Daniel B Mcclatchy

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2532720/publications.pdf

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471509 526287 1,064 28 17 27 citations h-index g-index papers 31 31 31 1453 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	15N Metabolic Labeling of Mammalian Tissue with Slow Protein Turnover. Journal of Proteome Research, 2007, 6, 2005-2010.	3.7	144
2	Quantification of the synaptosomal proteome of the rat cerebellum during post-natal development. Genome Research, 2007, 17, 1378-1388.	5.5	99
3	Direct Detection of Biotinylated Proteins by Mass Spectrometry. Journal of Proteome Research, 2014, 13, 3966-3978.	3.7	96
4	Pulsed Azidohomoalanine Labeling in Mammals (PALM) Detects Changes in Liver-Specific LKB1 Knockout Mice. Journal of Proteome Research, 2015, 14, 4815-4822.	3.7	69
5	Impact of the Identification Strategy on the Reproducibility of the DDA and DIA Results. Journal of Proteome Research, 2020, 19, 3153-3161.	3.7	61
6	Census 2: isobaric labeling data analysis. Bioinformatics, 2014, 30, 2208-2209.	4.1	59
7	Intracellular amyloid toxicity induces oxytosis/ferroptosis regulated cell death. Cell Death and Disease, 2020, 11, 828.	6.3	59
8	Acute Synthesis of CPEB Is Required for Plasticity of Visual Avoidance Behavior in Xenopus. Cell Reports, 2014, 6, 737-747.	6.4	58
9	PSEA-Quant: A Protein Set Enrichment Analysis on Label-Free and Label-Based Protein Quantification Data. Journal of Proteome Research, 2014, 13, 5496-5509.	3.7	57
10	Amyloid Accumulation Drives Proteome-wide Alterations in Mouse Models of Alzheimer's Disease-like Pathology. Cell Reports, 2017, 21, 2614-2627.	6.4	56
11	Altered network and rescue of human neurons derived from individuals with early-onset genetic epilepsy. Molecular Psychiatry, 2021, 26, 7047-7068.	7.9	38
12	HILAQ: A Novel Strategy for Newly Synthesized Protein Quantification. Journal of Proteome Research, 2017, 16, 2213-2220.	3.7	29
13	Differential Proteomic Analysis of Mammalian Tissues Using SILAM. PLoS ONE, 2011, 6, e16039.	2.5	29
14	Quantitative analysis of newly synthesized proteins. Nature Protocols, 2018, 13, 1744-1762.	12.0	28
15	Elongation Factor 1A Family Regulates the Recycling of the M4 Muscarinic Acetylcholine Receptor. Neurochemical Research, 2006, 31, 975-988.	3.3	23
16	Quantitative temporal analysis of protein dynamics in cardiac remodeling. Journal of Molecular and Cellular Cardiology, 2018, 121, 163-172.	1.9	22
17	Role of the visual experience-dependent nascent proteome in neuronal plasticity. ELife, 2018, 7, .	6.0	19
18	Stable Isotope Labeling in Mammals (SILAM). Methods in Molecular Biology, 2014, 1156, 133-146.	0.9	19

#	Article	IF	Citations
19	Stable Isotope Labeling of Mammals (SILAM). Cold Spring Harbor Protocols, 2008, 2008, pdb.prot4940.	0.3	17
20	From Synapse to Function: A Perspective on the Role of Neuroproteomics in Elucidating Mechanisms of Drug Addiction. Proteomes, 2018, 6, 50.	3.5	16
21	The Retinal Ganglion Cell Transportome Identifies Proteins Transported to Axons and Presynaptic Compartments in the Visual System InÂVivo. Cell Reports, 2019, 28, 1935-1947.e5.	6.4	16
22	Quantitative analysis of global protein stability rates in tissues. Scientific Reports, 2020, 10, 15983.	3.3	13
23	Proteomic screen reveals diverse protein transport between connected neurons in the visual system. Cell Reports, 2022, 38, 110287.	6.4	10
24	Temporal Quantitative Profiling of Newly Synthesized Proteins during $\hat{Al^2}$ Accumulation. Journal of Proteome Research, 2021, 20, 763-775.	3.7	9
25	Improving Proteomics Data Reproducibility with a Dual-Search Strategy. Analytical Chemistry, 2020, 92, 1697-1701.	6.5	8
26	Structural Analysis of Hippocampal Kinase Signal Transduction. ACS Chemical Neuroscience, 2018, 9, 3072-3085.	3.5	6
27	Interactome analysis illustrates diverse gene regulatory processes associated with LIN28A in human iPS cell-derived neural progenitor cells. IScience, 2021, 24, 103321.	4.1	2
28	Proteomics INTegrator (PINT): An Online Tool To Store, Query, and Visualize Large Proteomics Experiment Results. Journal of Proteome Research, 2019, 18, 2999-3008.	3.7	O