Krzysztof Zakrzewski

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Hotspot Mutations in H3F3A and IDH1 Define Distinct Epigenetic and Biological Subgroups of Glioblastoma. Cancer Cell, 2012, 22, 425-437.	16.8	1,551
2	Fusion of TTYH1 with the C19MC microRNA cluster drives expression of a brain-specific DNMT3B isoform in the embryonal brain tumor ETMR. Nature Genetics, 2014, 46, 39-44.	21.4	167
3	Germline and somatic FGFR1 abnormalities in dysembryoplastic neuroepithelial tumors. Acta Neuropathologica, 2016, 131, 847-863.	7.7	143
4	Arrested neural and advanced mesenchymal differentiation of glioblastoma cells-comparative study with neural progenitors. BMC Cancer, 2009, 9, 54.	2.6	40
5	Glioblastoma-derived spheroid cultures as an experimental model for analysis of EGFR anomalies. Journal of Neuro-Oncology, 2011, 102, 395-407.	2.9	27
6	Mutational analysis of hSNF5/INI1 and TP53 genes in choroid plexus carcinomas. Cancer Genetics and Cytogenetics, 2005, 156, 179-182.	1.0	25
7	Transcriptional profiles of pilocytic astrocytoma are related to their three different locations, but not to radiological tumor features. BMC Cancer, 2015, 15, 778.	2.6	22
8	Altered MicroRNA Expression Is Associated with Tumor Grade, Molecular Background and Outcome in Childhood Infratentorial Ependymoma. PLoS ONE, 2016, 11, e0158464.	2.5	20
9	Polycomb genes expression as a predictor of poor clinical outcome in children with medulloblastoma. Child's Nervous System, 2011, 27, 79-86.	1.1	17
10	Cranial and ventricular size following shunting or endoscopic third ventriculostomy (ETV) in infants with aqueductal stenosis: further insights from the International Infant Hydrocephalus Study (IIHS). Child's Nervous System, 2020, 36, 1407-1414.	1.1	15
11	Influence of Neuroendoscopic Third Ventriculostomy on the Size of Ventricles in Chronic Hydrocephalus. Journal of Child Neurology, 2004, 19, 579-587.	1.4	14
12	mRNA and miRNA Expression Analyses of the MYC/E2F/miR-17-92 Network in the Most Common Pediatric Brain Tumors. International Journal of Molecular Sciences, 2021, 22, 543.	4.1	11
13	Prevalence of mutated TP53 on cDNA (but not on DNA template) in pleomorphic xanthoastrocytoma with positive TP53 immunohistochemistry. Cancer Genetics and Cytogenetics, 2009, 193, 93-97.	1.0	9
14	Expression-based decision tree model reveals distinct microRNA expression pattern in pediatric neuronal and mixed neuronal-glial tumors. BMC Cancer, 2019, 19, 544.	2.6	9
15	Pilocytic astrocytoma as a predominant component of a recurrent complex type DNT. Folia Neuropathologica, 2009, 47, 284-8.	1.2	9
16	Expression of SOX11, PAX5, TTF-1 and ISL-1 in medulloblastoma. Pathology Research and Practice, 2016, 212, 965-971.	2.3	8
17	Sensitive detection of <i>FGFR1 N546K</i> mosaic mutation in patient with encephalocraniocutaneous lipomatosis and pilocytic astrocytoma. American Journal of Medical Genetics, Part A, 2019, 179, 1622-1627.	1.2	7
18	Novel Gene Expression Model for Outcome Prediction in Paediatric Medulloblastoma. Journal of Molecular Neuroscience, 2013, 51, 371-379.	2.3	6

#	Article	IF	CITATIONS
19	A hybrid technique for ventriculoatrial shunt implantation—technical note. Child's Nervous System, 2014, 30, 1729-1732.	1.1	5
20	The oscillatory flow of the cerebrospinal fluid in the Sylvian aqueduct and the prepontine cistern measured with phase contrast MRI in children with hydrocephalus—a preliminary report. Child's Nervous System, 2018, 34, 845-851.	1.1	5
21	microRNA interaction with MAPK and AKT pathways in paediatric brain tumours – preliminary results and review of the literature. Folia Neuropathologica, 2020, 58, 123-132.	1.2	5
22	Predicting conservative treatment failure in patients with lumbar disc herniation. Single center, case-control study. Clinical Neurology and Neurosurgery, 2020, 193, 105867.	1.4	5
23	Hedgehog signalling network gene status analysis in paediatric intracranial germ cell tumours. Folia Neuropathologica, 2019, 57, 227-238.	1.2	3