Sara C Larsen

List of Publications by Year in descending order

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759233 1199594 1,370 12 12 12 citations h-index g-index papers 16 16 16 2134 citing authors docs citations times ranked all docs

#	Article	IF	CITATION
1	An Optimized Shotgun Strategy for the Rapid Generation of Comprehensive Human Proteomes. Cell Systems, 2017, 4, 587-599.e4.	6.2	413
2	Proteome-wide analysis of arginine monomethylation reveals widespread occurrence in human cells. Science Signaling, 2016, 9, rs9.	3.6	241
3	Proteome-wide identification of the endogenous ADP-ribosylome of mammalian cells and tissue. Nature Communications, 2016, 7, 12917.	12.8	172
4	Systems-wide Analysis of Serine ADP-Ribosylation Reveals Widespread Occurrence and Site-Specific Overlap with Phosphorylation. Cell Reports, 2018, 24, 2493-2505.e4.	6.4	123
5	An Advanced Strategy for Comprehensive Profiling of ADP-ribosylation Sites Using Mass Spectrometry-based Proteomics*. Molecular and Cellular Proteomics, 2019, 18, 1010a-1026.	3.8	113
6	Mapping Physiological ADP-Ribosylation Using Activated Ion Electron Transfer Dissociation. Cell Reports, 2020, 32, 108176.	6.4	75
7	Treacle controls the nucleolar response to rDNA breaks via TOPBP1 recruitment and ATR activation. Nature Communications, 2020, 11 , 123 .	12.8	53
8	Biotin starvation causes mitochondrial protein hyperacetylation and partial rescue by the SIRT3-like deacetylase Hst4p. Nature Communications, 2015, 6, 7726.	12.8	47
9	The regulatory landscape of the human HPF1- and ARH3-dependent ADP-ribosylome. Nature Communications, 2021, 12, 5893.	12.8	45
10	Chemical genetics and proteome-wide site mapping reveal cysteine MARylation by PARP-7 on immune-relevant protein targets. ELife, 2021, 10 , .	6.0	43
11	Proteome-Wide Identification of In Vivo ADP-Ribose Acceptor Sites by Liquid Chromatography–Tandem Mass Spectrometry. Methods in Molecular Biology, 2017, 1608, 149-162.	0.9	24
12	Temporal and Site-Specific ADP-Ribosylation Dynamics upon Different Genotoxic Stresses. Cells, 2021, 10, 2927.	4.1	12