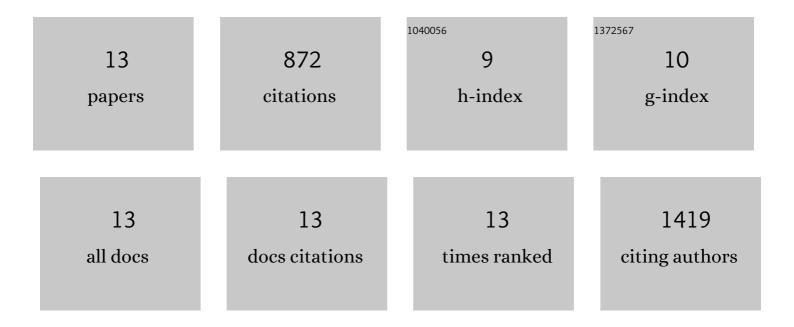
Ermir Qeli

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Generating and navigating proteome maps using mass spectrometry. Nature Reviews Molecular Cell Biology, 2010, 11, 789-801.	37.0	181
2	Deterministic protein inference for shotgun proteomics data provides new insights into Arabidopsis pollen development and function. Genome Research, 2009, 19, 1786-1800.	5.5	151
3	Identification and Functional Characterization of N-Terminally Acetylated Proteins in Drosophila melanogaster. PLoS Biology, 2009, 7, e1000236.	5.6	149
4	An integrated proteomics and transcriptomics reference data set provides new insights into the <i>Bradyrhizobium japonicum</i> bacteroid metabolism in soybean root nodules. Proteomics, 2010, 10, 1391-1400.	2.2	111
5	PeptideClassifier for protein inference and targeted quantitative proteomics. Nature Biotechnology, 2010, 28, 647-650.	17.5	92
6	A Mighty Small Heart: The Cardiac Proteome of Adult Drosophila melanogaster. PLoS ONE, 2011, 6, e18497.	2.5	81
7	Improved prediction of peptide detectability for targeted proteomics using a rank-based algorithm and organism-specific data. Journal of Proteomics, 2014, 108, 269-283.	2.4	43
8	Protein and gene model inference based on statistical modeling in k-partite graphs. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 12101-12106.	7.1	39
9	Visualizing regulatory interactions in metabolic networks. BMC Biology, 2007, 5, 46.	3.8	14
10	From Enzyme Kinetics to Metabolic Network Modeling – Visualization Tool for Enhanced Kinetic Analysis of Biochemical Network Models. Engineering in Life Sciences, 2006, 6, 155-162.	3.6	7
11	The time-dependent reorderable matrix method for visualizing evolving tabular data. , 2005, 5669, 199.		2
12	Customizable detection of changes for XML documents using XPath expressions. , 2006, , .		1
13	Filtering XML documents using XPath expressions and aspect-oriented programming. , 2006, , .		1