

# Daniel Jaschob

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4973398/publications.pdf>

Version: 2024-02-01

9  
papers

198  
citations

1683354

5  
h-index

1473754

9  
g-index

10  
all docs

10  
docs citations

10  
times ranked

427  
citing authors

#	ARTICLE	IF	CITATIONS
1	The molecular architecture of the Dam1 kinetochore complex is defined by cross-linking based structural modelling. <i>Nature Communications</i> , 2015, 6, 8673.	5.8	51
2	ProXL (Protein Cross-Linking Database): A Platform for Analysis, Visualization, and Sharing of Protein Cross-Linking Mass Spectrometry Data. <i>Journal of Proteome Research</i> , 2016, 15, 2863-2870.	1.8	51
3	MetaGOmics: A Web-Based Tool for Peptide-Centric Functional and Taxonomic Analysis of Metaproteomics Data. <i>Proteomes</i> , 2018, 6, 2.	1.7	43
4	Determining Protein Complex Structures Based on a Bayesian Model of in Vivo Förster Resonance Energy Transfer (FRET) Data. <i>Molecular and Cellular Proteomics</i> , 2014, 13, 2812-2823.	2.5	29
5	Discovery and Visualization of Uncharacterized Drug-Protein Adducts Using Mass Spectrometry. <i>Analytical Chemistry</i> , 2022, 94, 3501-3509.	3.2	7
6	Visualization and Dissemination of Multidimensional Proteomics Data Comparing Protein Abundance During <i>Caenorhabditis elegans</i> Development. <i>Journal of the American Society for Mass Spectrometry</i> , 2015, 26, 1827-1836.	1.2	5
7	Mason: a JavaScript web site widget for visualizing and comparing annotated features in nucleotide or protein sequences. <i>BMC Research Notes</i> , 2015, 8, 70.	0.6	4
8	Proxl (Protein Cross-Linking Database): A Public Server, QC Tools, and Other Major Updates. <i>Journal of Proteome Research</i> , 2019, 18, 759-764.	1.8	4
9	JobCenter: an open source, cross-platform, and distributed job queue management system optimized for scalability and versatility. <i>Source Code for Biology and Medicine</i> , 2012, 7, 8.	1.7	3