Shu-Huei Hsiao

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/820253/publications.pdf

Version: 2024-02-01

9 papers	222 citations	1478505 6 h-index	1474206 9 g-index
9 all docs	9 docs citations	9 times ranked	400 citing authors

#	Article	IF	Citations
1	Targeted Methylation of Two Tumor Suppressor Genes Is Sufficient to Transform Mesenchymal Stem Cells into Cancer Stem/Initiating Cells. Cancer Research, 2011, 71, 4653-4663.	0.9	91
2	DNA methylation of the Trip10 promoter accelerates mesenchymal stem cell lineage determination. Biochemical and Biophysical Research Communications, 2010, 400, 305-312.	2.1	45
3	Changes in DNA methylation are associated with the development of drug resistance in cervical cancer cells. Cancer Cell International, 2015, 15, 98.	4.1	44
4	Excavating relics of DNA methylation changes during the development of neoplasia. Seminars in Cancer Biology, 2009, 19, 198-208.	9.6	18
5	Early life ethanol exposure causes long-lasting disturbances in rat mesenchymal stem cells via epigenetic modifications. Biochemical and Biophysical Research Communications, 2014, 453, 338-344.	2.1	9
6	Methylation of the Tumor Suppressor Genes HIC1 and RassF1A Clusters Independently From the Methylation of Polycomb Target Genes in Colon Cancer. Annals of Surgical Oncology, 2017, 24, 578-585.	1.5	7
7	HIC1 and RassF1A Methylation Attenuates Tubulin Expression and Cell Stiffness in Cancer. International Journal of Molecular Sciences, 2018, 19, 2884.	4.1	4
8	JAK2V617F influences epigenomic changes in myeloproliferative neoplasms. Biochemical and Biophysical Research Communications, 2017, 494, 470-476.	2.1	3
9	Dysregulated HIC1 and RassF1A expression inÂvitro alters the cell cytoskeleton and exosomal Piwi-interacting RNA. Biochemical and Biophysical Research Communications, 2022, 594, 109-116.	2.1	1