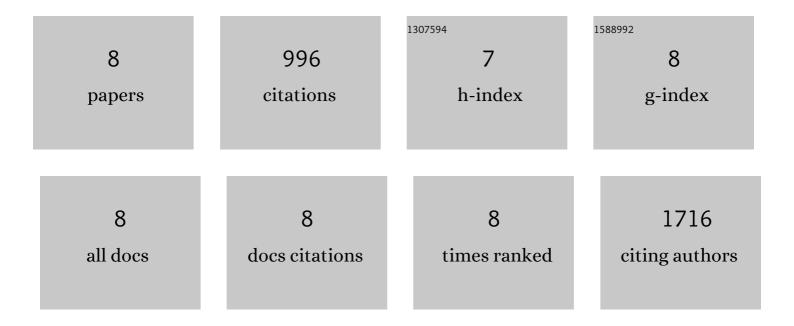
Daniel P Zolg

List of Publications by Year in descending order

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DANIEL P.ZOLC

#	Article	IF	CITATIONS
1	Mass-spectrometry-based draft of the Arabidopsis proteome. Nature, 2020, 579, 409-414.	27.8	328
2	Robust, reproducible and quantitative analysis of thousands of proteomes by micro-flow LC–MS/MS. Nature Communications, 2020, 11, 157.	12.8	218
3	Building ProteomeTools based on a complete synthetic human proteome. Nature Methods, 2017, 14, 259-262.	19.0	182
4	Mining the Human Tissue Proteome for Protein Citrullination. Molecular and Cellular Proteomics, 2018, 17, 1378-1391.	3.8	93
5	Deep learning boosts sensitivity of mass spectrometry-based immunopeptidomics. Nature Communications, 2021, 12, 3346.	12.8	90
6	High pH Reversed-Phase Micro-Columns for Simple, Sensitive, and Efficient Fractionation of Proteome and (TMT labeled) Phosphoproteome Digests. Methods in Molecular Biology, 2017, 1550, 83-98.	0.9	43
7	INFERYS rescoring: Boosting peptide identifications and scoring confidence of database search results. Rapid Communications in Mass Spectrometry, 2021, , e9128.	1.5	34
8	Prosit-TMT: Deep Learning Boosts Identification of TMT-Labeled Peptides. Analytical Chemistry, 2022, 94, 7181-7190.	6.5	8