

Viktória Koroknai

List of Publications by Year in descending order

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#	ARTICLE	IF	CITATIONS
1	Cytokine and Chemokine Receptor Patterns of Human Malignant Melanoma Cell Lines. International Journal of Molecular Sciences, 2022, 23, 2644.	4.1	3
2	Cell Proliferation Is Strongly Associated with the Treatment Conditions of an ER Stress Inducer New Anti-Melanoma Drug in Melanoma Cell Lines. Biomedicines, 2021, 9, 96.	3.2	5
3	Silencing Osteopontin Expression Inhibits Proliferation, Invasion and Induce Altered Protein Expression in Melanoma Cells. Pathology and Oncology Research, 2021, 27, 581395.	1.9	7
4	Molecular Alterations Associated with Acquired Drug Resistance during Combined Treatment with Encorafenib and Binimetinib in Melanoma Cell Lines. Cancers, 2021, 13, 6058.	3.7	7
5	DNA hypermethylation is associated with invasive phenotype of malignant melanoma. Experimental Dermatology, 2020, 29, 39-50.	2.9	30
6	Gene Expression Signature of BRAF Inhibitor Resistant Melanoma Spheroids. Pathology and Oncology Research, 2020, 26, 2557-2566.	1.9	5
7	Molecular alterations associated with acquired resistance to BRAFV600E targeted therapy in melanoma cells. Melanoma Research, 2019, 29, 390-400.	1.2	14
8	Detection of CCND1 Locus Amplification by Fluorescence In Situ Hybridization. Methods in Molecular Biology, 2018, 1726, 85-100.	0.9	3
9	Altered integrin expression patterns shown by microarray in human cutaneous melanoma. Melanoma Research, 2017, 27, 180-188.	1.2	20
10	Genomic profiling of invasive melanoma cell lines by array comparative genomic hybridization. Melanoma Research, 2016, 26, 100-107.	1.2	9
11	The role of osteopontin expression in melanoma progression. Tumor Biology, 2015, 36, 7841-7847.	1.8	23
12	DNA Methylation Characteristics of Primary Melanomas with Distinct Biological Behaviour. PLoS ONE, 2014, 9, e96612.	2.5	27
13	The role of CCND1 alterations during the progression of cutaneous malignant melanoma. Tumor Biology, 2012, 33, 2189-2199.	1.8	45