

Michael D Taylor

List of Publications by Year in descending order

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Version: 2024-02-01

419
papers

47,691
citations

1893

102
h-index

2178

202
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448
all docs

448
docs citations

448
times ranked

35961
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | DNA methylation-based classification of central nervous system tumours. <i>Nature</i> , 2018, 555, 469-474. | 27.8 | 1,872 |
| 2 | Hotspot Mutations in H3F3A and IDH1 Define Distinct Epigenetic and Biological Subgroups of Glioblastoma. <i>Cancer Cell</i> , 2012, 22, 425-437. | 16.8 | 1,551 |
| 3 | Molecular subgroups of medulloblastoma: the current consensus. <i>Acta Neuropathologica</i> , 2012, 123, 465-472. | 7.7 | 1,536 |
| 4 | Medulloblastoma Comprises Four Distinct Molecular Variants. <i>Journal of Clinical Oncology</i> , 2011, 29, 1408-1414. | 1.6 | 1,131 |
| 5 | Molecular Classification of Ependymal Tumors across All CNS Compartments, Histopathological Grades, and Age Groups. <i>Cancer Cell</i> , 2015, 27, 728-743. | 16.8 | 933 |
| 6 | Molecular subgroups of medulloblastoma: an international meta-analysis of transcriptome, genetic aberrations, and clinical data of WNT, SHH, Group 3, and Group 4 medulloblastomas. <i>Acta Neuropathologica</i> , 2012, 123, 473-484. | 7.7 | 863 |
| 7 | Intertumoral Heterogeneity within Medulloblastoma Subgroups. <i>Cancer Cell</i> , 2017, 31, 737-754.e6. | 16.8 | 836 |
| 8 | The whole-genome landscape of medulloblastoma subtypes. <i>Nature</i> , 2017, 547, 311-317. | 27.8 | 787 |
| 9 | Dissecting the genomic complexity underlying medulloblastoma. <i>Nature</i> , 2012, 488, 100-105. | 27.8 | 765 |
| 10 | Subgroup-specific structural variation across 1,000 medulloblastoma genomes. <i>Nature</i> , 2012, 488, 49-56. | 27.8 | 761 |
| 11 | Radial glia cells are candidate stem cells of ependymoma. <i>Cancer Cell</i> , 2005, 8, 323-335. | 16.8 | 758 |
| 12 | Genome Sequencing of Pediatric Medulloblastoma Links Catastrophic DNA Rearrangements with TP53 Mutations. <i>Cell</i> , 2012, 148, 59-71. | 28.9 | 743 |
| 13 | Mutations in SUFU predispose to medulloblastoma. <i>Nature Genetics</i> , 2002, 31, 306-310. | 21.4 | 722 |
| 14 | New Brain Tumor Entities Emerge from Molecular Classification of CNS-PNETs. <i>Cell</i> , 2016, 164, 1060-1072. | 28.9 | 702 |
| 15 | Medulloblastoma exome sequencing uncovers subtype-specific somatic mutations. <i>Nature</i> , 2012, 488, 106-110. | 27.8 | 675 |
| 16 | Recurrent somatic alterations of FGFR1 and NTRK2 in pilocytic astrocytoma. <i>Nature Genetics</i> , 2013, 45, 927-932. | 21.4 | 674 |
| 17 | The Genetic Landscape of the Childhood Cancer Medulloblastoma. <i>Science</i> , 2011, 331, 435-439. | 12.6 | 652 |
| 18 | Genome Sequencing of SHH Medulloblastoma Predicts Genotype-Related Response to Smoothed Inhibition. <i>Cancer Cell</i> , 2014, 25, 393-405. | 16.8 | 627 |

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|----|--|------|-----------|
| 19 | Genomics Identifies Medulloblastoma Subgroups That Are Enriched for Specific Genetic Alterations. <i>Journal of Clinical Oncology</i> , 2006, 24, 1924-1931. | 1.6 | 617 |
| 20 | Comprehensive Analysis of Hypermutation in Human Cancer. <i>Cell</i> , 2017, 171, 1042-1056.e10. | 28.9 | 596 |
| 21 | Medulloblastomics: the end of the beginning. <i>Nature Reviews Cancer</i> , 2012, 12, 818-834. | 28.4 | 560 |
| 22 | Challenges to curing primary brain tumours. <i>Nature Reviews Clinical Oncology</i> , 2019, 16, 509-520. | 27.6 | 540 |
| 23 | Extrachromosomal oncogene amplification drives tumour evolution and genetic heterogeneity. <i>Nature</i> , 2017, 543, 122-125. | 27.8 | 530 |
| 24 | Enhancer hijacking activates GFI1 family oncogenes in medulloblastoma. <i>Nature</i> , 2014, 511, 428-434. | 27.8 | 520 |
| 25 | Delineation of Two Clinically and Molecularly Distinct Subgroups of Posterior Fossa Ependymoma. <i>Cancer Cell</i> , 2011, 20, 143-157. | 16.8 | 494 |
| 26 | Risk stratification of childhood medulloblastoma in the molecular era: the current consensus. <i>Acta Neuropathologica</i> , 2016, 131, 821-831. | 7.7 | 478 |
| 27 | 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 |
| 28 | Awake craniotomy with brain mapping as the routine surgical approach to treating patients with supratentorial intraaxial tumors: a prospective trial of 200 cases. <i>Journal of Neurosurgery</i> , 1999, 90, 35-41. | 1.6 | 429 |
| 29 | Multiple recurrent genetic events converge on control of histone lysine methylation in medulloblastoma. <i>Nature Genetics</i> , 2009, 41, 465-472. | 21.4 | 391 |
| 30 | Subgroup-Specific Prognostic Implications of TP53 Mutation in Medulloblastoma. <i>Journal of Clinical Oncology</i> , 2013, 31, 2927-2935. | 1.6 | 381 |
| 31 | Decoding the regulatory landscape of medulloblastoma using DNA methylation sequencing. <i>Nature</i> , 2014, 510, 537-541. | 27.8 | 378 |
| 32 | Clonal selection drives genetic divergence of metastatic medulloblastoma. <i>Nature</i> , 2012, 482, 529-533. | 27.8 | 376 |
| 33 | Medulloblastoma. <i>Nature Reviews Disease Primers</i> , 2019, 5, 11. | 30.5 | 376 |
| 34 | The eEF2 Kinase Confers Resistance to Nutrient Deprivation by Blocking Translation Elongation. <i>Cell</i> , 2013, 153, 1064-1079. | 28.9 | 348 |
| 35 | YAP1 is amplified and up-regulated in hedgehog-associated medulloblastomas and mediates Sonic hedgehog-driven neural precursor proliferation. <i>Genes and Development</i> , 2009, 23, 2729-2741. | 5.9 | 332 |
| 36 | Cross-species genomics matches driver mutations and cell compartments to model ependymoma. <i>Nature</i> , 2010, 466, 632-636. | 27.8 | 324 |

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|----|--|------|-----------|
| 37 | Fate mapping of human glioblastoma reveals an invariant stem cell hierarchy. <i>Nature</i> , 2017, 549, 227-232. | 27.8 | 321 |
| 38 | Rapid, reliable, and reproducible molecular sub-grouping of clinical medulloblastoma samples. <i>Acta Neuropathologica</i> , 2012, 123, 615-626. | 7.7 | 318 |
| 39 | Recurrence patterns across medulloblastoma subgroups: an integrated clinical and molecular analysis. <i>Lancet Oncology</i> , The, 2013, 14, 1200-1207. | 10.7 | 307 |
| 40 | Combined hereditary and somatic mutations of replication error repair genes result in rapid onset of ultra-hypermutated cancers. <i>Nature Genetics</i> , 2015, 47, 257-262. | 21.4 | 306 |
| 41 | Disrupting the CD47-SIRP α anti-phagocytic axis by a humanized anti-CD47 antibody is an efficacious treatment for malignant pediatric brain tumors. <i>Science Translational Medicine</i> , 2017, 9, . | 12.4 | 306 |
| 42 | Childhood cerebellar tumours mirror conserved fetal transcriptional programs. <i>Nature</i> , 2019, 572, 67-73. | 27.8 | 293 |
| 43 | Prognostic value of medulloblastoma extent of resection after accounting for molecular subgroup: a retrospective integrated clinical and molecular analysis. <i>Lancet Oncology</i> , The, 2016, 17, 484-495. | 10.7 | 274 |
| 44 | The miR-17/92 Polycistron Is Up-regulated in Sonic Hedgehog-Driven Medulloblastomas and Induced by N-myc in Sonic Hedgehog-Treated Cerebellar Neural Precursors. <i>Cancer Research</i> , 2009, 69, 3249-3255. | 0.9 | 273 |
| 45 | The current consensus on the clinical management of intracranial ependymoma and its distinct molecular variants. <i>Acta Neuropathologica</i> , 2017, 133, 5-12. | 7.7 | 271 |
| 46 | Spectrum and prevalence of genetic predisposition in medulloblastoma: a retrospective genetic study and prospective validation in a clinical trial cohort. <i>Lancet Oncology</i> , The, 2018, 19, 785-798. | 10.7 | 268 |
| 47 | An Animal Model of MYC-Driven Medulloblastoma. <i>Cancer Cell</i> , 2012, 21, 155-167. | 16.8 | 267 |
| 48 | Divergent clonal selection dominates medulloblastoma at recurrence. <i>Nature</i> , 2016, 529, 351-357. | 27.8 | 266 |
| 49 | Cytogenetic Prognostication Within Medulloblastoma Subgroups. <i>Journal of Clinical Oncology</i> , 2014, 32, 886-896. | 1.6 | 263 |
| 50 | The clinical implications of medulloblastoma subgroups. <i>Nature Reviews Neurology</i> , 2012, 8, 340-351. | 10.1 | 261 |
| 51 | Robust molecular subgrouping and copy-number profiling of medulloblastoma from small amounts of archival tumour material using high-density DNA methylation arrays. <i>Acta Neuropathologica</i> , 2013, 125, 913-916. | 7.7 | 244 |
| 52 | <i>BRAF</i> Mutation and <i>CDKN2A</i> Deletion Define a Clinically Distinct Subgroup of Childhood Secondary High-Grade Glioma. <i>Journal of Clinical Oncology</i> , 2015, 33, 1015-1022. | 1.6 | 244 |
| 53 | Integrated Molecular and Clinical Analysis of 1,000 Pediatric Low-Grade Gliomas. <i>Cancer Cell</i> , 2020, 37, 569-583.e5. | 16.8 | 244 |
| 54 | Quiescent Sox2+ Cells Drive Hierarchical Growth and Relapse in Sonic Hedgehog Subgroup Medulloblastoma. <i>Cancer Cell</i> , 2014, 26, 33-47. | 16.8 | 241 |

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|----|---|------|-----------|
| 55 | MicroRNA-199b-5p Impairs Cancer Stem Cells through Negative Regulation of HES1 in Medulloblastoma. PLoS ONE, 2009, 4, e4998. | 2.5 | 233 |
| 56 | Therapeutic and Prognostic Implications of BRAF V600E in Pediatric Low-Grade Gliomas. Journal of Clinical Oncology, 2017, 35, 2934-2941. | 1.6 | 232 |
| 57 | Adult Medulloblastoma Comprises Three Major Molecular Variants. Journal of Clinical Oncology, 2011, 29, 2717-2723. | 1.6 | 215 |
| 58 | Methylation of the TERT promoter and risk stratification of childhood brain tumours: an integrative genomic and molecular study. Lancet Oncology, The, 2013, 14, 534-542. | 10.7 | 212 |
| 59 | Frequent Amplification of a chr19q13.41 MicroRNA Polycistron in Aggressive Primitive Neuroectodermal Brain Tumors. Cancer Cell, 2009, 16, 533-546. | 16.8 | 207 |
| 60 | HDAC and PI3K Antagonists Cooperate to Inhibit Growth of MYC- Driven Medulloblastoma. Cancer Cell, 2016, 29, 311-323. | 16.8 | 204 |
| 61 | Alterations in ALK/ROS1/NTRK/MET drive a group of infantile hemispheric gliomas. Nature Communications, 2019, 10, 4343. | 12.8 | 200 |
| 62 | Molecular heterogeneity and CXorf67 alterations in posterior fossa group A (PFA) ependymomas. Acta Neuropathologica, 2018, 136, 211-226. | 7.7 | 199 |
| 63 | Pediatric and adult sonic hedgehog medulloblastomas are clinically and molecularly distinct. Acta Neuropathologica, 2011, 122, 231-240. | 7.7 | 195 |
| 64 | Embryonal tumor with abundant neuropil and true rosettes (ETANTR), ependymoblastoma, and medulloepithelioma share molecular similarity and comprise a single clinicopathological entity. Acta Neuropathologica, 2014, 128, 279-289. | 7.7 | 191 |
| 65 | Integrated (epi)-Genomic Analyses Identify Subgroup-Specific Therapeutic Targets in CNS Rhabdoid Tumors. Cancer Cell, 2016, 30, 891-908. | 16.8 | 191 |
| 66 | Roadmap for the Emerging Field of Cancer Neuroscience. Cell, 2020, 181, 219-222. | 28.9 | 182 |
| 67 | Second-generation molecular subgrouping of medulloblastoma: an international meta-analysis of Group 3 and Group 4 subtypes. Acta Neuropathologica, 2019, 138, 309-326. | 7.7 | 180 |
| 68 | Distinct Neural Stem Cell Populations Give Rise to Disparate Brain Tumors in Response to N-MYC. Cancer Cell, 2012, 21, 601-613. | 16.8 | 177 |
| 69 | Impact of Craniospinal Dose, Boost Volume, and Neurologic Complications on Intellectual Outcome in Patients With Medulloblastoma. Journal of Clinical Oncology, 2014, 32, 1760-1768. | 1.6 | 177 |
| 70 | HDAC5 and HDAC9 in Medulloblastoma: Novel Markers for Risk Stratification and Role in Tumor Cell Growth. Clinical Cancer Research, 2010, 16, 3240-3252. | 7.0 | 175 |
| 71 | Familial Posterior Fossa Brain Tumors of Infancy Secondary to Germline Mutation of the hSNF5 Gene. American Journal of Human Genetics, 2000, 66, 1403-1406. | 6.2 | 170 |
| 72 | Therapeutic targeting of ependymoma as informed by oncogenic enhancer profiling. Nature, 2018, 553, 101-105. | 27.8 | 170 |

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|----|---|------|-----------|
| 73 | Aberrant patterns of H3K4 and H3K27 histone lysine methylation occur across subgroups in medulloblastoma. <i>Acta Neuropathologica</i> , 2013, 125, 373-384. | 7.7 | 169 |
| 74 | Immunohistochemical analysis of H3K27me3 demonstrates global reduction in group-A childhood posterior fossa ependymoma and is a powerful predictor of outcome. <i>Acta Neuropathologica</i> , 2017, 134, 705-714. | 7.7 | 168 |
| 75 | Fusion of TTYH1 with the C19MC microRNA cluster drives expression of a brain-specific DNMT3B isoform in the embryonal brain tumor ETMR. <i>Nature Genetics</i> , 2014, 46, 39-44. | 21.4 | 167 |
| 76 | Therapeutic Impact of Cytoreductive Surgery and Irradiation of Posterior Fossa Ependymoma in the Molecular Era: A Retrospective Multicohort Analysis. <i>Journal of Clinical Oncology</i> , 2016, 34, 2468-2477. | 1.6 | 160 |
| 77 | Medulloblastoma in the age of molecular subgroups: a review. <i>Journal of Neurosurgery: Pediatrics</i> , 2019, 24, 353-363. | 1.3 | 153 |
| 78 | Awake Craniotomy for Removal of Intracranial Tumor: Considerations for Early Discharge. <i>Anesthesia and Analgesia</i> , 2001, 92, 89-94. | 2.2 | 149 |
| 79 | Universal Poor Survival in Children With Medulloblastoma Harboring Somatic <i>TP53</i> Mutations. <i>Journal of Clinical Oncology</i> , 2010, 28, 1345-1350. | 1.6 | 148 |
| 80 | Identification of GPC2 as an Oncoprotein and Candidate Immunotherapeutic Target in High-Risk Neuroblastoma. <i>Cancer Cell</i> , 2017, 32, 295-309.e12. | 16.8 | 148 |
| 81 | Molecular subgroups of atypical teratoid rhabdoid tumours in children: an integrated genomic and clinicopathological analysis. <i>Lancet Oncology</i> , The, 2015, 16, 569-582. | 10.7 | 147 |
| 82 | Pleiotropic role for <i>MYCN</i> in medulloblastoma. <i>Genes and Development</i> , 2010, 24, 1059-1072. | 5.9 | 146 |
| 83 | TERT promoter mutations are highly recurrent in SHH subgroup medulloblastoma. <i>Acta Neuropathologica</i> , 2013, 126, 917-929. | 7.7 | 146 |
| 84 | Identification of differentially expressed and developmentally regulated genes in medulloblastoma using suppression subtraction hybridization. <i>Oncogene</i> , 2004, 23, 3444-3453. | 5.9 | 144 |
| 85 | <i>FSTL5</i> Is a Marker of Poor Prognosis in Non-WNT/Non-SHH Medulloblastoma. <i>Journal of Clinical Oncology</i> , 2011, 29, 3852-3861. | 1.6 | 143 |
| 86 | Superior Intellectual Outcomes After Proton Radiotherapy Compared With Photon Radiotherapy for Pediatric Medulloblastoma. <i>Journal of Clinical Oncology</i> , 2020, 38, 454-461. | 1.6 | 143 |
| 87 | Molecular subgroups of medulloblastoma. <i>Expert Review of Neurotherapeutics</i> , 2012, 12, 871-884. | 2.8 | 142 |
| 88 | CNS-PNETs with C19MC amplification and/or LIN28 expression comprise a distinct histogenetic diagnostic and therapeutic entity. <i>Acta Neuropathologica</i> , 2014, 128, 291-303. | 7.7 | 141 |
| 89 | Locoregional delivery of CAR T cells to the cerebrospinal fluid for treatment of metastatic medulloblastoma and ependymoma. <i>Nature Medicine</i> , 2020, 26, 720-731. | 30.7 | 141 |
| 90 | Stalled developmental programs at the root of pediatric brain tumors. <i>Nature Genetics</i> , 2019, 51, 1702-1713. | 21.4 | 136 |

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|-----|--|------|-----------|
| 91 | DNA methylation profiling of medulloblastoma allows robust subclassification and improved outcome prediction using formalin-fixed biopsies. <i>Acta Neuropathologica</i> , 2013, 125, 359-371. | 7.7 | 133 |
| 92 | DNA hypermethylation within TERT promoter upregulates TERT expression in cancer. <i>Journal of Clinical Investigation</i> , 2018, 129, 223-229. | 8.2 | 130 |
| 93 | Medulloblastoma: From Myth to Molecular. <i>Journal of Clinical Oncology</i> , 2017, 35, 2355-2363. | 1.6 | 129 |
| 94 | Recurrent noncoding U1 snRNA mutations drive cryptic splicing in SHH medulloblastoma. <i>Nature</i> , 2019, 574, 707-711. | 27.8 | 129 |
| 95 | The U1 spliceosomal RNA is recurrently mutated in multiple cancers. <i>Nature</i> , 2019, 574, 712-716. | 27.8 | 128 |
| 96 | Prognostic significance of clinical, histopathological, and molecular characteristics of medulloblastomas in the prospective HIT2000 multicenter clinical trial cohort. <i>Acta Neuropathologica</i> , 2014, 128, 137-149. | 7.7 | 125 |
| 97 | Molecular cytogenetic analysis of medulloblastomas and supratentorial primitive neuroectodermal tumors by using conventional banding, comparative genomic hybridization, and spectral karyotyping. <i>Journal of Neurosurgery</i> , 2000, 93, 437-448. | 1.6 | 124 |
| 98 | Molecular Insights into Pediatric Brain Tumors Have the Potential to Transform Therapy. <i>Clinical Cancer Research</i> , 2014, 20, 5630-5640. | 7.0 | 124 |
| 99 | Myocardial Fibrosis Burden Predicts Left Ventricular Ejection Fraction and Is Associated With Age and Steroid Treatment Duration in Duchenne Muscular Dystrophy. <i>Journal of the American Heart Association</i> , 2015, 4, . | 3.7 | 114 |
| 100 | Clinical and neuroanatomical predictors of cerebellar mutism syndrome. <i>Neuro-Oncology</i> , 2012, 14, 1294-1303. | 1.2 | 112 |
| 101 | Medulloblastoma subgroup-specific outcomes in irradiated children: who are the true high-risk patients?. <i>Neuro-Oncology</i> , 2016, 18, 291-297. | 1.2 | 112 |
| 102 | Spatial heterogeneity in medulloblastoma. <i>Nature Genetics</i> , 2017, 49, 780-788. | 21.4 | 112 |
| 103 | Survival Benefit for Pediatric Patients With Recurrent Ependymoma Treated With Reirradiation. <i>International Journal of Radiation Oncology Biology Physics</i> , 2012, 83, 1541-1548. | 0.8 | 111 |
| 104 | Molecular Insight into Medulloblastoma and Central Nervous System Primitive Neuroectodermal Tumor Biology from Hereditary Syndromes: A Review. <i>Neurosurgery</i> , 2000, 47, 888-901. | 1.1 | 110 |
| 105 | The G protein α subunit G_{i1} is a tumor suppressor in Sonic hedgehog-driven medulloblastoma. <i>Nature Medicine</i> , 2014, 20, 1035-1042. | 30.7 | 110 |
| 106 | PINK1 Is a Negative Regulator of Growth and the Warburg Effect in Glioblastoma. <i>Cancer Research</i> , 2016, 76, 4708-4719. | 0.9 | 107 |
| 107 | Genome-Wide Profiles of Extra-cranial Malignant Rhabdoid Tumors Reveal Heterogeneity and Dysregulated Developmental Pathways. <i>Cancer Cell</i> , 2016, 29, 394-406. | 16.8 | 105 |
| 108 | OTX2 Is Critical for the Maintenance and Progression of Shh-Independent Medulloblastomas. <i>Cancer Research</i> , 2010, 70, 181-191. | 0.9 | 104 |

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|-----|---|------|-----------|
| 109 | Aberrant ERBB4-SRC Signaling as a Hallmark of Group 4 Medulloblastoma Revealed by Integrative Phosphoproteomic Profiling. <i>Cancer Cell</i> , 2018, 34, 379-395.e7. | 16.8 | 104 |
| 110 | Inhibition of BRD4 attenuates tumor cell self-renewal and suppresses stem cell signaling in MYC driven medulloblastoma. <i>Oncotarget</i> , 2014, 5, 2355-2371. | 1.8 | 103 |
| 111 | <i>TP53</i> Mutation Is Frequently Associated With <i>CTNNB1</i> Mutation or <i>MYCN</i> Amplification and Is Compatible With Long-Term Survival in Medulloblastoma. <i>Journal of Clinical Oncology</i> , 2010, 28, 5188-5196. | 1.6 | 100 |
| 112 | MicroRNA 218 Acts as a Tumor Suppressor by Targeting Multiple Cancer Phenotype-associated Genes in Medulloblastoma. <i>Journal of Biological Chemistry</i> , 2013, 288, 1918-1928. | 3.4 | 100 |
| 113 | Single-Cell Transcriptomics in Medulloblastoma Reveals Tumor-Initiating Progenitors and Oncogenic Cascades during Tumorigenesis and Relapse. <i>Cancer Cell</i> , 2019, 36, 302-318.e7. | 16.8 | 96 |
| 114 | An Epigenetic Genome-Wide Screen Identifies <i>SPINT2</i> as a Novel Tumor Suppressor Gene in Pediatric Medulloblastoma. <i>Cancer Research</i> , 2008, 68, 9945-9953. | 0.9 | 95 |
| 115 | Clinical, Pathological, and Molecular Characterization of Infant Medulloblastomas Treated with Sequential High-Dose Chemotherapy. <i>Pediatric Blood and Cancer</i> , 2016, 63, 1527-1534. | 1.5 | 94 |
| 116 | The molecular landscape of ETMR at diagnosis and relapse. <i>Nature</i> , 2019, 576, 274-280. | 27.8 | 94 |
| 117 | Histone H3.3G34-Mutant Interneuron Progenitors Co-opt PDGFRA for Gliomagenesis. <i>Cell</i> , 2020, 183, 1617-1633.e22. | 28.9 | 93 |
| 118 | MLL5 Orchestrates a Cancer Self-Renewal State by Repressing the Histone Variant H3.3 and Globally Reorganizing Chromatin. <i>Cancer Cell</i> , 2015, 28, 715-729. | 16.8 | 90 |
| 119 | MR Imaging-Based Radiomic Signatures of Distinct Molecular Subgroups of Medulloblastoma. <i>American Journal of Neuroradiology</i> , 2019, 40, 154-161. | 2.4 | 87 |
| 120 | Heterogeneity within the PF-EPN-B ependymoma subgroup. <i>Acta Neuropathologica</i> , 2018, 136, 227-237. | 7.7 | 86 |
| 121 | A Hematogenous Route for Medulloblastoma Leptomeningeal Metastases. <i>Cell</i> , 2018, 172, 1050-1062.e14. | 28.9 | 85 |
| 122 | Molecular Characterization of Choroid Plexus Tumors Reveals Novel Clinically Relevant Subgroups. <i>Clinical Cancer Research</i> , 2015, 21, 184-192. | 7.0 | 84 |
| 123 | Molecular diagnostics of CNS embryonal tumors. <i>Acta Neuropathologica</i> , 2010, 120, 553-566. | 7.7 | 83 |
| 124 | Medulloblastoma subgroups remain stable across primary and metastatic compartments. <i>Acta Neuropathologica</i> , 2015, 129, 449-457. | 7.7 | 80 |
| 125 | Personalizing the Treatment of Pediatric Medulloblastoma: Polo-like Kinase 1 as a Molecular Target in High-Risk Children. <i>Cancer Research</i> , 2013, 73, 6734-6744. | 0.9 | 79 |
| 126 | Metabolic Regulation of the Epigenome Drives Lethal Infantile Ependymoma. <i>Cell</i> , 2020, 181, 1329-1345.e24. | 28.9 | 79 |

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|-----|--|------|-----------|
| 127 | scRNA-seq in medulloblastoma shows cellular heterogeneity and lineage expansion support resistance to SHH inhibitor therapy. <i>Nature Communications</i> , 2019, 10, 5829. | 12.8 | 77 |
| 128 | An epigenetic gateway to brain tumor cell identity. <i>Nature Neuroscience</i> , 2016, 19, 10-19. | 14.8 | 76 |
| 129 | Therapeutic radiation for childhood cancer drives structural aberrations of NF2 in meningiomas. <i>Nature Communications</i> , 2017, 8, 186. | 12.8 | 76 |
| 130 | Failure of a medulloblastoma-derived mutant of SUFU to suppress WNT signaling. <i>Oncogene</i> , 2004, 23, 4577-4583. | 5.9 | 75 |
| 131 | EAG2 potassium channel with evolutionarily conserved function as a brain tumor target. <i>Nature Neuroscience</i> , 2015, 18, 1236-1246. | 14.8 | 74 |
| 132 | Significance of molecular classification of ependymomas: C11orf95-RELA fusion-negative supratentorial ependymomas are a heterogeneous group of tumors. <i>Acta Neuropathologica Communications</i> , 2018, 6, 134. | 5.2 | 74 |
| 133 | Identification and Analyses of Extra-Cranial and Cranial Rhabdoid Tumor Molecular Subgroups Reveal Tumors with Cytotoxic T Cell Infiltration. <i>Cell Reports</i> , 2019, 29, 2338-2354.e7. | 6.4 | 74 |
| 134 | The RAG-1/2 endonuclease causes genomic instability and controls CNS complications of lymphoblastic leukemia in p53/Prkdc-deficient mice. <i>Cancer Cell</i> , 2003, 3, 37-50. | 16.8 | 73 |
| 135 | The genetic and epigenetic basis of ependymoma. <i>Child's Nervous System</i> , 2009, 25, 1195-1201. | 1.1 | 73 |
| 136 | The RNA-Binding Protein Musashi1 Affects Medulloblastoma Growth via a Network of Cancer-Related Genes and Is an Indicator of Poor Prognosis. <i>American Journal of Pathology</i> , 2012, 181, 1762-1772. | 3.8 | 73 |
| 137 | Intellectual Outcome in Molecular Subgroups of Medulloblastoma. <i>Journal of Clinical Oncology</i> , 2016, 34, 4161-4170. | 1.6 | 72 |
| 138 | Targeting the enhancer of zeste homologue 2 in medulloblastoma. <i>International Journal of Cancer</i> , 2012, 131, 1800-1809. | 5.1 | 71 |
| 139 | Shh Signaling Protects Atoh1 from Degradation Mediated by the E3 Ubiquitin Ligase Huwe1 in Neural Precursors. <i>Developmental Cell</i> , 2014, 29, 649-661. | 7.0 | 71 |
| 140 | The Genetics of Pediatric Brain Tumors. <i>Current Neurology and Neuroscience Reports</i> , 2010, 10, 215-223. | 4.2 | 69 |
| 141 | Genetic and Epigenetic Inactivation of Kruppel-like Factor 4 in Medulloblastoma. <i>Neoplasia</i> , 2010, 12, 20-27. | 5.3 | 69 |
| 142 | Rapid Diagnosis of Medulloblastoma Molecular Subgroups. <i>Clinical Cancer Research</i> , 2011, 17, 1883-1894. | 7.0 | 69 |
| 143 | Polo-like kinase 1 (PLK1) inhibition suppresses cell growth and enhances radiation sensitivity in medulloblastoma cells. <i>BMC Cancer</i> , 2012, 12, 80. | 2.6 | 69 |
| 144 | A C19MC-LIN28A-MYCN Oncogenic Circuit Driven by Hijacked Super-enhancers Is a Distinct Therapeutic Vulnerability in ETMRs: A Lethal Brain Tumor. <i>Cancer Cell</i> , 2019, 36, 51-67.e7. | 16.8 | 69 |

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|-----|---|------|-----------|
| 145 | Voltage-gated potassium channel EAG2 controls mitotic entry and tumor growth in medulloblastoma via regulating cell volume dynamics. <i>Genes and Development</i> , 2012, 26, 1780-1796. | 5.9 | 68 |
| 146 | Hypermethylation of the Inactive X Chromosome Is a Frequent Event in Cancer. <i>Cell</i> , 2013, 155, 567-581. | 28.9 | 67 |
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