

Thomas Menneteau

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8058002/publications.pdf>

Version: 2024-02-01

9
papers

269
citations

1162889

8
h-index

1474057

9
g-index

11
all docs

11
docs citations

11
times ranked

445
citing authors

#	ARTICLE	IF	CITATIONS
1	Deciphering preferential interactions within supramolecular protein complexes: the proteasome case. <i>Molecular Systems Biology</i> , 2015, 11, 771.	3.2	75
2	Comparison of label-free quantification methods for the determination of protein complexes subunits stoichiometry. <i>EuPA Open Proteomics</i> , 2014, 4, 82-86.	2.5	45
3	FAN1 controls mismatch repair complex assembly via MLH1 retention to stabilize CAG repeat expansion in Huntington's disease. <i>Cell Reports</i> , 2021, 36, 109649.	2.9	32
4	PIP30/FAM192A is a novel regulator of the nuclear proteasome activator PA28 β . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E6477-E6486.	3.3	29
5	Computational Strategies and Challenges for Using Native Ion Mobility Mass Spectrometry in Biophysics and Structural Biology. <i>Analytical Chemistry</i> , 2020, 92, 10872-10880.	3.2	24
6	Conformational maps of human 20S proteasomes reveal PA28- and immuno-dependent inter-ring crosstalks. <i>Nature Communications</i> , 2020, 11, 6140.	5.8	23
7	Mass Spectrometry-based Absolute Quantification of 20S Proteasome Status for Controlled Ex-vivo Expansion of Human Adipose-derived Mesenchymal Stromal/Stem Cells. <i>Molecular and Cellular Proteomics</i> , 2019, 18, 744-759.	2.5	17
8	Software Requirements for the Analysis and Interpretation of Native Ion Mobility Mass Spectrometry Data. <i>Analytical Chemistry</i> , 2020, 92, 10881-10890.	3.2	17
9	Proteasome complexes experience profound structural and functional rearrangements throughout mammalian spermatogenesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, e2116826119.	3.3	6