

# Mark Thompson

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6048282/publications.pdf>

Version: 2024-02-01

16  
papers

9,280  
citations

840776

11  
h-index

1058476

14  
g-index

20  
all docs

20  
docs citations

20  
times ranked

20563  
citing authors

#	ARTICLE	IF	CITATIONS
1	FAIR Principles: Interpretations and Implementation Considerations. <i>Data Intelligence</i> , 2020, 2, 10-29.	1.5	149
2	A Generic Workflow for the Data FAIRification Process. <i>Data Intelligence</i> , 2020, 2, 56-65.	1.5	59
3	Making FAIR Easy with FAIR Tools: From Creolization to Convergence. <i>Data Intelligence</i> , 2020, 2, 87-95.	1.5	21
4	The “œAœof FAIR œœ“ As Open as Possible, as Closed as Necessary. <i>Data Intelligence</i> , 2020, 2, 47-55.	1.5	29
5	BioHackathon 2015: Semantics of data for life sciences and reproducible research. <i>F1000Research</i> , 2020, 9, 136.	1.6	5
6	The RD-Connect Registry & Biobank Finder: a tool for sharing aggregated data and metadata among rare disease researchers. <i>European Journal of Human Genetics</i> , 2018, 26, 631-643.	2.8	33
7	Consent Codes: Upholding Standard Data Use Conditions. <i>PLoS Genetics</i> , 2016, 12, e1005772.	3.5	65
8	The Implicitome: A Resource for Rationalizing Gene-Disease Associations. <i>PLoS ONE</i> , 2016, 11, e0149621.	2.5	22
9	The FAIR Guiding Principles for scientific data management and stewardship. <i>Scientific Data</i> , 2016, 3, 160018.	5.3	8,670
10	Nanopublications for exposing experimental data in the life-sciences: a Huntington's Disease case study. <i>Journal of Biomedical Semantics</i> , 2015, 6, 5.	1.6	12
11	Multidisciplinary Collaboration to Facilitate Hypotheses Generation in Huntington's Disease. , 2015, , .		3
12	From Peer-Reviewed to Peer-Reproduced in Scholarly Publishing: The Complementary Roles of Data Models and Workflows in Bioinformatics. <i>PLoS ONE</i> , 2015, 10, e0127612.	2.5	27
13	Structuring research methods and data with the research object model: genomics workflows as a case study. <i>Journal of Biomedical Semantics</i> , 2014, 5, 41.	1.6	26
14	Preserving sequence annotations across reference sequences. <i>Journal of Biomedical Semantics</i> , 2014, 5, S6.	1.6	3
15	Interoperability and FAIRness through a novel combination of Web technologies. <i>PeerJ Computer Science</i> , 0, 3, e110.	4.5	58
16	BioHackathon series in 2013 and 2014: improvements of semantic interoperability in life science data and services. <i>F1000Research</i> , 0, 8, 1677.	1.6	0