

# David P Nusinow

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6137481/publications.pdf>

Version: 2024-02-01

14  
papers

3,563  
citations

840776

11  
h-index

1058476

14  
g-index

16  
all docs

16  
docs citations

16  
times ranked

6401  
citing authors

#	ARTICLE	IF	CITATIONS
1	Dual proteome-scale networks reveal cell-specific remodeling of the human interactome. <i>Cell</i> , 2021, 184, 3022-3040.e28.	28.9	455
2	Reimagining high-throughput profiling of reactive cysteines for cell-based screening of large electrophile libraries. <i>Nature Biotechnology</i> , 2021, 39, 630-641.	17.5	142
3	Quantitative Proteomics of the Cancer Cell Line Encyclopedia. <i>Cell</i> , 2020, 180, 387-402.e16.	28.9	596
4	Investigation of Proteomic and Phosphoproteomic Responses to Signaling Network Perturbations Reveals Functional Pathway Organizations in Yeast. <i>Cell Reports</i> , 2019, 29, 2092-2104.e4.	6.4	41
5	Reply to "Selective effects of heterozygous protein-truncating variants"™. <i>Nature Genetics</i> , 2019, 51, 3-4.	21.4	6
6	Active Instrument Engagement Combined with a Real-Time Database Search for Improved Performance of Sample Multiplexing Workflows. <i>Journal of Proteome Research</i> , 2019, 18, 1299-1306.	3.7	109
7	Estimating the selective effects of heterozygous protein-truncating variants from human exome data. <i>Nature Genetics</i> , 2017, 49, 806-810.	21.4	157
8	A mass-tolerant database search identifies a large proportion of unassigned spectra in shotgun proteomics as modified peptides. <i>Nature Biotechnology</i> , 2015, 33, 743-749.	17.5	371
9	MultiNotch MS3 Enables Accurate, Sensitive, and Multiplexed Detection of Differential Expression across Cancer Cell Line Proteomes. <i>Analytical Chemistry</i> , 2014, 86, 7150-7158.	6.5	1,130
10	Quantitative Temporal Viromics: An Approach to Investigate Host-Pathogen Interaction. <i>Cell</i> , 2014, 157, 1460-1472.	28.9	409
11	Essential roles for <i>stat92E</i> in expanding and patterning the proximodistal axis of the <i>Drosophila</i> wing imaginal disc. <i>Developmental Biology</i> , 2013, 378, 38-50.	2.0	15
12	Network-based inference from complex proteomic mixtures using SNIPE. <i>Bioinformatics</i> , 2012, 28, 3115-3122.	4.1	9
13	Networked-based Characterization of Extracellular Matrix Proteins from Adult Mouse Pulmonary and Aortic Valves. <i>Journal of Proteome Research</i> , 2011, 10, 812-823.	3.7	36
14	Reciprocal roles for <i>bowl</i> and <i>lines</i> in specifying the peripodial epithelium and the disc proper of the <i>Drosophila</i> wing primordium. <i>Development (Cambridge)</i> , 2008, 135, 3031-3041.	2.5	17