Mohammed Bangash

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1790288/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Comprehensive molecular biomarker identification in breast cancer brain metastases. Journal of Translational Medicine, 2017, 15, 269.	4.4	80
2	Microarray Expression Data Identify DCC as a Candidate Gene for Early Meningioma Progression. PLoS ONE, 2016, 11, e0153681.	2.5	40
3	Pleomorphism and drug resistant cancer stem cells are characteristic of aggressive primary meningioma cell lines. Cancer Cell International, 2017, 17, 72.	4.1	21
4	In situ characterization of stem cells-like biomarkers in meningiomas. Cancer Cell International, 2018, 18, 77.	4.1	15
5	Epidemiological Trends of Histopathologically WHO Classified CNS Tumors in Developing Countries: Systematic Review. Asian Pacific Journal of Cancer Prevention, 2015, 16, 205-216.	1.2	10
6	Microarray expression profiling identifies genes, including cytokines, and biofunctions, as diapedesis, associated with a brain metastasis from a papillary thyroid carcinoma. American Journal of Cancer Research, 2016, 6, 2140-2161.	1.4	10
7	Malignant Transformation and Spine Metastasis of an Intracranial Grade I Meningioma: In Situ Immunofluorescence Analysis of Cancer Stem Cells Case Report and Literature Review. World Neurosurgery, 2018, 120, 274-289.	1.3	5
8	Overlapping variants in the blood, tissues and cell lines for patients with intracranial meningiomas are predominant in stem cell-related genes. Heliyon, 2020, 6, e05632.	3.2	4