glioblastoma. Neuro-Oncology, 2021, 23, 175-176.	1.2	1
27	NTRK2 Fusion driven pediatric glioblastoma: Identification of oncogenic Drivers via integrative Genome and transcriptome profiling. Clinical Case Reports (discontinued), 2021, 9, 1472-1477.	0.5	3
28	Molecular Signatures of Chromosomal Instability Correlate With Copy Number Variation Patterns and Patient Outcome in IDH-Mutant and IDH-Wildtype Astrocytomas. Journal of Neuropathology and Experimental Neurology, 2021, 80, 354-365.	1.7	12
29	Molecular classification of a complex structural rearrangement of the RB1 locus in an infant with sporadic, isolated, intracranial, sellar region retinoblastoma. Acta Neuropathologica Communications, 2021, 9, 61.	5 . 2	5
30	Cross-Species Genomics Reveals Oncogenic Dependencies in ZFTA/C11orf95 Fusion–Positive Supratentorial Ependymomas. Cancer Discovery, 2021, 11, 2230-2247.	9.4	39
31	Genome-wide association study to reveal novel germline markers of melanoma survival Journal of Clinical Oncology, 2021, 39, 9581-9581.	1.6	O
32	Molecular analysis of encapsulated papillary carcinoma of the breast with and without invasion. Human Pathology, 2021, 111, 67-74.	2.0	7
33	Spatial progression and molecular heterogeneity of IDH-mutant glioblastoma determined by DNA methylation-based mapping. Acta Neuropathologica Communications, 2021, 9, 120.	5.2	6
34	Abstract CT015: A phase 1 dose escalation study of TB-403 in pediatric relapsed or refractory medulloblastoma, neuroblastoma, Ewing sarcoma, or alveolar rhabdomyosarcoma., 2021, , .		1
35	Comparison of solid tissue sequencing and liquid biopsy accuracy in identification of clinically relevant gene mutations and rearrangements in lung adenocarcinomas. Modern Pathology, 2021, 34, 2168-2174.	5.5	21
36	Functional Characterization of Brain Tumor-Initiating Cells and Establishment of GBM Preclinical Models that Incorporate Heterogeneity, Therapy, and Sex Differences. Molecular Cancer Therapeutics, 2021, 20, 2585-2597.	4.1	16

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37	Recurrent fusions in PLAGL1 define a distinct subset of pediatric-type supratentorial neuroepithelial tumors. Acta Neuropathologica, 2021, 142, 827-839.	7.7	33
38	Molecular Pathology of Gliomas. Surgical Pathology Clinics, 2021, 14, 379-386.	1.7	18
39	Sarcoma classification by DNA methylation profiling. Nature Communications, 2021, 12, 498.	12.8	237
40	Clear cell meningiomas are defined by a highly distinct DNA methylation profile and mutations in SMARCE1. Acta Neuropathologica, 2021, 141, 281-290.	7.7	31
41	YAP1-FAM118B Fusion Defines a Rare Subset of Childhood and Young Adulthood Meningiomas. American Journal of Surgical Pathology, 2021, 45, 329-340.	3.7	14
42	Integrated Molecular-Morphologic Meningioma Classification: A Multicenter Retrospective Analysis, Retrospectively and Prospectively Validated. Journal of Clinical Oncology, 2021, 39, 3839-3852.	1.6	93
43	Subgroup-specific outcomes of children with malignant childhood brain tumors treated with an irradiation-sparing protocol. Child's Nervous System, 2020, 36, 133-144.	1.1	3
44	Near real-time intraoperative brain tumor diagnosis using stimulated Raman histology and deep neural networks. Nature Medicine, 2020, 26, 52-58.	30.7	413
45	Genomic Molecular Classification of CNS Malignancies. Advances in Anatomic Pathology, 2020, 27, 44-50.	4.3	5
46	Molecular subgrouping of primary pineal parenchymal tumors reveals distinct subtypes correlated with clinical parameters and genetic alterations. Acta Neuropathologica, 2020, 139, 243-257.	7.7	50
47	Using methylation profiling to diagnose systemic metastases of pleomorphic xanthoastrocytoma. Neuro-Oncology Advances, 2020, 2, vdz057.	0.7	2
48	Sequencing identifies multiple early introductions of SARS-CoV-2 to the New York City region. Genome Research, 2020, 30, 1781-1788.	5 . 5	66
49	Functional Precision Medicine Identifies New Therapeutic Candidates for Medulloblastoma. Cancer Research, 2020, 80, 5393-5407.	0.9	38
50	Expression profiling of the adhesion G protein-coupled receptor GPR133 (ADGRD1) in glioma subtypes. Neuro-Oncology Advances, 2020, 2, vdaa053.	0.7	13
51	Molecular Correlates of Long Survival in IDH-Wildtype Glioblastoma Cohorts. Journal of Neuropathology and Experimental Neurology, 2020, 79, 843-854.	1.7	32
52	Molecular and clinicopathologic features of gliomas harboring NTRK fusions. Acta Neuropathologica Communications, 2020, 8, 107.	5 . 2	84
53	Genome-Wide DNA Methylation Profiles in Community Members Exposed to the World Trade Center Disaster. International Journal of Environmental Research and Public Health, 2020, 17, 5493.	2.6	13
54	Anaplastic Transformation in Myxopapillary Ependymoma: A Report of 2 Cases and Review of the Literature. Journal of Neuropathology and Experimental Neurology, 2020, 79, 1044-1053.	1.7	4

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55	COVID-19-Induced Neurovascular Injury: a Case Series with Emphasis on Pathophysiological Mechanisms. SN Comprehensive Clinical Medicine, 2020, 2, 2109-2125.	0.6	19
56	Correlative study of epigenetic regulation of tumor microenvironment in spindle cell melanomas and cutaneous malignant peripheral nerve sheath tumors. Scientific Reports, 2020, 10, 12996.	3.3	6
57	Limited Environmental Serine and Glycine Confer Brain Metastasis Sensitivity to PHGDH Inhibition. Cancer Discovery, 2020, 10, 1352-1373.	9.4	145
58	MiRâ€1253 exerts tumorâ€suppressive effects in medulloblastoma via inhibition of CDK6 and CD276 (B7â€H3). Brain Pathology, 2020, 30, 732-745.	4.1	35
59	Methylation Profiling of Medulloblastoma in a Clinical Setting Permits Sub-classification and Reveals New Outcome Predictions. Frontiers in Neurology, 2020, 11, 167.	2.4	7
60	Tumor Microenvironment Is Critical for the Maintenance of Cellular States Found in Primary Glioblastomas. Cancer Discovery, 2020, 10, 964-979.	9.4	102
61	Exploring DNA Methylation for Prognosis and Analyzing the Tumor Microenvironment in Pleomorphic Xanthoastrocytoma. Journal of Neuropathology and Experimental Neurology, 2020, 79, 880-890.	1.7	9
62	Association of Initial Viral Load in Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2) Patients with Outcome and Symptoms. American Journal of Pathology, 2020, 190, 1881-1887.	3.8	155
63	Feasibility and clinical utility of a pan-solid tumor targeted RNA fusion panel: A single center experience. Experimental and Molecular Pathology, 2020, 114, 104403.	2.1	9
64	Ganglioglioma in a Survivor of Infantile Glioblastoma. Journal of Pediatric Hematology/Oncology, 2020, 42, e56-e60.	0.6	2
65	Novel EWSR1â€VGLL1 fusion in a pediatric neuroepithelial neoplasm. Clinical Genetics, 2020, 97, 791-792.	2.0	5
66	WNT-Activated Medulloblastomas With Hybrid Molecular Subtypes. JCO Precision Oncology, 2020, 4, 348-354.	3.0	5
67	Genomics of MPNST (GeM) Consortium: Rationale and Study Design for Multi-Omic Characterization of NF1-Associated and Sporadic MPNSTs. Genes, 2020, 11, 387.	2.4	16
68	Diffuse midline glioma with novel, potentially targetable, <i>FGFR2–VPS35</i> fusion. Journal of Physical Education and Sports Management, 2020, 6, a005660.	1.2	5
69	Dissecting the immunosuppressive tumor microenvironments in Glioblastoma-on-a-Chip for optimized PD-1 immunotherapy. ELife, 2020, 9, .	6.0	81
70	MBCL-01. METHYLATION PROFILING OF PEDIATRIC MEDULLOBLASTOMA IN SAUDI ARABIA IN A CLINICAL SETTING PERMITS SUB-CLASSIFICATION AND REVEALS NEW OUTCOME PREDICTIONS. Neuro-Oncology, 2020, 22, iii386-iii387.	1,2	0
71	EPEN-18. CROSS-SPECIES GENOMICS IDENTIFIES GLI2 AS AN ONCOGENE OF C11orf95 FUSION-POSITIVE SUPRATENTORIAL EPENDYMOMA. Neuro-Oncology, 2020, 22, iii311-iii311.	1.2	O
72	Establishing a prognostic threshold for total copy number variation within adult IDH-mutant grade II/III astrocytomas. Acta Neuropathologica Communications, 2019, 7, 121.	5.2	16

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73	Chronic Dengue Virus Panencephalitis in a Patient with Progressive Dementia with Extrapyramidal Features. Annals of Neurology, 2019, 86, 695-703.	5.3	24
74	Histone H3K36I mutation in a metastatic histiocytic tumor of the skull and response to sarcoma chemotherapy. Journal of Physical Education and Sports Management, 2019, 5, a004606.	1.2	8
7 5	Intraosseous Petrous Apex Schwannoma: Case Report and Review of Literature. World Neurosurgery, 2019, 132, 182-187.	1.3	4
76	Revisiting multifocal breast cancer: a clonality study of ductal carcinoma using whole exome sequencing. Human Pathology, 2019, 94, 71-77.	2.0	0
77	Long-term clinical and visual outcomes after surgical resection of pediatric pilocytic/pilomyxoid optic pathway gliomas. Journal of Neurosurgery: Pediatrics, 2019, 24, 166-173.	1.3	17
78	GOPC-ROS1 Fusion Due to Microdeletion at 6q22 Is an Oncogenic Driver in a Subset of Pediatric Gliomas and Glioneuronal Tumors. Journal of Neuropathology and Experimental Neurology, 2019, 78, 1089-1099.	1.7	17
79	Primary CNS Alveolar Rhabdomyosarcoma: Importance of Epigenetic and Transcriptomic Assays for Accurate Diagnosis. Journal of Neuropathology and Experimental Neurology, 2019, 78, 1073-1075.	1.7	6
80	DNA methylation-based classification of sinonasal undifferentiated carcinoma. Modern Pathology, 2019, 32, 1447-1459.	5.5	82
81	MR imaging phenotype correlates with extent of genome-wide copy number abundance in IDH mutant gliomas. Neuroradiology, 2019, 61, 1023-1031.	2.2	8
82	Polysomy is associated with poor outcome in $1p/19q$ codeleted oligodendroglial tumors. Neuro-Oncology, 2019, 21, 1164-1174.	1.2	12
83	Whole transcriptome analysis identifies upregulated genes and pathways in ductal carcinoma in situ mimicking usual ductal hyperplasia. Human Pathology: Case Reports, 2019, 17, 200308.	0.2	0
84	Total copy number variation as a prognostic factor in adult astrocytoma subtypes. Acta Neuropathologica Communications, 2019, 7, 92.	5.2	48
85	GENE-06. DISTINCT MOLECULAR SUBGROUPS OF TUMORS OF THE PINEAL REGION CORRELATE WITH CLINICAL PARAMETERS AND GENETIC ALTERATIONS. Neuro-Oncology, 2019, 21, ii81-ii82.	1.2	O
86	Genome-Wide Analysis of Glioblastoma Patients with Unexpectedly Long Survival. Journal of Neuropathology and Experimental Neurology, 2019, 78, 501-507.	1.7	15
87	Modeling Patient-Derived Glioblastoma with Cerebral Organoids. Cell Reports, 2019, 26, 3203-3211.e5.	6.4	293
88	Recurrent <i>EP300-BCOR</i> Fusions in Pediatric Gliomas With Distinct Clinicopathologic Features. Journal of Neuropathology and Experimental Neurology, 2019, 78, 305-314.	1.7	29
89	ATIM-37. PHASE II, OPEN-LABEL, SINGLE ARM, MULTICENTER STUDY OF AVELUMAB WITH HYPOFRACTIONATED RADIATION (HFRT) FOR ADULT PATIENTS WITH SECONDARILY TRANSFORMED IDH-MUTANT GLIOBLASTOMA (GBM). Neuro-Oncology, 2019, 21, vi9-vi10.	1.2	3
90	PATH-40. PROFILING PLEOMORPHIC XANTHROASTROCYTOMA WITH DNA METHYLATION AND EXPLORING THE TUMOR IMMUNE CELL-TYPE COMPOSITION WITH METHYLATION-BASED DECONVOLUTION. Neuro-Oncology, 2019, 21, vi152-vi152.	1.2	O

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91	Functional and topographic effects on DNA methylation in IDH1/2 mutant cancers. Scientific Reports, $2019, 9, 16830$.	3.3	29
92	GENE-62. IDENTIFICATION OF FGFR4 p.G388R VARIANT IN CEREBELLAR HEMANGIOBLASTOMAS. Neuro-Oncology, 2019, 21, vill1-vill1.	1.2	0
93	NIMG-09. NONINVASIVE PERFUSION IMAGING BIOMARKER OF MALIGNANT GENOTYPE IN ISOCITRATE DEHYDROGENASE MUTANT GLIOMAS. Neuro-Oncology, 2019, 21, vi163-vi163.	1.2	0
94	The molecular landscape of ETMR at diagnosis and relapse. Nature, 2019, 576, 274-280.	27.8	94
95	There is an exception to every ruleâ€"T2-FLAIR mismatch sign in gliomas. Neuroradiology, 2019, 61, 225-227.	2.2	52
96	Cell Surface Notch Ligand DLL3 is a Therapeutic Target in Isocitrate Dehydrogenase–mutant Glioma. Clinical Cancer Research, 2019, 25, 1261-1271.	7.0	50
97	BCAT1 and miR-2504: novel methylome signature distinguishes spindle/desmoplastic melanoma from superficial malignant peripheral nerve sheath tumor. Modern Pathology, 2019, 32, 338-345.	5. 5	8
98	Plasma cell-free circulating tumor DNA (ctDNA) detection in longitudinally followed glioblastoma patients using <i>TERT</i> promoter mutation-specific droplet digital PCR assays Journal of Clinical Oncology, 2019, 37, 2026-2026.	1.6	11
99	Genomic Applications in Brain Tumors. , 2019, , 289-308.		0
100	Single cell analysis of urothelial carcinoma (UC) liver metastases identifies epithelial-mesenchymal transition (EMT) as a potential mechanism of resistance to immunotherapy Journal of Clinical Oncology, 2019, 37, e16018-e16018.	1.6	0
101	A recurrent kinase domain mutation in PRKCA defines chordoid glioma of the third ventricle. Nature Communications, 2018, 9, 810.	12.8	56
102	Cardiac arrhythmia and neuroexcitability gene variants in resected brain tissue from patients with sudden unexpected death in epilepsy (SUDEP). Npj Genomic Medicine, 2018, 3, 9.	3.8	43
103	Loss of histone H3K27me3 identifies a subset of meningiomas with increased risk of recurrence. Acta Neuropathologica, 2018, 135, 955-963.	7.7	109
104	Hacking macrophage-associated immunosuppression for regulating glioblastoma angiogenesis. Biomaterials, 2018, 161, 164-178.	11.4	184
105	Whole Genome DNA Methylation Analysis of Human Glioblastoma Using Illumina BeadArrays. Methods in Molecular Biology, 2018, 1741, 31-51.	0.9	36
106	Programmed death ligand 1 expression and tumor infiltrating lymphocytes in neurofibromatosis type 1 and 2 associated tumors. Journal of Neuro-Oncology, 2018, 138, 183-190.	2.9	54
107	DNA methylation-based classification of central nervous system tumours. Nature, 2018, 555, 469-474.	27.8	1,872
108	Rapid Intraoperative Diagnosis of Pediatric Brain Tumors Using Stimulated Raman Histology. Cancer Research, 2018, 78, 278-289.	0.9	98

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109	Pre-treatment lymphopenia and indication of tumor-induced systemic immunosuppression in medulloblastoma. Journal of Neuro-Oncology, 2018, 136, 541-544.	2.9	14
110	GENE-14. DNA METHYLATION AND PROTEOMIC ALTERATIONS IDENTIFY HISTOLOGICALLY-DEFINED TUMOR CELL POPULATIONS AND CHARACTERIZE INTRATUMOR HETEROGENEITY IN GLIOBLASTOMA. Neuro-Oncology, 2018, 20, vi105-vi105.	1.2	0
111	TBIO-16. AUTOMATED CELL ENRICHMENT AND DIGITAL CELL SORTING USING DIELECTROPHORETIC ARRAYS FOR ISOLATION OF CIRCULATING TUMOR CELLS IN PEDIATRIC BRAIN TUMOR PATIENTS. Neuro-Oncology, 2018, 20, i183-i183.	1.2	O
112	GENE-16. CLINICALLY AGGRESSIVE MENINGIOMAS ARE CHARACTERIZED BY MUTATIONAL SIGNATURES ASSOCIATED WITH DEFECTIVE DNA REPAIR AND MUTATIONS IN CHROMATIN REMODELING GENES. Neuro-Oncology, 2018, 20, vi106-vi106.	1.2	0
113	PDTM-38. PEDIATRIC MENINGIOMAS ARE CHARACTERIZED BY DISTINCT METHYLATION PROFILES DIFFERENT FROM ADULT MENINGIOMAS. Neuro-Oncology, 2018, 20, vi212-vi212.	1.2	1
114	MNGI-14. LOSS OF HISTONE H3K27me3 IDENTIFIES A SUBSET OF MENINGIOMAS WITH INCREASED RISK OF RECURRENCE. Neuro-Oncology, 2018, 20, vi151-vi151.	1.2	0
115	High-Grade Glioma, Including Diffuse Intrinsic Pontine Glioma. , 2018, , 193-221.		0
116	Classification and mutation prediction from non–small cell lung cancer histopathology images using deep learning. Nature Medicine, 2018, 24, 1559-1567.	30.7	1,768
117	Predicting Genotype and Survival in Glioma Using Standard Clinical MR Imaging Apparent Diffusion Coefficient Images: A Pilot Study from The Cancer Genome Atlas. American Journal of Neuroradiology, 2018, 39, 1814-1820.	2.4	53
118	NFM-11. PEDIATRIC MENINGIOMAS ARE MOLECULARLY DISTINCT FROM ADULT COUNTERPARTS. Neuro-Oncology, 2018, 20, i144-i145.	1.2	1
119	Aspartate is a limiting metabolite for cancer cell proliferation under hypoxia and in tumours. Nature Cell Biology, 2018, 20, 775-781.	10.3	311
120	Recurrent homozygous deletion of DROSHA and microduplication of PDE4DIP in pineoblastoma. Nature Communications, 2018, 9, 2868.	12.8	54
121	A case of molecularly profiled extraneural medulloblastoma metastases in a child. BMC Medical Genetics, 2018, 19, 10.	2.1	3
122	Genetic and Epigenetic Features of Rapidly Progressing IDH-Mutant Astrocytomas. Journal of Neuropathology and Experimental Neurology, 2018, 77, 542-548.	1.7	34
123	Primary intracranial spindle cell sarcoma with rhabdomyosarcoma-like features share a highly distinct methylation profile and DICER1 mutations. Acta Neuropathologica, 2018, 136, 327-337.	7.7	104
124	DNA methylation-based classifier for diagnosis of endometrial cancer Journal of Clinical Oncology, 2018, 36, e17570-e17570.	1.6	0
125	Rapid intraoperative histology of unprocessed surgical specimens via fibre-laser-based stimulated Raman scattering microscopy. Nature Biomedical Engineering, 2017, 1, .	22.5	374
126	Mutant IDH1 and seizures in patients with glioma. Neurology, 2017, 88, 1805-1813.	1.1	167

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127	Rapid progression to glioblastoma in a subset of IDH-mutated astrocytomas: a genome-wide analysis. Journal of Neuro-Oncology, 2017, 133, 183-192.	2.9	30
128	Low-Grade Astrocytoma Mutations in IDH1, P53, and ATRX Cooperate to Block Differentiation of Human Neural Stem Cells via Repression of SOX2. Cell Reports, 2017, 21, 1267-1280.	6.4	95
129	Immunohistochemical analysis of H3K27me3 demonstrates global reduction in group-A childhood posterior fossa ependymoma and is a powerful predictor of outcome. Acta Neuropathologica, 2017, 134, 705-714.	7.7	168
130	T2–FLAIR Mismatch, an Imaging Biomarker for IDH and 1p/19q Status in Lower-grade Gliomas: A TCGA/TCIA Project. Clinical Cancer Research, 2017, 23, 6078-6085.	7. O	285
131	Osimertinib Dose Escalation Induces Regression of Progressive EGFR T790M–Mutant Leptomeningeal Lung Adenocarcinoma. Journal of Thoracic Oncology, 2017, 12, e188-e190.	1.1	13
132	Endothelium-Independent Primitive Myxoid Vascularization Creates Invertebrate-Like Channels to Maintain Blood Supply in Optic Gliomas. American Journal of Pathology, 2017, 187, 1867-1878.	3.8	4
133	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. Acta Neuropathologica, 2017, 133, 417-429.	7.7	172
134	GENE-02. PERIPHERAL BLOOD DNA METHYLATION PROFILES IDENTIFY IDH1/2 MUTATION STATUS IN ADULTS WITH DIFFUSE GLIOMA. Neuro-Oncology, 2017, 19, vi92-vi92.	1.2	0
135	DNA Methylation–Based Classifier for Accurate Molecular Diagnosis of Bone Sarcomas. JCO Precision Oncology, 2017, 2017, 1-11.	3.0	37
136	Apolipoprotein L1 risk variants associate with prevalent atherosclerotic disease in African American systemic lupus erythematosus patients. PLoS ONE, 2017, 12, e0182483.	2.5	21
137	Notch signaling regulates metabolic heterogeneity in glioblastoma stem cells. Oncotarget, 2017, 8, 64932-64953.	1.8	58
138	A DNA methylation-based classifier for accurate molecular diagnosis of bone sarcomas Journal of Clinical Oncology, 2017, 35, 11034-11034.	1.6	0
139	IMST-40. REPROGRAMMING OF THE TUMOR IMMUNE MICROENVIRONMENT BY AN ANG-2/VEGF BISPECIFIC ANTIBODY DELAYS TUMOR GROWTH AND PROLONGS SURVIVAL IN PRECLINICAL GBM MODELS. Neuro-Oncology, 2016, 18, vi95-vi95.	1.2	0
140	Adult Primary Spinal Epidural Extraosseous Ewing's Sarcoma: A Case Report and Review of the Literature. Case Reports in Neurological Medicine, 2016, 2016, 1-8.	0.4	9
141	HG-127ANAPLASTIC PLEOMORPHIC XANTHOASTROCYTOMAS: A CLINICOPATHOLOGIC AND MOLECULAR PROFILE. Neuro-Oncology, 2016, 18, iii77.3-iii77.	1.2	0
142	HG-73SAFETY AND FEASIBILITY OF A MULTI-INSTITUTIONAL PHASE II TRIAL INCOPORATING BIOPSY AND MOLECULARLY DETERMINED TREATMENT OF CHILDREN AND YOUNG ADULTS WITH NEWLY DIAGNOSED DIFFUSE INTRINSIC PONTINE GLIOMAS (DIPG). Neuro-Oncology, 2016, 18, iii65.1-iii65.	1.2	0
143	LG-67MIDBRAIN GLIOMAS: A LARGE SERIES OF CLINICALLY AND RADIOGRAPHICALLY HETEROGENEOUS TUMORS. Neuro-Oncology, 2016, 18, iii94.2-iii94.	1.2	0
144	LG-74STRATEGIES FOR THE SURGICAL MANAGEMENT OF PEDIATRIC OPTIC PATHWAY GLIOMAS - EXPERIENCE WITH 100 PATIENTS. Neuro-Oncology, 2016, 18, iii95.5-iii96.	1.2	0

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145	TB-27SUBGROUP-SPECIFIC OUTCOMES OF CHILDREN WITH MALIGNANT CHILDHOOD BRAIN TUMORS TREATED WITH AN IRRADIATION-SPARING PROTOCOL. Neuro-Oncology, 2016, 18, iii173.3-iii173.	1.2	O
146	MPTH-34. THE PROGNOSTIC VALUE OF POLYSOMY IN OLIGODENDROGLIAL TUMORS. Neuro-Oncology, 2016, 18, vi113-vi113.	1.2	0
147	MPTH-59. ANAPLASTIC PLEOMORPHIC XANTHOASTROCYTOMAS: AÂCLINICOPATHOLOGIC AND MOLECULAR PROFILE. Neuro-Oncology, 2016, 18, vi118-vi118.	1.2	O
148	NIMG-76. MIDBRAIN GLIOMAS: AÂLARGE SERIES THAT IDENTIFIES FEATURES CORRESPONDING WITH OUTCOME. Neuro-Oncology, 2016, 18, vi141-vi141.	1.2	0
149	STMC-21. ASTROCYTOMA MUTATIONS IDH1, p53 AND ATRX COOPERATE TO BLOCK DIFFERENTIATION OF NEURAL STEM CELLS VIA Sox2. Neuro-Oncology, 2016, 18, vi187-vi187.	1.2	O
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