

Thomas Menneteau

List of Publications by Year in descending order

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Version: 2024-02-01

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papers

269
citations

1162889
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11
all docs

11
docs citations

11
times ranked

445
citing authors

#	ARTICLE	IF	CITATIONS
1	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
2	FAN1 controls mismatch repair complex assembly via MLH1 retention to stabilize CAG repeat expansion in Huntingtonâ€™s disease. <i>Cell Reports</i> , 2021, 36, 109649.	2.9	32
3	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
4	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
5	Conformational maps of human 20S proteasomes reveal PA28- and immuno-dependent inter-ring crosstalks. <i>Nature Communications</i> , 2020, 11, 6140.	5.8	23
6	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
7	PIP30/FAM192A is a novel regulator of the nuclear proteasome activator PA28 ^{Î³} . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E6477-E6486.	3.3	29
8	Deciphering preferential interactions within supramolecular protein complexes: the proteasome case. <i>Molecular Systems Biology</i> , 2015, 11, 771.	3.2	75
9	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