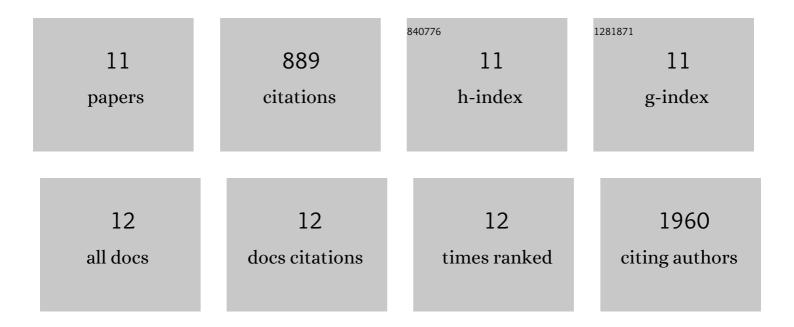
Tasha L Johnson

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/105977/publications.pdf Version: 2024-02-01



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