Christof Taxis

List of Publications by Citations

Source: https://exaly.com/author-pdf/455839/christof-taxis-publications-by-citations.pdf

Version: 2024-04-26

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

2,932 17 35 g-index

35 ext. papers ext. citations 5.6 avg, IF L-index

#	Paper	IF	Citations
33	A versatile toolbox for PCR-based tagging of yeast genes: new fluorescent proteins, more markers and promoter substitution cassettes. <i>Yeast</i> , 2004 , 21, 947-62	3.4	1404
32	Protein dislocation from the ER requires polyubiquitination and the AAA-ATPase Cdc48. <i>Nature Cell Biology</i> , 2002 , 4, 134-9	23.4	447
31	A LOV2 domain-based optogenetic tool to control protein degradation and cellular function. <i>Chemistry and Biology</i> , 2013 , 20, 619-26		181
30	Use of modular substrates demonstrates mechanistic diversity and reveals differences in chaperone requirement of ERAD. <i>Journal of Biological Chemistry</i> , 2003 , 278, 35903-13	5.4	155
29	System of centromeric, episomal, and integrative vectors based on drug resistance markers for Saccharomyces cerevisiae. <i>BioTechniques</i> , 2006 , 40, 73-8	2.5	136
28	ER-golgi traffic is a prerequisite for efficient ER degradation. <i>Molecular Biology of the Cell</i> , 2002 , 13, 180	0 6. 48	97
27	Efficient protein depletion by genetically controlled deprotection of a dormant N-degron. <i>Molecular Systems Biology</i> , 2009 , 5, 267	12.2	78
26	Spore number control and breeding in Saccharomyces cerevisiae: a key role for a self-organizing system. <i>Journal of Cell Biology</i> , 2005 , 171, 627-40	7.3	60
25	Targeted protein depletion in Saccharomyces cerevisiae by activation of a bidirectional degron. <i>BMC Systems Biology</i> , 2010 , 4, 176	3.5	46
24	Photo-sensitive degron variants for tuning protein stability by light. <i>BMC Systems Biology</i> , 2014 , 8, 128	3.5	44
23	The deca-GX3 proteins Yae1-Lto1 function as adaptors recruiting the ABC protein Rli1 for iron-sulfur cluster insertion. <i>ELife</i> , 2015 , 4, e08231	8.9	42
22	Dynamic organization of the actin cytoskeleton during meiosis and spore formation in budding yeast. <i>Traffic</i> , 2006 , 7, 1628-42	5.7	33
21	Cytokinesis in yeast meiosis depends on the regulated removal of Ssp1p from the prospore membrane. <i>EMBO Journal</i> , 2007 , 26, 1843-52	13	25
20	TIPI: TEV protease-mediated induction of protein instability. <i>Methods in Molecular Biology</i> , 2012 , 832, 611-26	1.4	23
19	A tobacco etch virus protease with increased substrate tolerance at the P1aposition. <i>PLoS ONE</i> , 2013 , 8, e67915	3.7	23
18	Nud1p, the yeast homolog of Centriolin, regulates spindle pole body inheritance in meiosis. <i>EMBO Journal</i> , 2006 , 25, 3856-68	13	23
17	Acetate regulation of spore formation is under the control of the Ras/cyclic AMP/protein kinase A pathway and carbon dioxide in Saccharomyces cerevisiae. <i>Eukaryotic Cell</i> , 2012 , 11, 1021-32		18

LIST OF PUBLICATIONS

16	Optogenetic Downregulation of Protein Levels with an Ultrasensitive Switch. <i>ACS Synthetic Biology</i> , 2019 , 8, 1026-1036	5.7	15
15	The Mitotic Exit Network Regulates Spindle Pole Body Selection During Sporulation of. <i>Genetics</i> , 2017 , 206, 919-937	4	13
14	Synthetic Control of Protein Degradation during Cell Proliferation and Developmental Processes. <i>ACS Omega</i> , 2019 , 4, 2766-2778	3.9	13
13	An Optogenetic Tool for Induced Protein Stabilization Based on the Phaeodactylum tricornutum Aureochrome 1a Light-Oxygen-Voltage Domain. <i>Journal of Molecular Biology</i> , 2020 , 432, 1880-1900	6.5	11
12	Controlling Protein Activity and Degradation Using Blue Light. <i>Methods in Molecular Biology</i> , 2016 , 1408, 67-78	1.4	10
11	Proteasome Activity Is Influenced by the HECT_2 Protein Ipa1 in Budding Yeast. <i>Genetics</i> , 2018 , 209, 157-171	4	9
10	Strategies to investigate protein turnover with fluorescent protein reporters in eukaryotic organisms. <i>AIMS Biophysics</i> , 2020 , 7, 90-118	0.8	7
9	Development of a Synthetic Switch to Control Protein Stability in Eukaryotic Cells with Light. <i>Methods in Molecular Biology</i> , 2017 , 1596, 241-255	1.4	5
8	Degradation of integral membrane proteins modified with the photosensitive degron module requires the cytosolic endoplasmic reticulum-associated degradation pathway. <i>Molecular Biology of the Cell</i> , 2019 , 30, 2558-2570	3.5	4
7	Regulation of exocytotic events by centrosome-analogous structures. <i>Topics in Current Genetics</i> , 2004 , 193-207		4
6	Biophotography: concepts, applications and perspectives. <i>Applied Microbiology and Biotechnology</i> , 2016 , 100, 3415-20	5.7	3
5	An Optogenetic Toolbox for Synergistic Regulation of Protein Abundance. <i>ACS Synthetic Biology</i> , 2021 ,	5.7	1
4	Development of an Optogenetic Tool to Regulate Protein Stability In Vivo118-131		
3	Lichtsignale f⊞die Hefe. <i>BioSpektrum</i> , 2019 , 25, 628-630	0.1	
2	Synthetische Biologie l\(\text{St} Mikroorganismen \text{\text{Behen}}\(\text{\text{BioSpektrum}}, \text{ 2015}, 21, 380-381	0.1	
1	Light-induced fermenter production of derivatives of the sweet protein monellin is maximized in prestationary Saccharomyces cerevisiae cultures <i>Biotechnology Journal</i> , 2022 , e2100676	5.6	_