

# Chris F Taylor

## List of Publications by Year in descending order

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Version: 2024-02-01

24  
papers

3,524  
citations

331259

21  
h-index

525886

27  
g-index

28  
all docs

28  
docs citations

28  
times ranked

5943  
citing authors

#	ARTICLE	IF	CITATIONS
1	The Ontology for Biomedical Investigations. PLoS ONE, 2016, 11, e0154556.	1.1	217
2	Data Standards for Omics Data: The Basis of Data Sharing and Reuse. Methods in Molecular Biology, 2011, 719, 31-69.	0.4	73
3	Recombinant protein quality evaluation: proposal for a minimal information standard. Standards in Genomic Sciences, 2011, 5, 195-197.	1.5	8
4	Guidelines for reporting the use of capillary electrophoresis in proteomics. Nature Biotechnology, 2010, 28, 654-655.	9.4	24
5	Guidelines for reporting the use of gel image informatics in proteomics. Nature Biotechnology, 2010, 28, 655-656.	9.4	22
6	Debunking minimum information myths: one hat need not fit all. New Biotechnology, 2009, 25, 171-172.	2.4	5
7	'Omics Data Sharing. Science, 2009, 326, 234-236.	6.0	136
8	Standards for Functional Genomics. , 2009, , 293-329.		0
9	Towards interoperable reporting standards for omics data: hopes and hurdles. Summit on Translational Bioinformatics, 2009, 2009, 112-5.	0.7	1
10	Promoting coherent minimum reporting guidelines for biological and biomedical investigations: the MIBBI project. Nature Biotechnology, 2008, 26, 889-896.	9.4	506
11	Guidelines for reporting the use of mass spectrometry in proteomics. Nature Biotechnology, 2008, 26, 860-861.	9.4	82
12	Guidelines for reporting the use of mass spectrometry informatics in proteomics. Nature Biotechnology, 2008, 26, 862-862.	9.4	62
13	Guidelines for reporting the use of gel electrophoresis in proteomics. Nature Biotechnology, 2008, 26, 863-864.	9.4	61
14	The minimum information about a proteomics experiment (MIAPE). Nature Biotechnology, 2007, 25, 887-893.	9.4	694
15	The Functional Genomics Experiment model (FuGE): an extensible framework for standards in functional genomics. Nature Biotechnology, 2007, 25, 1127-1133.	9.4	96
16	Standards for reporting bioscience data: a forward look. Drug Discovery Today, 2007, 12, 527-533.	3.2	17
17	A roadmap for the establishment of standard data exchange structures for metabolomics. Metabolomics, 2007, 3, 243-248.	1.4	35
18	Minimum Reporting Requirements for Proteomics: A MIAPE Primer. Proteomics, 2006, 6, 39-44.	1.3	52

#	ARTICLE	IF	CITATIONS
19	PRIDE: a public repository of protein and peptide identifications for the proteomics community. Nucleic Acids Research, 2006, 34, D659-D663.	6.5	254
20	The Work of the Human Proteome Organisation's Proteomics Standards Initiative (HUPO PSI). OMICS A Journal of Integrative Biology, 2006, 10, 145-151.	1.0	64
21	Pedro: a configurable data entry tool for XML. Bioinformatics, 2004, 20, 2463-2465.	1.8	24
22	A common open representation of mass spectrometry data and its application to proteomics research. Nature Biotechnology, 2004, 22, 1459-1466.	9.4	724
23	Advances in the development of common interchange standards for proteomic data. Proteomics, 2004, 4, 2363-2365.	1.3	29
24	A systematic approach to modeling, capturing, and disseminating proteomics experimental data. Nature Biotechnology, 2003, 21, 247-254.	9.4	246