

# Attila Csordas

## List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

19  
papers

11,699  
citations

15  
h-index

21  
g-index

21  
ext. papers

15,504  
ext. citations

12.6  
avg, IF

5.84  
L-index

#	Paper	IF	Citations
19	The PRIDE database and related tools and resources in 2019: improving support for quantification data. <i>Nucleic Acids Research</i> , <b>2019</b> , 47, D442-D450	20.1	3856
18	The ProteomeXchange consortium in 2017: supporting the cultural change in proteomics public data deposition. <i>Nucleic Acids Research</i> , <b>2017</b> , 45, D1100-D1106	20.1	568
17	2016 update of the PRIDE database and its related tools. <i>Nucleic Acids Research</i> , <b>2016</b> , 44, D447-56	20.1	2440
16	Exploring the potential of public proteomics data. <i>Proteomics</i> , <b>2016</b> , 16, 214-25	4.8	54
15	PRIDE Inspector Toolsuite: Moving Toward a Universal Visualization Tool for Proteomics Data Standard Formats and Quality Assessment of ProteomeXchange Datasets. <i>Molecular and Cellular Proteomics</i> , <b>2016</b> , 15, 305-17	7.6	91
14	ProteomeXchange provides globally coordinated proteomics data submission and dissemination. <i>Nature Biotechnology</i> , <b>2014</b> , 32, 223-6	44.5	2053
13	How to submit MS proteomics data to ProteomeXchange via the PRIDE database. <i>Proteomics</i> , <b>2014</b> , 14, 2233-41	4.8	41
12	The PRoteomics IDentifications (PRIDE) database and associated tools: status in 2013. <i>Nucleic Acids Research</i> , <b>2013</b> , 41, D1063-9	20.1	1587
11	From Peptidome to PRIDE: public proteomics data migration at a large scale. <i>Proteomics</i> , <b>2013</b> , 13, 1692-58	4.8	11
10	Hydra: a scalable proteomic search engine which utilizes the Hadoop distributed computing framework. <i>BMC Bioinformatics</i> , <b>2012</b> , 13, 324	3.6	43
9	PRIDE Inspector: a tool to visualize and validate MS proteomics data. <i>Nature Biotechnology</i> , <b>2012</b> , 30, 135-7	44.5	103
8	The PRoteomics IDentification (PRIDE) Converter 2 framework: an improved suite of tools to facilitate data submission to the PRIDE database and the ProteomeXchange consortium. <i>Molecular and Cellular Proteomics</i> , <b>2012</b> , 11, 1682-9	7.6	96
7	PRIDE: quality control in a proteomics data repository. <i>Database: the Journal of Biological Databases and Curation</i> , <b>2012</b> , 2012, bas004	5	30
6	MitoWheel, visualizing the human mitochondrial genome. <i>Nature Precedings</i> , <b>2009</b> ,		3
5	Human heart mitochondria do not produce physiologically relevant quantities of nitric oxide. <i>Life Sciences</i> , <b>2007</b> , 80, 633-7	6.8	13
4	Mitochondrial transfer between eukaryotic animal cells and its physiologic role. <i>Rejuvenation Research</i> , <b>2006</b> , 9, 450-4	2.6	12
3	Mitochondrial NO and reactive nitrogen species production: does mtNOS exist?. <i>Nitric Oxide - Biology and Chemistry</i> , <b>2006</b> , 14, 162-8	5	93

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|---|---|-----|----|
| 2 | The novel red-fluorescent probe DAR-4M measures reactive nitrogen species rather than NO.<br><i>Journal of Pharmacological and Toxicological Methods</i> , <b>2005</b> , 52, 335-40 | 1.7 | 21 |
| 1 | MitoWheel, visualizing the human mitochondrial genome   |     | 1  |