

Hsueh-Chi Yen

List of Publications by Citations

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

14
papers

1,118
citations

11
h-index

14
g-index

14
ext. papers

1,285
ext. citations

20.4
avg, IF

3.95
L-index

#	Paper	IF	Citations
14	Global identification of modular cullin-RING ligase substrates. <i>Cell</i> , 2011 , 147, 459-74	56.2	321
13	Global protein stability profiling in mammalian cells. <i>Science</i> , 2008 , 322, 918-23	33.3	318
12	Identification of SCF ubiquitin ligase substrates by global protein stability profiling. <i>Science</i> , 2008 , 322, 923-9	33.3	147
11	Schizosaccharomyces pombe Int6 and Ras homologs regulate cell division and mitotic fidelity via the proteasome. <i>Cell</i> , 2003 , 112, 207-17	56.2	74
10	C-Terminal End-Directed Protein Elimination by CRL2 Ubiquitin Ligases. <i>Molecular Cell</i> , 2018 , 70, 602-613.e3	17.3	59
9	Yin6, a fission yeast Int6 homolog, complexes with Moe1 and plays a role in chromosome segregation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2000 , 97, 14370-5	11.5	44
8	SELENOPROTEINS. CRL2 aids elimination of truncated selenoproteins produced by failed UGA/Sec decoding. <i>Science</i> , 2015 , 349, 91-5	33.3	42
7	Rpn5 is a conserved proteasome subunit and required for proper proteasome localization and assembly. <i>Journal of Biological Chemistry</i> , 2003 , 278, 30669-76	5.4	32
6	Recognition of the Diglycine C-End Degron by CRL2 Ubiquitin Ligase. <i>Molecular Cell</i> , 2018 , 72, 813-822.e4	17.6	30
5	Transcriptional activation of C/EBPbeta gene by c-Jun and ATF2. <i>DNA and Cell Biology</i> , 2002 , 21, 551-60	3.6	18
4	Isolation of the Schizosaccharomyces pombe proteasome subunit Rpn7 and a structure-function study of the proteasome-COP9-initiation factor domain. <i>Journal of Biological Chemistry</i> , 2007 , 282, 32414-23	5.4	16
3	INT6: A Link Between the Proteasome and Tumorigenesis. <i>Cell Cycle</i> , 2003 , 2, 80-82	4.7	11
2	The C-degron pathway eliminates mislocalized proteins and products of deubiquitinating enzymes. <i>EMBO Journal</i> , 2021 , 40, e105846	13	4
1	A quantitative model for the rate-limiting process of UGA alternative assignments to stop and selenocysteine codons. <i>PLoS Computational Biology</i> , 2017 , 13, e1005367	5	2