

Jean-Baptiste Boyer

List of Publications by Year in descending order

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Version: 2024-02-01

11
papers

177
citations

1307594

7
h-index

1281871

11
g-index

11
all docs

11
docs citations

11
times ranked

250
citing authors

#	ARTICLE	IF	CITATIONS
1	Dual lysine and N-terminal acetyltransferases reveal the complexity underpinning protein acetylation. <i>Molecular Systems Biology</i> , 2020, 16, e9464.	7.2	53
2	O-GlcNAcylation site mapping by (azide-alkyne) click chemistry and mass spectrometry following intensive fractionation of skeletal muscle cells proteins. <i>Journal of Proteomics</i> , 2018, 186, 83-97.	2.4	27
3	NAA50 Is an Enzymatically Active ϵ -Acetyltransferase That Is Crucial for Development and Regulation of Stress Responses. <i>Plant Physiology</i> , 2020, 183, 1502-1516.	4.8	23
4	Integrative proteomic and phosphoproteomic profiling of prostate cell lines. <i>PLoS ONE</i> , 2019, 14, e0224148.	2.5	14
5	Lysine-specific acetylated proteome from the archaeon <i>Thermococcus gammatolerans</i> reveals the presence of acetylated histones. <i>Journal of Proteomics</i> , 2021, 232, 104044.	2.4	12
6	How may targeted proteomics complement genomic data in breast cancer?. <i>Expert Review of Proteomics</i> , 2017, 14, 43-54.	3.0	11
7	HYPK promotes the activity of the ϵ -acetyltransferase A complex to determine proteostasis of nonAc-X ² /N-degron-containing proteins. <i>Science Advances</i> , 2022, 8, .	10.3	11
8	N- and O-acetylation of threonine residues in the context of proteomics. <i>Journal of Proteomics</i> , 2014, 108, 369-372.	2.4	8
9	N-acetylation of secreted proteins in Apicomplexa is widespread and is independent of the ER acetyl-CoA transporter AT1. <i>Journal of Cell Science</i> , 2022, 135, .	2.0	7
10	A Continuous Assay Set to Screen and Characterize Novel Protein N-Acetyltransferases Unveils Rice General Control Non-repressible 5-Related N-Acetyltransferase2 Activity. <i>Frontiers in Plant Science</i> , 2022, 13, 832144.	3.6	6
11	Prioritizing targets for structural biology through the lens of proteomics: The archaeal protein TGAM_1934 from <i>Thermococcus gammatolerans</i> . <i>Proteomics</i> , 2015, 15, 114-123.	2.2	5