

Marco Faini

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1500483/publications.pdf>

Version: 2024-02-01

10
papers

1,359
citations

1040056

9
h-index

1281871

11
g-index

14
all docs

14
docs citations

14
times ranked

2604
citing authors

#	ARTICLE	IF	CITATIONS
1	A repository of assays to quantify 10,000 human proteins by SWATH-MS. <i>Scientific Data</i> , 2014, 1, 140031.	5.3	370
2	Crosslinking and Mass Spectrometry: An Integrated Technology to Understand the Structure and Function of Molecular Machines. <i>Trends in Biochemical Sciences</i> , 2016, 41, 20-32.	7.5	330
3	Structures of human PRC2 with its cofactors AEBP2 and JARID2. <i>Science</i> , 2018, 359, 940-944.	12.6	170
4	JARID2 and AEBP2 regulate PRC2 in the presence of H2AK119ub1 and other histone modifications. <i>Science</i> , 2021, 371, .	12.6	137
5	Systematic proteome and proteostasis profiling in human Trisomy 21 fibroblast cells. <i>Nature Communications</i> , 2017, 8, 1212.	12.8	112
6	Structural basis of AAUAAA polyadenylation signal recognition by the human CPSF complex. <i>Nature Structural and Molecular Biology</i> , 2018, 25, 135-138.	8.2	96
7	Structural insights into the assembly and polyA signal recognition mechanism of the human CPSF complex. <i>ELife</i> , 2017, 6, .	6.0	71
8	The Evolving Contribution of Mass Spectrometry to Integrative Structural Biology. <i>Journal of the American Society for Mass Spectrometry</i> , 2016, 27, 966-974.	2.8	36
9	Molecular architecture of the human tRNA ligase complex. <i>ELife</i> , 2021, 10, .	6.0	22
10	The MLL1 trimeric catalytic complex is a dynamic conformational ensemble stabilized by multiple weak interactions. <i>Nucleic Acids Research</i> , 2019, 47, 9433-9447.	14.5	8