Charles A S Banks

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2588153/publications.pdf

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		1040056	1199594	
13	371	9	12	
papers	citations	h-index	g-index	
13	13	13	689	
all docs	docs citations	times ranked	citing authors	

#	Article	IF	CITATIONS
1	Multiple roles for PARP1 in ALC1-dependent nucleosome remodeling. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	3
2	Perturbation of BRMS1 interactome reveals pathways that impact metastasis. PLoS ONE, 2021, 16, e0259128.	2.5	0
3	Driving integrative structural modeling with serial capture affinity purification. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 31861-31870.	7.1	8
4	Differential Complex Formation via Paralogs in the Human Sin3 Protein Interaction Network. Molecular and Cellular Proteomics, 2020, 19, 1468-1484.	3.8	26
5	Integrative Modeling of a Sin3/HDAC Complex Sub-structure. Cell Reports, 2020, 31, 107516.	6.4	29
6	Purification and enzymatic assay of class I histone deacetylase enzymes. Methods in Enzymology, 2019, 626, 23-40.	1.0	4
7	A Structured Workflow for Mapping Human Sin3 Histone Deacetylase Complex Interactions Using Halo-MudPIT Affinity-Purification Mass Spectrometry. Molecular and Cellular Proteomics, 2018, 17, 1432-1447.	3.8	27
8	Differential HDAC1/2 network analysis reveals a role for prefoldin/CCT in HDAC1/2 complex assembly. Scientific Reports, 2018, 8, 13712.	3.3	32
9	Advancement of mass spectrometry-based proteomics technologies to explore triple negative breast cancer. Molecular BioSystems, 2017, 13, 42-55.	2.9	19
10	SONAR Discovers RNA-Binding Proteins from Analysis of Large-Scale Protein-Protein Interactomes. Molecular Cell, 2016, 64, 282-293.	9.7	155
11	Proteins interacting with cloning scars: a source of false positive protein-protein interactions. Scientific Reports, 2015, 5, 8530.	3.3	19
12	Affinity purification of protein complexes for analysis by multidimensional protein identification technology. Protein Expression and Purification, 2012, 86, 105-119.	1.3	24
13	Identification and Characterization of a Schizosaccharomyces pombe RNA Polymerase II Elongation Factor with Similarity to the Metazoan Transcription Factor ELL. Journal of Biological Chemistry, 2007, 282, 5761-5769.	3.4	25