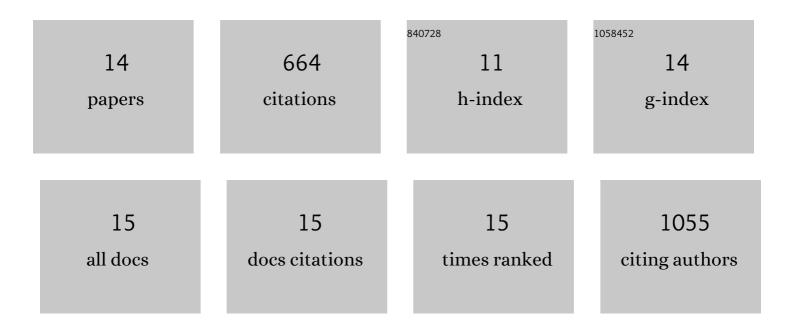
Daria Gawron

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	PROTEOFORMER: deep proteome coverage through ribosome profiling and MS integration. Nucleic Acids Research, 2015, 43, e29-e29.	14.5	132
2	N-terminal Proteomics and Ribosome Profiling Provide a Comprehensive View of the Alternative Translation Initiation Landscape in Mice and Men. Molecular and Cellular Proteomics, 2014, 13, 1245-1261.	3.8	123
3	Positional proteomics reveals differences in Nâ€ŧerminal proteoform stability. Molecular Systems Biology, 2016, 12, 858.	7.2	79
4	A proteogenomics approach integrating proteomics and ribosome profiling increases the efficiency of protein identification and enables the discovery of alternative translation start sites. Proteomics, 2014, 14, 2688-2698.	2.2	66
5	Site-specific methylation of 18S ribosomal RNA by SNORD42A is required for acute myeloid leukemia cell proliferation. Blood, 2020, 135, 2059-2070.	1.4	52
6	elF1 modulates the recognition of suboptimal translation initiation sites and steers gene expression via uORFs. Nucleic Acids Research, 2017, 45, 7997-8013.	14.5	51
7	An affinity-enhanced, broadly neutralizing heavy chain–only antibody protects against SARS-CoV-2 infection in animal models. Science Translational Medicine, 2021, 13, eabi7826.	12.4	41
8	The proteome under translational control. Proteomics, 2014, 14, 2647-2662.	2.2	38
9	Omics Assisted N-terminal Proteoform and Protein Expression Profiling On Methionine Aminopeptidase 1 (MetAP1) Deletion. Molecular and Cellular Proteomics, 2018, 17, 694-708.	3.8	21
10	Mass spectrometry and the cellular surfaceome. Mass Spectrometry Reviews, 2022, 41, 804-841.	5.4	19
11	Matrix metalloproteinase (MMP)-2 and MMP-9 Expression in Tumor Infiltrating CD3 Lymphocytes From Women With Endometrial Cancer. International Journal of Gynecological Cancer, 2012, 22, 1303-1309.	2.5	16
12	Bacterial riboproteogenomics: the era of N-terminal proteoform existence revealed. FEMS Microbiology Reviews, 2020, 44, 418-431.	8.6	12
13	Limited Evidence for Protein Products of Noncoding Transcripts in the HEK293T Cellular Cytosol. Molecular and Cellular Proteomics, 2022, 21, 100264.	3.8	11
14	Orthogonal proteomics methods to unravel the HOTAIR interactome. Scientific Reports, 2022, 12, 1513.	3.3	3