

# Meng-Chang Hsiao

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11236775/publications.pdf>

Version: 2024-02-01

11  
papers

281  
citations

1307594

7  
h-index

1372567

10  
g-index

11  
all docs

11  
docs citations

11  
times ranked

674  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genotype-Phenotype Correlation in NF1: Evidence for a More Severe Phenotype Associated with Missense Mutations Affecting NF1 Codons 844-848. <i>American Journal of Human Genetics</i> , 2018, 102, 69-87.	6.2	144
2	Risk Factors for Development of Acute Kidney Injury in Patients with Urinary Tract Infection. <i>PLoS ONE</i> , 2015, 10, e0133835.	2.5	40
3	Decoding NF1 Intragenic Copy-Number Variations. <i>American Journal of Human Genetics</i> , 2015, 97, 238-249.	6.2	24
4	Risk Factors for Development of Septic Shock in Patients with Urinary Tract Infection. <i>BioMed Research International</i> , 2015, 2015, 1-7.	1.9	23
5	Urolithiasis Is a Risk Factor for Uroseptic Shock and Acute Kidney Injury in Patients With Urinary Tract Infection. <i>Frontiers in Medicine</i> , 2019, 6, 288.	2.6	19
6	Palindrome-Mediated and Replication-Dependent Pathogenic Structural Rearrangements within the <i>NF1</i> Gene. <i>Human Mutation</i> , 2014, 35, 891-898.	2.5	13
7	Risk factors for uroseptic shock in hospitalized patients aged over 80 years with urinary tract infection. <i>Annals of Translational Medicine</i> , 2020, 8, 477-477.	1.7	9
8	Molecular profiling of CNS tumors for the treatment and management of disease. <i>Journal of Clinical Neuroscience</i> , 2020, 71, 311-315.	1.5	6
9	Molecular profiling of gynecologic cancers for treatment and management of disease “ demonstrating clinical significance using the AMP/ASCO/CAP guidelines for interpretation and reporting of somatic variants. <i>Cancer Genetics</i> , 2020, 242, 25-34.	0.4	2
10	PATH-46. DETECTING MISDIAGNOSED ATYPICAL TERATOID/RHABDOID TUMOR (ATRT) BY DNA METHYLATION-BASED TUMOR CLASSIFICATION. <i>Neuro-Oncology</i> , 2019, 21, vi153-vi154.	1.2	1
11	Alternative outcomes of pathogenic complex somatic structural variations in the genomes of NF1 and NF2 patients. <i>Neurogenetics</i> , 2017, 18, 169-174.	1.4	0