

Yu Mei

List of Publications by Year in descending order

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papers

701
citations

759233

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#	ARTICLE	IF	CITATIONS
1	Immune profiling of pituitary tumors reveals variations in immune infiltration and checkpoint molecule expression. <i>Pituitary</i> , 2021, 24, 359-373.	2.9	12
2	Immune Microenvironment of Vestibular Schwannomas. <i>Journal of Neurological Surgery, Part B: Skull Base</i> , 2018, 79, S1-S188.	0.8	1
3	Immune Microenvironment of Pituitary Adenomas. <i>Journal of Neurological Surgery, Part B: Skull Base</i> , 2018, 79, S1-S188.	0.8	0
4	Clinical Identification of Oncogenic Drivers and Copy-Number Alterations in Pituitary Tumors. <i>Endocrinology</i> , 2017, 158, 2284-2291.	2.8	53
5	Genomic landscape of high-grade meningiomas. <i>Npj Genomic Medicine</i> , 2017, 2, .	3.8	130
6	Landscape of Genomic Alterations in Pituitary Adenomas. <i>Clinical Cancer Research</i> , 2017, 23, 1841-1851.	7.0	94
7	Osteoglycin promotes meningioma development through downregulation of NF2 and activation of mTOR signaling. <i>Cell Communication and Signaling</i> , 2017, 15, 34.	6.5	21
8	Genomic profile of human meningioma cell lines. <i>PLoS ONE</i> , 2017, 12, e0178322.	2.5	44
9	Increased expression of programmed death ligand 1 (PD-L1) in human pituitary tumors. <i>Oncotarget</i> , 2016, 7, 76565-76576.	1.8	100
10	Meningioma Genomics: Diagnostic, Prognostic, and Therapeutic Applications. <i>Frontiers in Surgery</i> , 2016, 3, 40.	1.4	70
11	Genomic landscape of intracranial meningiomas. <i>Journal of Neurosurgery</i> , 2016, 125, 525-535.	1.6	104
12	Genomic and Epigenomic Landscape in Meningioma. <i>Neurosurgery Clinics of North America</i> , 2016, 27, 167-179.	1.7	31
13	High incidence of TERT mutation in brain tumor cell lines. <i>Brain Tumor Pathology</i> , 2016, 33, 222-227.	1.7	26