

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