

# Marta Iannuccelli

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3532386/publications.pdf>

Version: 2024-02-01

20  
papers

3,477  
citations

759233

12  
h-index

752698

20  
g-index

22  
all docs

22  
docs citations

22  
times ranked

6172  
citing authors

#	ARTICLE	IF	CITATIONS
1	The IntAct database: efficient access to fine-grained molecular interaction data. <i>Nucleic Acids Research</i> , 2022, 50, D648-D653.	14.5	89
2	The Intricacy of the Viral-Human Protein Interaction Networks: Resources, Data, and Analyses. <i>Frontiers in Microbiology</i> , 2022, 13, 849781.	3.5	2
3	A Resource to Infer Molecular Paths Linking Cancer Mutations to Perturbation of Cell Metabolism. <i>Frontiers in Molecular Biosciences</i> , 2022, 9, .	3.5	3
4	Phosphomatics: interactive interrogation of substrateâ€“kinase networks in global phosphoproteomics datasets. <i>Bioinformatics</i> , 2021, 37, 1635-1636.	4.1	12
5	Integrating Patient-Specific Information into Logic Models of Complex Diseases: Application to Acute Myeloid Leukemia. <i>Journal of Personalized Medicine</i> , 2021, 11, 117.	2.5	3
6	A Resource for the Network Representation of Cell Perturbations Caused by SARS-CoV-2 Infection. <i>Genes</i> , 2021, 12, 450.	2.4	7
7	COVID19 Disease Map, a computational knowledge repository of virusâ€“host interaction mechanisms. <i>Molecular Systems Biology</i> , 2021, 17, e10387.	7.2	53
8	SIGNOR 2.0, the SIGnaling Network Open Resource 2.0: 2019 update. <i>Nucleic Acids Research</i> , 2020, 48, D504-D510.	14.5	160
9	CancerGeneNet: linking driver genes to cancer hallmarks. <i>Nucleic Acids Research</i> , 2020, 48, D416-D421.	14.5	29
10	Towards a unified open access dataset of molecular interactions. <i>Nature Communications</i> , 2020, 11, 6144.	12.8	49
11	Using the MINT Database to Search Protein Interactions. <i>Current Protocols in Bioinformatics</i> , 2020, 69, e93.	25.8	13
12	Myo-REG: A Portal for Signaling Interactions in Muscle Regeneration. <i>Frontiers in Physiology</i> , 2019, 10, 1216.	2.8	8
13	Selectivity of the CUBAN domain in the recognition of ubiquitin and NEDD8. <i>FEBS Journal</i> , 2019, 286, 653-677.	4.7	22
14	DISNOR: a disease network open resource. <i>Nucleic Acids Research</i> , 2018, 46, D527-D534.	14.5	42
15	SIGNOR: a database of causal relationships between biological entities. <i>Nucleic Acids Research</i> , 2016, 44, D548-D554.	14.5	243
16	The MIntAct projectâ€“IntAct as a common curation platform for 11 molecular interaction databases. <i>Nucleic Acids Research</i> , 2014, 42, D358-D363.	14.5	1,634
17	HuPho: the human phosphatase portal. <i>FEBS Journal</i> , 2013, 280, 379-387.	4.7	55
18	MINT, the molecular interaction database: 2012 update. <i>Nucleic Acids Research</i> , 2012, 40, D857-D861.	14.5	917

#	ARTICLE	IF	CITATIONS
19	The Protein-Protein Interaction tasks of BioCreative III: classification/ranking of articles and linking bio-ontology concepts to full text. BMC Bioinformatics, 2011, 12, S3.	2.6	121
20	Benchmarking of the 2010 BioCreative Challenge III text-mining competition by the BioGRID and MINT interaction databases. BMC Bioinformatics, 2011, 12, S8.	2.6	11