Tobias Ternent

List of Publications by Year in Descending Order

12 papers	7,856 citations	13 h-index	14 g-index
14 ext. papers	11,200 ext. citations	13.4 avg, IF	5.65 L-index

#	Paper	IF	Citations
12	Quantitative Proteomics Data in the Public Domain: Challenges and Opportunities. *Methods in Molecular Biology*, **2019**, 1977, 217-235	1.4	4
11	The PRIDE database and related tools and resources in 2019: improving support for quantification data. *Nucleic Acids Research*, **2019**, 47, D442-D450	20.1	3856
10	The proBAM and proBed standard formats: enabling a seamless integration of genomics and proteomics data. *Genome Biology*, **2018**, 19, 12	18.3	16
9	Discovering and linking public omics data sets using the Omics Discovery Index. *Nature Biotechnology*, **2017**, 35, 406-409	44.5	105
8	The ProteomeXchange consortium in 2017: supporting the cultural change in proteomics public data deposition. *Nucleic Acids Research*, **2017**, 45, D1100-D1106	20.1	568
7	OLS Client and OLS Dialog: Open Source Tools to Annotate Public Omics Datasets. *Proteomics*, **2017**, 17, 1700244	4.8	13
6	The mzIdentML Data Standard Version 1.2, Supporting Advances in Proteome Informatics. *Molecular and Cellular Proteomics*, **2017**, 16, 1275-1285	7.6	37
5	2016 update of the PRIDE database and its related tools. *Nucleic Acids Research*, **2016**, 44, D447-56	20.1	2440
4	PRIDE Inspector Toolsuite: Moving Toward a Universal Visualization Tool for Proteomics Data Standard Formats and Quality Assessment of ProteomeXchange Datasets. *Molecular and Cellular Proteomics*, **2016**, 15, 305-17	7.6	91
3	ms-data-core-api: an open-source, metadata-oriented library for computational proteomics. *Bioinformatics*, **2015**, 31, 2903-5	7.2	26
2	Introducing the PRIDE Archive RESTful web services. *Nucleic Acids Research*, **2015**, 43, W599-604	20.1	15
1	How to submit MS proteomics data to ProteomeXchange via the PRIDE database. *Proteomics*, **2014**, 14, 2233-41	4.8	41