

# Daniel S Ziemianowicz

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5345878/publications.pdf>

Version: 2024-02-01

12  
papers

298  
citations

1163117

8  
h-index

1199594

12  
g-index

12  
all docs

12  
docs citations

12  
times ranked

489  
citing authors

#	ARTICLE	IF	CITATIONS
1	First Community-Wide, Comparative Cross-Linking Mass Spectrometry Study. <i>Analytical Chemistry</i> , 2019, 91, 6953-6961.	6.5	100
2	Structure of the mycobacterial ESX-5 type VII secretion system pore complex. <i>Science Advances</i> , 2021, 7, .	10.3	45
3	Amino Acid Insertion Frequencies Arising from Photoproducts Generated Using Aliphatic Diazirines. <i>Journal of the American Society for Mass Spectrometry</i> , 2017, 28, 2011-2021.	2.8	42
4	Photo-Cross-Linking Mass Spectrometry and Integrative Modeling Enables Rapid Screening of Antigen Interactions Involving Bacterial Transferrin Receptors. <i>Journal of Proteome Research</i> , 2019, 18, 934-946.	3.7	20
5	Simultaneous Proteoform Analysis of Histones H3 and H4 with a Simplified Middle-Down Proteomics Method. <i>Analytical Chemistry</i> , 2018, 90, 3083-3090.	6.5	17
6	The substrate specificity of the human TRAPPII complexâ€™s Rab-guanine nucleotide exchange factor activity. <i>Communications Biology</i> , 2020, 3, 735.	4.4	16
7	Quantitative Analysis of Protein Covalent Labeling Mass Spectrometry Data in the Mass Spec Studio. <i>Analytical Chemistry</i> , 2019, 91, 8492-8499.	6.5	15
8	Aqueous production of oxygen atoms from hydroxyl radicals. <i>Physical Chemistry Chemical Physics</i> , 2014, 16, 26094-26102.	2.8	13
9	Mechanism of O( <sup>3</sup> P) Formation from a Hydroxyl Radical Pair in Aqueous Solution. <i>Journal of Chemical Theory and Computation</i> , 2015, 11, 4740-4748.	5.3	9
10	Correlation between Labeling Yield and Surface Accessibility in Covalent Labeling Mass Spectrometry. <i>Journal of the American Society for Mass Spectrometry</i> , 2020, 31, 207-216.	2.8	8
11	Harmonizing structural mass spectrometry analyses in the mass spec studio. <i>Journal of Proteomics</i> , 2020, 225, 103844.	2.4	7
12	IMProv: A Resource for Cross-link-Driven Structure Modeling that Accommodates Protein Dynamics. <i>Molecular and Cellular Proteomics</i> , 2021, 20, 100139.	3.8	6