Sem Tamara

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5297449/publications.pdf

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687363 752698 22 674 13 20 citations h-index g-index papers 25 25 25 764 docs citations citing authors all docs times ranked

#	Article	IF	CITATIONS
1	High-Resolution Native Mass Spectrometry. Chemical Reviews, 2022, 122, 7269-7326.	47.7	164
2	Identifying glycation hot-spots in bovine milk proteins during production and storage of skim milk powder. International Dairy Journal, 2022, 129, 105340.	3.0	6
3	A perspective toward mass spectrometry-based de novo sequencing of endogenous antibodies. MAbs, 2022, 14, .	5.2	17
4	Generating Informative Sequence Tags from Antigen-Binding Regions of Heavily Glycosylated IgA1 Antibodies by Native Top-Down Electron Capture Dissociation. Journal of the American Society for Mass Spectrometry, 2021, 32, 1326-1335.	2.8	15
5	Human Milk from Previously COVID-19-Infected Mothers: The Effect of Pasteurization on Specific Antibodies and Neutralization Capacity. Nutrients, 2021, 13, 1645.	4.1	54
6	Structure of the human signal peptidase complex reveals the determinants for signal peptide cleavage. Molecular Cell, 2021, 81, 3934-3948.e11.	9.7	51
7	Human plasma IgG1 repertoires are simple, unique, and dynamic. Cell Systems, 2021, 12, 1131-1143.e5.	6.2	37
8	Single-particle mass analysis of intact ribosomes by mass photometry and Orbitrap-based charge detection mass spectrometry. IScience, 2021, 24, 103211.	4.1	22
9	Huntingtin structure is orchestrated by HAP40 and shows a polyglutamine expansion-specific interaction with exon 1. Communications Biology, 2021, 4, 1374.	4.4	22
10	Selectivity over coverage in <i>de novo</i> sequencing of IgGs. Chemical Science, 2020, 11, 11886-11896.	7.4	13
11	A wealth of genotype-specific proteoforms fine-tunes hemoglobin scavenging by haptoglobin. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 15554-15564.	7.1	31
12	How paired PSII–LHCII supercomplexes mediate the stacking of plant thylakoid membranes unveiled by structural mass-spectrometry. Nature Communications, 2020, 11, 1361.	12.8	57
13	Structural Proteomics Applied to Plant Membrane Protein Complexes. Trends in Plant Science, 2020, 25, 945-946.	8.8	O
14	Distinct Stabilities of the Structurally Homologous Heptameric Co-Chaperonins GroES and gp31. Journal of the American Society for Mass Spectrometry, 2019, 30, 7-15.	2.8	5
15	Expanding the mass range for UVPD-based native top-down mass spectrometry. Chemical Science, 2019, 10, 7163-7171.	7.4	29
16	A Colorful Pallet of B-Phycoerythrin Proteoforms Exposed by a Multimodal Mass Spectrometry Approach. CheM, 2019, 5, 1302-1317.	11.7	10
17	Dissecting ribosomal particles throughout the kingdoms of life using advanced hybrid mass spectrometry methods. Nature Communications, 2018, 9, 2493.	12.8	67
18	Phosphate Transfer in Activated Protein Complexes Reveals Interaction Sites. Angewandte Chemie, 2017, 129, 13829-13832.	2.0	2

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#	Article	IF	CITATION
19	Phosphate Transfer in Activated Protein Complexes Reveals Interaction Sites. Angewandte Chemie - International Edition, 2017, 56, 13641-13644.	13.8	4
20	Symmetry of Charge Partitioning in Collisional and UV Photon-Induced Dissociation of Protein Assemblies. Journal of the American Chemical Society, 2016, 138, 10860-10868.	13.7	42
21	Spatial distribution of metabolites in the human lens. Experimental Eye Research, 2016, 143, 68-74.	2.6	17
22	Human Plasma IgG1 Repertoires are Simple, Unique, and Dynamic. SSRN Electronic Journal, 0, , .	0.4	4