

Tasha L Johnson

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/105977/publications.pdf>

Version: 2024-02-01

11
papers

889
citations

840776

11
h-index

1281871

11
g-index

12
all docs

12
docs citations

12
times ranked

1960
citing authors

#	ARTICLE	IF	CITATIONS
1	Global post-translational modification profiling of HIV-1-infected cells reveals mechanisms of host cellular pathway remodeling. <i>Cell Reports</i> , 2022, 39, 110690.	6.4	12
2	Multiple Routes to Oncogenesis Are Promoted by the Human Papillomavirusâ€™Host Protein Network. <i>Cancer Discovery</i> , 2018, 8, 1474-1489.	9.4	67
3	CRL4 ^{AMBRA1} targets Elongin C for ubiquitination and degradation to modulate CRL5 signaling. <i>EMBO Journal</i> , 2018, 37, .	7.8	13
4	An Mtb-Human Protein-Protein Interaction Map Identifies a Switch between Host Antiviral and Antibacterial Responses. <i>Molecular Cell</i> , 2018, 71, 637-648.e5.	9.7	100
5	Centriolar satellites assemble centrosomal microcephaly proteins to recruit CDK2 and promote centriole duplication. <i>ELife</i> , 2015, 4, .	6.0	118
6	Acetylome Profiling Reveals Overlap in the Regulation of Diverse Processes by Sirtuins, Gcn5, and Esa1. <i>Molecular and Cellular Proteomics</i> , 2015, 14, 162-176.	3.8	59
7	Global Mapping of Herpesvirus-Host Protein Complexes Reveals a Transcription Strategy for Late Genes. <i>Molecular Cell</i> , 2015, 57, 349-360.	9.7	165
8	A Combined Proteomics/Genomics Approach Links Hepatitis C Virus Infection with Nonsense-Mediated mRNA Decay. <i>Molecular Cell</i> , 2015, 57, 329-340.	9.7	124
9	Global Mapping of the Inc-Human Interactome Reveals that Retromer Restricts Chlamydia Infection. <i>Cell Host and Microbe</i> , 2015, 18, 109-121.	11.0	174
10	Lineage-Specific Viral Hijacking of Non-canonical E3ÂUbiquitin Ligase Cofactors in the Evolution of Vif Anti-APOBEC3 Activity. <i>Cell Reports</i> , 2015, 11, 1236-1250.	6.4	42
11	Prediction of Functionally Important Phospho-Regulatory Events in <i>Xenopus laevis</i> Oocytes. <i>PLoS Computational Biology</i> , 2015, 11, e1004362.	3.2	14