## Jean-Baptiste Boyer

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/127029/publications.pdf

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11	177	7	11
papers	citations	h-index	g-index
11	11	11	250 citing authors
all docs	docs citations	times ranked	

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