

Jesus M Arizmendi

List of Publications by Year in descending order

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Version: 2024-02-01

12
papers

244
citations

1163117

8
h-index

1281871

11
g-index

12
all docs

12
docs citations

12
times ranked

482
citing authors

#	ARTICLE	IF	CITATIONS
1	Proteomic analysis of mycelium and secretome of different <i>Botrytis cinerea</i> wild-type strains. <i>Journal of Proteomics</i> , 2014, 97, 195-221.	2.4	74
2	Nuclear Phosphoproteomic Screen Uncovers ACLY as Mediator of IL-2-induced Proliferation of CD4+ T lymphocytes. <i>Molecular and Cellular Proteomics</i> , 2016, 15, 2076-2092.	3.8	40
3	Quantitative proteomics reveals neuronal ubiquitination of Rngo/Ddi1 and several proteasomal subunits by Ube3a, accounting for the complexity of Angelman syndrome. <i>Human Molecular Genetics</i> , 2018, 27, 1955-1971.	2.9	30
4	PAnalyzer: A software tool for protein inference in shotgun proteomics. <i>BMC Bioinformatics</i> , 2012, 13, 288.	2.6	28
5	In-Depth Proteomic Characterization of Classical and Non-Classical Monocyte Subsets. <i>Proteomes</i> , 2018, 6, 8.	3.5	18
6	Detailed Dissection of UBE3A-Mediated DDI1 Ubiquitination. <i>Frontiers in Physiology</i> , 2019, 10, 534.	2.8	17
7	Neuronal Proteomic Analysis of the Ubiquitinated Substrates of the Disease-Linked E3 Ligases Parkin and Ube3a. <i>BioMed Research International</i> , 2018, 2018, 1-14.	1.9	12
8	Multi-Omics Integration Highlights the Role of Ubiquitination in CCl4-Induced Liver Fibrosis. <i>International Journal of Molecular Sciences</i> , 2020, 21, 9043.	4.1	12
9	A Proteomic Approach for Systematic Mapping of Substrates of Human Deubiquitinating Enzymes. <i>International Journal of Molecular Sciences</i> , 2021, 22, 4851.	4.1	6
10	Composition of the ferredoxin-nitrite reductase from the cyanobacterium <i>Phormidium laminosum</i> . <i>Biochemical Society Transactions</i> , 1990, 18, 637-638.	3.4	3
11	Characterization of Receptor-Associated Protein Complex Assembly in Interleukin (IL)-2- and IL-15-Activated T-Cell Lines. <i>Journal of Proteome Research</i> , 2017, 16, 106-121.	3.7	3
12	Multi-laboratory experiment PME11 for the standardization of phosphoproteome analysis. <i>Journal of Proteomics</i> , 2022, 251, 104409.	2.4	1