

Charles A S Banks

List of Publications by Year in descending order

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

Version: 2024-02-01

13
papers

371
citations

1040056

9
h-index

1199594

12
g-index

13
all docs

13
docs citations

13
times ranked

689
citing authors

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