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List of Publications by Year in descending order

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29
papers

1,281
citations

361413

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477307

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docs citations

29
times ranked

2085
citing authors

#	ARTICLE	IF	CITATIONS
1	Tumor infiltrating immune cells in gliomas and meningiomas. <i>Brain, Behavior, and Immunity</i> , 2016, 53, 1-15.	4.1	228
2	Molecular and Genomic Alterations in Glioblastoma Multiforme. <i>American Journal of Pathology</i> , 2015, 185, 1820-1833.	3.8	141
3	Proposal for a new risk stratification classification for meningioma based on patient age, WHO tumor grade, size, localization, and karyotype. <i>Neuro-Oncology</i> , 2014, 16, 735-747.	1.2	98
4	Early recurrences in histologically benign/grade I meningiomas are associated with large tumors and coexistence of monosomy 14 and del(1p36) in the ancestral tumor cell clone. <i>Neuro-Oncology</i> , 2007, 9, 438-446.	1.2	74
5	New Classification Scheme for the Prognostic Stratification of Meningioma on the Basis of Chromosome 14 Abnormalities, Patient Age, and Tumor Histopathology. <i>Journal of Clinical Oncology</i> , 2003, 21, 3285-3295.	1.6	71
6	Genetic/molecular alterations of meningiomas and the signaling pathways targeted. <i>Oncotarget</i> , 2015, 6, 10671-10688.	1.8	58
7	Detailed Characterization of Alterations of Chromosomes 7, 9, and 10 in Glioblastomas as Assessed by Single-Nucleotide Polymorphism Arrays. <i>Journal of Molecular Diagnostics</i> , 2011, 13, 634-647.	2.8	55
8	Microarray-Based Analysis of Spinal versus Intracranial Meningiomas: Different Clinical, Biological, and Genetic Characteristics Associated with Distinct Patterns of Gene Expression. <i>Journal of Neuropathology and Experimental Neurology</i> , 2006, 65, 445-454.	1.7	48
9	Gene expression profiles of human glioblastomas are associated with both tumor cytogenetics and histopathology. <i>Neuro-Oncology</i> , 2010, 12, 991-1003.	1.2	45
10	Intratumoral Patterns of Clonal Evolution in Meningiomas as Defined by Multicolor Interphase Fluorescence in Situ Hybridization (FISH). <i>Journal of Molecular Diagnostics</i> , 2004, 6, 316-325.	2.8	42
11	Gene Expression Profiles of Meningiomas are Associated with Tumor Cytogenetics and Patient Outcome. <i>Brain Pathology</i> , 2009, 19, 409-420.	4.1	42
12	Amplified and Homozygously Deleted Genes in Glioblastoma: Impact on Gene Expression Levels. <i>PLoS ONE</i> , 2012, 7, e46088.	2.5	38
13	The Cytogenetic Relationship between Primary and Recurrent Meningiomas Points to the Need for New Treatment Strategies in Cases at High Risk of Relapse. <i>Clinical Cancer Research</i> , 2006, 12, 772-780.	7.0	37
14	Patient Gender Is Associated with Distinct Patterns of Chromosomal Abnormalities and Sex Chromosome-Linked Gene-Expression Profiles in Meningiomas. <i>Oncologist</i> , 2007, 12, 1225-1236.	3.7	37
15	Gains of chromosome 22 by fluorescence in situ hybridization in the context of an hyperdiploid karyotype are associated with aggressive clinical features in meningioma patients. <i>Cancer</i> , 2001, 92, 377-385.	4.1	34
16	Characterization of Chromosome 14 Abnormalities by Interphase In Situ Hybridization and Comparative Genomic Hybridization in 124 Meningiomas. <i>American Journal of Clinical Pathology</i> , 2005, 123, 744-751.	0.7	34
17	Association between Inflammatory Infiltrates and Isolated Monosomy 22/del(22q) in Meningiomas. <i>PLoS ONE</i> , 2013, 8, e74798.	2.5	31
18	Incidence of numerical chromosome aberrations in meningioma tumors as revealed by fluorescence in situ hybridization using 10 chromosome-specific probes. <i>Cytometry</i> , 2002, 50, 153-159.	1.8	28

#	ARTICLE	IF	CITATIONS
19	Tumor cell and immune cell profiles in primary human glioblastoma: Impact on patient outcome. <i>Brain Pathology</i> , 2021, 31, 365-380.	4.1	27
20	Intratumoral patterns of clonal evolution in gliomas. <i>Neurogenetics</i> , 2010, 11, 227-239.	1.4	23
21	Association between mutation of the NF2 gene and monosomy 22 in menopausal women with sporadic meningiomas. <i>BMC Medical Genetics</i> , 2013, 14, 114.	2.1	16
22	Delineation of commonly deleted chromosomal regions in meningiomas by high-density single nucleotide polymorphism genotyping arrays. <i>Genes Chromosomes and Cancer</i> , 2012, 51, 606-617.	2.8	15
23	Heterogeneous EGFR, CDK4, MDM4, and PDGFRA Gene Expression Profiles in Primary GBM: No Association with Patient Survival. <i>Cancers</i> , 2020, 12, 231.	3.7	13
24	Multivariate analysis reveals differentially expressed genes among distinct subtypes of diffuse astrocytic gliomas: diagnostic implications. <i>Scientific Reports</i> , 2020, 10, 11270.	3.3	11
25	The protein expression profile of meningioma cells is associated with distinct cytogenetic tumour subgroups. <i>Neuropathology and Applied Neurobiology</i> , 2015, 41, 319-332.	3.2	10
26	Characterization of chromosome 14 abnormalities by interphase in situ hybridization and comparative genomic hybridization in 124 meningiomas: correlation with clinical, histopathologic, and prognostic features. <i>American Journal of Clinical Pathology</i> , 2005, 123, 744-51.	0.7	9
27	Array-based comparative genomic hybridization of mapped BAC DNA clones to screen for chromosome 14 copy number abnormalities in meningiomas. <i>European Journal of Human Genetics</i> , 2008, 16, 1450-1458.	2.8	6
28	Prognostic stratification of adult primary glioblastoma multiforme patients based on their tumor gene amplification profiles. <i>Oncotarget</i> , 2018, 9, 28083-28102.	1.8	5
29	Whole-Exome Sequencing Reveals Recurrent but Heterogeneous Mutational Profiles in Sporadic WHO Grade 1 Meningiomas. <i>Frontiers in Oncology</i> , 2021, 11, 740782.	2.8	5