Marta Iannuccelli

List of Publications by Year
in descending order

20	3,477	759233	752698
papers	citations	12	20
		h-index	g-index

22	22	22	6172
all docs	docs citations	times ranked	citing authors

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

#	Article	IF	Citations
19	A Resource to Infer Molecular Paths Linking Cancer Mutations to Perturbation of Cell Metabolism. Frontiers in Molecular Biosciences, 2022, 9, .	3.5	3
20	The Intricacy of the Viral-Human Protein Interaction Networks: Resources, Data, and Analyses. Frontiers in Microbiology, 2022, 13, 849781.	3.5	2