

# Gwendolyn M Jang

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1344349/publications.pdf>

Version: 2024-02-01

17  
papers

6,884  
citations

567281  
15  
h-index

888059  
17  
g-index

24  
all docs

24  
docs citations

24  
times ranked

13727  
citing authors

#	ARTICLE	IF	CITATIONS
1	A SARS-CoV-2 protein interaction map reveals targets for drug repurposing. <i>Nature</i> , 2020, 583, 459-468.	27.8	3,542
2	The Global Phosphorylation Landscape of SARS-CoV-2 Infection. <i>Cell</i> , 2020, 182, 685-712.e19.	28.9	825
3	Global landscape of HIV-human protein complexes. <i>Nature</i> , 2012, 481, 365-370.	27.8	651
4	Comparative host-coronavirus protein interaction networks reveal pan-viral disease mechanisms. <i>Science</i> , 2020, 370, .	12.6	508
5	Comparative Flavivirus-Host Protein Interaction Mapping Reveals Mechanisms of Dengue and Zika Virus Pathogenesis. <i>Cell</i> , 2018, 175, 1931-1945.e18.	28.9	252
6	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
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	Protein Interaction Mapping Identifies RBBP6 as a Negative Regulator of Ebola Virus Replication. <i>Cell</i> , 2018, 175, 1917-1930.e13.	28.9	108
9	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
10	Multiple Routes to Oncogenesis Are Promoted by the Human Papillomavirus-Host Protein Network. <i>Cancer Discovery</i> , 2018, 8, 1474-1489.	9.4	67
11	Identification of antiviral roles for the exon-junction complex and nonsense-mediated decay in flaviviral infection. <i>Nature Microbiology</i> , 2019, 4, 985-995.	13.3	52
12	Enterovirus pathogenesis requires the host methyltransferase SETD3. <i>Nature Microbiology</i> , 2019, 4, 2523-2537.	13.3	51
13	PJA2 ubiquitinates the HIV-1 Tat protein with atypical chain linkages to activate viral transcription. <i>Scientific Reports</i> , 2017, 7, 45394.	3.3	30
14	The HIV-1 Tat protein recruits a ubiquitin ligase to reorganize the 7SK snRNP for transcriptional activation. <i>ELife</i> , 2018, 7, .	6.0	29
15	PEAK3/C19orf35 pseudokinase, a new NFK3 kinase family member, inhibits Crkl through dimerization. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 15495-15504.	7.1	19
16	A Quantitative Genetic Interaction Map of HIV Infection. <i>Molecular Cell</i> , 2020, 78, 197-209.e7.	9.7	17
17	CRL4 <sup>AMBRA1</sup> targets Elongin C for ubiquitination and degradation to modulate CRL5 signaling. <i>EMBO Journal</i> , 2018, 37, .	7.8	13