## Yih-Chii Hwang

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/11911471/publications.pdf

Version: 2024-02-01

		1937685	2053705	
5	216	4	5	
papers	citations	h-index	g-index	
			202	
6	6	6	393	
all docs	docs citations	times ranked	citing authors	

#	Article	IF	CITATIONS
1	Predicting essential genes based on network and sequence analysis. Molecular BioSystems, 2009, 5, 1672.	2.9	109
2	HIPPIE: a high-throughput identification pipeline for promoter interacting enhancer elements. Bioinformatics, 2015, 31, 1290-1292.	4.1	52
3	McEnhancer: predicting gene expression via semi-supervised assignment of enhancers to target genes. Genome Biology, 2017, 18, 199.	8.8	27
4	High-throughput identification of long-range regulatory elements and their target promoters in the human genome. Nucleic Acids Research, 2013, 41, 4835-4846.	14.5	26
5	HIPPIE2: a method for fine-scale identification of physically interacting chromatin regions. NAR Genomics and Bioinformatics, 2020, 2, Iqaa022.	3.2	2