

# Ulrich SchÄ¼ller

## List of Publications by Year in descending order

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Version: 2024-02-01

102  
papers

10,089  
citations

101543

36  
h-index

37204

96  
g-index

104  
all docs

104  
docs citations

104  
times ranked

11796  
citing authors

#	ARTICLE	IF	CITATIONS
1	Primary central nervous system sarcoma with <i>DICER1</i> mutation—treatment results of a novel molecular entity in pediatric Peruvian patients. <i>Cancer</i> , 2022, 128, 697-707.	4.1	14
2	Machine learning models predict the primary sites of head and neck squamous cell carcinoma metastases based on <i>DNA</i> methylation. <i>Journal of Pathology</i> , 2022, 256, 378-387.	4.5	19
3	ALK inhibition as a salvage therapy for a relapsed unclassifiable sarcomatous CNS tumor with EML4/ALK fusion in an infant. <i>Pediatric Blood and Cancer</i> , 2022, 69, e29594.	1.5	0
4	A new amplicon-based gene panel for next generation sequencing characterization of meningiomas. <i>Brain Pathology</i> , 2022, 32, e13046.	4.1	7
5	Single-cell transcriptomics identifies potential cells of origin of MYC rhabdoid tumors. <i>Nature Communications</i> , 2022, 13, 1544.	12.8	9
6	Updates in the classification of ependymal neoplasms: The 2021 WHO Classification and beyond. <i>Brain Pathology</i> , 2022, 32, e13068.	4.1	29
7	Comprehensive profiling of myxopapillary ependymomas identifies a distinct molecular subtype with relapsing disease. <i>Neuro-Oncology</i> , 2022, 24, 1689-1699.	1.2	11
8	Increased replication stress and R-loop accumulation in EGFRvIII-expressing glioblastoma present new therapeutic opportunities. <i>Neuro-Oncology Advances</i> , 2022, 4, vdab180.	0.7	2
9	Disruption of GMNC-MCIDAS multiciliogenesis program is critical in choroid plexus carcinoma development. <i>Cell Death and Differentiation</i> , 2022, 29, 1596-1610.	11.2	7
10	SMARCB1-deficient and SMARCA4-deficient Malignant Brain Tumors With Complex Copy Number Alterations and TP53 Mutations May Represent the First Clinical Manifestation of Li-Fraumeni Syndrome. <i>American Journal of Surgical Pathology</i> , 2022, 46, 1277-1283.	3.7	3
11	DNA methylation subclass receptor tyrosine kinase II (RTK II) is predictive for seizure development in glioblastoma patients. <i>Neuro-Oncology</i> , 2022, 24, 1886-1897.	1.2	7
12	TBIO-07. Pediatric tumor classification through genome-wide methylation profiling of extracellular vesicle DNA. <i>Neuro-Oncology</i> , 2022, 24, i184-i184.	1.2	0
13	OLIG2 Is a Determinant for the Relapse of <i>MYC</i> -Amplified Medulloblastoma. <i>Clinical Cancer Research</i> , 2022, 28, 4278-4291.	7.0	3
14	Ependymoma relapse goes along with a relatively stable epigenome, but a severely altered tumor morphology. <i>Brain Pathology</i> , 2021, 31, 33-44.	4.1	8
15	Atypical teratoid/rhabdoid tumors (ATRTs) with SMARCA4 mutation are molecularly distinct from SMARCB1-deficient cases. <i>Acta Neuropathologica</i> , 2021, 141, 291-301.	7.7	47
16	Mutations within FGFR1 are associated with superior outcome in a series of 83 diffuse midline gliomas with H3F3A K27M mutations. <i>Acta Neuropathologica</i> , 2021, 141, 323-325.	7.7	20
17	DIMEimmune: Robust estimation of infiltrating lymphocytes in CNS tumors from DNA methylation profiles. <i>Oncimmunology</i> , 2021, 10, 1932365.	4.6	17
18	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

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19	The transcriptional landscape of Shh medulloblastoma. <i>Nature Communications</i> , 2021, 12, 1749.	12.8	47
20	Cauda equina paragangliomas express HOXB13. <i>Neuropathology and Applied Neurobiology</i> , 2021, 47, 889-890.	3.2	9
21	Neurofibromatosis type 2 predisposes to ependymomas of various localization, histology, and molecular subtype. <i>Acta Neuropathologica</i> , 2021, 141, 971-974.	7.7	12
22	Cross-Species Genomics Reveals Oncogenic Dependencies in ZFTA/C11orf95 Fusion-Positive Supratentorial Ependymomas. <i>Cancer Discovery</i> , 2021, 11, 2230-2247.	9.4	39
23	Histopathological patterns in atypical teratoid/rhabdoid tumors are related to molecular subgroup. <i>Brain Pathology</i> , 2021, 31, e12967.	4.1	16
24	SMARCA4-deficient rhabdoid tumours show intermediate molecular features between SMARCB1-deficient rhabdoid tumours and small cell carcinomas of the ovary, hypercalcaemic type. <i>Journal of Pathology</i> , 2021, 255, 1-15.	4.5	14
25	Evidence for a low-penetrant extended phenotype of rhabdoid tumor predisposition syndrome type 1 from a kindred with gain of SMARCB1 exon 6. <i>Pediatric Blood and Cancer</i> , 2021, 68, e29185.	1.5	0
26	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
27	Malignant gliomas with H3F3A G34R mutation or MYCN amplification in pediatric patients with Li Fraumeni syndrome. <i>Acta Neuropathologica</i> , 2021, 142, 591-593.	7.7	5
28	IDH2 R172 Mutations Across Poorly Differentiated Sinonasal Tract Malignancies. <i>American Journal of Surgical Pathology</i> , 2021, 45, 1190-1204.	3.7	26
29	Comprehensive molecular characterization of pediatric radiation-induced high-grade glioma. <i>Nature Communications</i> , 2021, 12, 5531.	12.8	31
30	Double adenomas of the pituitary reveal distinct lineage markers, copy number alterations, and epigenetic profiles. <i>Pituitary</i> , 2021, 24, 904-913.	2.9	4
31	Molecular profiling of pediatric meningiomas shows tumor characteristics distinct from adult meningiomas. <i>Acta Neuropathologica</i> , 2021, 142, 873-886.	7.7	12
32	Simultaneous Brg1 Knockout and MYCN Overexpression in Cerebellar Granule Neuron Precursors Is Insufficient to Drive Tumor Formation but Temporarily Enhances their Proliferation and Delays their Migration. <i>Cerebellum</i> , 2021, 20, 410-419.	2.5	4
33	Sarcoma classification by DNA methylation profiling. <i>Nature Communications</i> , 2021, 12, 498.	12.8	237
34	Local and systemic therapy of recurrent ependymoma in children and adolescents: short- and long-term results of the E-HIT-REZ 2005 study. <i>Neuro-Oncology</i> , 2021, 23, 1012-1023.	1.2	19
35	Systemic chemotherapy of pediatric recurrent ependymomas: results from the German HIT-REZ studies. <i>Journal of Neuro-Oncology</i> , 2021, 155, 193-202.	2.9	6
36	The H3.3K27M oncohistone affects replication stress outcome and provokes genomic instability in pediatric glioma. <i>PLoS Genetics</i> , 2021, 17, e1009868.	3.5	14

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37	Co-activation of Sonic hedgehog and Wnt signaling in murine retinal precursor cells drives ocular lesions with features of intraocular medulloepithelioma. <i>Oncogenesis</i> , 2021, 10, 78.	4.9	0
38	Brahma-related gene 1 has time-specific roles during brain and eye development. <i>Development (Cambridge)</i> , 2021, 148, .	2.5	5
39	Preferential sensitivity to HDAC inhibitors in tumors with CREBBP mutation. <i>Cancer Gene Therapy</i> , 2020, 27, 294-300.	4.6	29
40	Molecular characterization of histopathological ependymoma variants. <i>Acta Neuropathologica</i> , 2020, 139, 305-318.	7.7	43
41	Macrophage-tumor cell interaction promotes ATRT progression and chemoresistance. <i>Acta Neuropathologica</i> , 2020, 139, 913-936.	7.7	24
42	Treatment response of CNS high-grade neuroepithelial tumors with MN1 alteration. <i>Pediatric Blood and Cancer</i> , 2020, 67, e28627.	1.5	5
43	An 8-year-old Girl with Posterior Fossa Mass. <i>Brain Pathology</i> , 2020, 30, 713-714.	4.1	1
44	Genome-wide DNA methylation profiles distinguish silent from non-silent ACTH adenomas. <i>Acta Neuropathologica</i> , 2020, 140, 95-97.	7.7	7
45	Molecular profiling of an osseous metastasis in glioblastoma during checkpoint inhibition: potential mechanisms of immune escape. <i>Acta Neuropathologica Communications</i> , 2020, 8, 28.	5.2	24
46	EGFRVIII upregulates DNA mismatch repair resulting in increased temozolomide sensitivity of MGMT promoter methylated glioblastoma. <i>Oncogene</i> , 2020, 39, 3041-3055.	5.9	42
47	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
48	OUP accepted manuscript. <i>Cerebral Cortex</i> , 2020, 30, 1382-1392.	2.9	4
49	Immunologic Profiling of Mutational and Transcriptional Subgroups in Pediatric and Adult High-Grade Gliomas. <i>Cancer Immunology Research</i> , 2019, 7, 1401-1411.	3.4	35
50	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
51	Gliosarcoma Is Driven by Alterations in PI3K/Akt, RAS/MAPK Pathways and Characterized by Collagen Gene Expression Signature. <i>Cancers</i> , 2019, 11, 284.	3.7	18
52	TCF4 (E2-2) harbors tumor suppressive functions in SHH medulloblastoma. <i>Acta Neuropathologica</i> , 2019, 137, 657-673.	7.7	20
53	The transcriptional coactivator and histone acetyltransferase CBP regulates neural precursor cell development and migration. <i>Acta Neuropathologica Communications</i> , 2019, 7, 199.	5.2	13
54	The molecular landscape of ETMR at diagnosis and relapse. <i>Nature</i> , 2019, 576, 274-280.	27.8	94

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55	DNA methylation-based reclassification of olfactory neuroblastoma. <i>Acta Neuropathologica</i> , 2018, 136, 255-271.	7.7	59
56	Germline variants in SMARCB1 and other members of the BAF chromatin-remodeling complex across human disease entities: a meta-analysis. <i>European Journal of Human Genetics</i> , 2018, 26, 1083-1093.	2.8	30
57	DNA methylation-based classification of central nervous system tumours. <i>Nature</i> , 2018, 555, 469-474.	27.8	1,872
58	Opposing Effects of CREBBP Mutations Govern the Phenotype of Rubinstein-Taybi Syndrome and Adult SHH Medulloblastoma. <i>Developmental Cell</i> , 2018, 44, 709-724.e6.	7.0	35
59	Identification of time-to-peak on dynamic 18F-FET-PET as a prognostic marker specifically in IDH1/2 mutant diffuse astrocytoma. <i>Neuro-Oncology</i> , 2018, 20, 279-288.	1.2	71
60	Group 3 medulloblastoma in a patient with a GYS2 germline mutation and glycogen storage disease 0a. <i>Child's Nervous System</i> , 2018, 34, 581-584.	1.1	2
61	Subgroup-specific immune and stromal microenvironment in medulloblastoma. <i>Oncolmmunology</i> , 2018, 7, e1462430.	4.6	77
62	Heterogeneity within the PF-EPN-B ependymoma subgroup. <i>Acta Neuropathologica</i> , 2018, 136, 227-237.	7.7	86
63	Primary intracranial spindle cell sarcoma with rhabdomyosarcoma-like features share a highly distinct methylation profile and DICER1 mutations. <i>Acta Neuropathologica</i> , 2018, 136, 327-337.	7.7	104
64	Overexpression of Lin28b in Neural Stem Cells is Insufficient for Brain Tumor Formation, but Induces Pathological Lobulation of the Developing Cerebellum. <i>Cerebellum</i> , 2017, 16, 122-131.	2.5	14
65	Spatial heterogeneity in medulloblastoma. <i>Nature Genetics</i> , 2017, 49, 780-788.	21.4	112
66	A mouse model for embryonal tumors with multilayered rosettes uncovers the therapeutic potential of Sonic-hedgehog inhibitors. <i>Nature Medicine</i> , 2017, 23, 1191-1202.	30.7	38
67	Medulloblastoma: experimental models and reality. <i>Acta Neuropathologica</i> , 2017, 134, 679-689.	7.7	25
68	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
69	Integrated (epi)-Genomic Analyses Identify Subgroup-Specific Therapeutic Targets in CNS Rhabdoid Tumors. <i>Cancer Cell</i> , 2016, 30, 891-908.	16.8	191
70	Somatostatin-receptor-targeted radionuclide therapy for progressive meningioma: benefit linked to <sup>68</sup> Ga-DOTATATE/TOC uptake. <i>Neuro-Oncology</i> , 2016, 18, now060.	1.2	79
71	Divergent clonal selection dominates medulloblastoma at recurrence. <i>Nature</i> , 2016, 529, 351-357.	27.8	266
72	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

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73	Distinct Histomorphology in Molecular Subgroups of Glioblastomas in Young Patients. <i>Journal of Neuropathology and Experimental Neurology</i> , 2016, 75, 408-414.	1.7	35
74	Sonic Hedgehog promotes proliferation of Notch-dependent monociliated choroid plexus tumour cells. <i>Nature Cell Biology</i> , 2016, 18, 418-430.	10.3	59
75	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
76	New Brain Tumor Entities Emerge from Molecular Classification of CNS-PNETs. <i>Cell</i> , 2016, 164, 1060-1072.	28.9	702
77	Histologically distinct neuroepithelial tumors with histone 3 G34 mutation are molecularly similar and comprise a single nosologic entity. <i>Acta Neuropathologica</i> , 2016, 131, 137-146.	7.7	162
78	Canonical Wnt Signaling Drives Tumor-Like Lesions from Sox2-Positive Precursors of the Murine Olfactory Epithelium. <i>PLoS ONE</i> , 2016, 11, e0166690.	2.5	7
79	Characterization of pancreatic glucagon-producing tumors and pituitary gland tumors in transgenic mice overexpressing <i>MYCN</i> in <i>hGFAP</i> -positive cells. <i>Oncotarget</i> , 2016, 7, 74415-74426.	1.8	21
80	Comparative integrated molecular analysis of intraocular medulloepitheliomas and central nervous system embryonal tumors with multilayered rosettes confirms that they are distinct nosologic entities. <i>Neuropathology</i> , 2015, 35, 538-544.	1.2	38
81	Histological subtype of medulloblastoma frequently changes upon recurrence. <i>Acta Neuropathologica</i> , 2015, 129, 459-461.	7.7	10
82	Wnt/ $\beta$ -catenin signaling inhibits the Shh pathway and impairs tumor growth in Shh-dependent medulloblastoma. <i>Acta Neuropathologica</i> , 2014, 127, 605-607.	7.7	33
83	Subgroup-specific localization of human medulloblastoma based on pre-operative MRI. <i>Acta Neuropathologica</i> , 2014, 127, 931-933.	7.7	53
84	Genome Sequencing of SHH Medulloblastoma Predicts Genotype-Related Response to Smoothed Inhibition. <i>Cancer Cell</i> , 2014, 25, 393-405.	16.8	627
85	Loss of SmarC Proteins Impairs Cerebellar Development. <i>Journal of Neuroscience</i> , 2014, 34, 13486-13491.	3.6	36
86	SMARCA4-mutated atypical teratoid/rhabdoid tumors are associated with inherited germline alterations and poor prognosis. <i>Acta Neuropathologica</i> , 2014, 128, 453-456.	7.7	155
87	Genomic and transcriptomic analyses match medulloblastoma mouse models to their human counterparts. <i>Acta Neuropathologica</i> , 2014, 128, 123-136.	7.7	54
88	Recurrence patterns across medulloblastoma subgroups: an integrated clinical and molecular analysis. <i>Lancet Oncology</i> , The, 2013, 14, 1200-1207.	10.7	307
89	Constitutive activation of $\beta$ -Catenin in neural progenitors results in disrupted proliferation and migration of neurons within the central nervous system. <i>Developmental Biology</i> , 2013, 374, 319-332.	2.0	37
90	TERT promoter mutations are highly recurrent in SHH subgroup medulloblastoma. <i>Acta Neuropathologica</i> , 2013, 126, 917-929.	7.7	146

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91	LIN28A immunoreactivity is a potent diagnostic marker of embryonal tumor with multilayered rosettes (ETMR). <i>Acta Neuropathologica</i> , 2012, 124, 875-881.	7.7	115
92	Dissecting the genomic complexity underlying medulloblastoma. <i>Nature</i> , 2012, 488, 100-105.	27.8	765
93	Proper cerebellar development requires expression of $\beta$ 1-integrin in Bergmann glia, but not in granule neurons. <i>Glia</i> , 2012, 60, 820-832.	4.9	26
94	Sonic hedgehog-associated medulloblastoma arising from the cochlear nuclei of the brainstem. <i>Acta Neuropathologica</i> , 2012, 123, 601-614.	7.7	71
95	Severe Alterations of Cerebellar Cortical Development after Constitutive Activation of Wnt Signaling in Granule Neuron Precursors. <i>Molecular and Cellular Biology</i> , 2011, 31, 3326-3338.	2.3	55
96	Dexamethasone Destabilizes Nmyc to Inhibit the Growth of Hedgehog-Associated Medulloblastoma. <i>Cancer Research</i> , 2010, 70, 5220-5225.	0.9	19
97	Medulloblastoma Can Be Initiated by Deletion of Patched in Lineage-Restricted Progenitors or Stem Cells. <i>Cancer Cell</i> , 2008, 14, 135-145.	16.8	606
98	Acquisition of Granule Neuron Precursor Identity Is a Critical Determinant of Progenitor Cell Competence to Form Shh-Induced Medulloblastoma. <i>Cancer Cell</i> , 2008, 14, 123-134.	16.8	572
99	Forkhead Transcription Factor FoxM1 Regulates Mitotic Entry and Prevents Spindle Defects in Cerebellar Granule Neuron Precursors. <i>Molecular and Cellular Biology</i> , 2007, 27, 8259-8270.	2.3	84
100	$\beta$ -catenin function is required for cerebellar morphogenesis. <i>Brain Research</i> , 2007, 1140, 161-169.	2.2	46
101	Cerebellar $\beta$ -transcriptome™ reveals cell-type and stage-specific expression during postnatal development and tumorigenesis. <i>Molecular and Cellular Neurosciences</i> , 2006, 33, 247-259.	2.2	42
102	Insulin-Like Growth Factor II Is Involved in the Proliferation Control of Medulloblastoma and Its Cerebellar Precursor Cells. <i>American Journal of Pathology</i> , 2005, 166, 1153-1162.	3.8	106