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List of Publications by Year in descending order

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257101 233125 4,892 45 43 24 citations h-index g-index papers 50 50 50 7993 citing authors docs citations times ranked all docs

#	Article	IF	Citations
1	The IntAct database: efficient access to fine-grained molecular interaction data. Nucleic Acids Research, 2022, 50, D648-D653.	6.5	89
2	The gene regulation knowledge commons: the action area of GREEKC. Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms, 2022, 1865, 194768.	0.9	3
3	The RDâ€Connect Genomeâ€Phenome Analysis Platform: Accelerating diagnosis, research, and gene discovery for rare diseases. Human Mutation, 2022, , .	1.1	18
4	The Intricacy of the Viral-Human Protein Interaction Networks: Resources, Data, and Analyses. Frontiers in Microbiology, 2022, 13, 849781.	1.5	2
5	IMEx Databases: Displaying Molecular Interactions into a Single, Standards-Compliant Dataset. Methods in Molecular Biology, 2022, 2449, 27-42.	0.4	4
6	A Resource to Infer Molecular Paths Linking Cancer Mutations to Perturbation of Cell Metabolism. Frontiers in Molecular Biosciences, 2022, 9, .	1.6	3
7	Phosphomatics: interactive interrogation of substrate–kinase networks in global phosphoproteomics datasets. Bioinformatics, 2021, 37, 1635-1636.	1.8	12
8	The Minimum Information about a Molecular Interaction CAusal STatement (MI2CAST). Bioinformatics, 2021, 36, 5712-5718.	1.8	14
9	Integrating Patient-Specific Information into Logic Models of Complex Diseases: Application to Acute Myeloid Leukemia. Journal of Personalized Medicine, 2021, 11, 117.	1.1	3
10	A Resource for the Network Representation of Cell Perturbations Caused by SARS-CoV-2 Infection. Genes, 2021, 12, 450.	1.0	7
11	COVID19 Disease Map, a computational knowledge repository of virus–host interaction mechanisms. Molecular Systems Biology, 2021, 17, e10387.	3.2	53
12	SIGNOR 2.0, the SIGnaling Network Open Resource 2.0: 2019 update. Nucleic Acids Research, 2020, 48, D504-D510.	6.5	160
13	CancerGeneNet: linking driver genes to cancer hallmarks. Nucleic Acids Research, 2020, 48, D416-D421.	6.5	29
14	Towards a unified open access dataset of molecular interactions. Nature Communications, 2020, 11, 6144.	5.8	49
15	The ELIXIR Core Data Resources: fundamental infrastructure for the life sciences. Bioinformatics, 2020, 36, 2636-2642.	1.8	47
16	Using the MINT Database to Search Protein Interactions. Current Protocols in Bioinformatics, 2020, 69, e93.	25.8	13
17	The IMEx coronavirus interactome: an evolving map of <i> Coronaviridae < /i > $\hat{a} \in \text{``host molecular'}$ interactions. Database: the Journal of Biological Databases and Curation, 2020, 2020, .</i>	1.4	34
18	Myo-REC: A Portal for Signaling Interactions in Muscle Regeneration. Frontiers in Physiology, 2019, 10, 1216.	1.3	8

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19	CausalTAB: the PSI-MITAB 2.8 updated format for signalling data representation and dissemination. Bioinformatics, 2019, 35, 3779-3785.	1.8	32
20	DISNOR: a disease network open resource. Nucleic Acids Research, 2018, 46, D527-D534.	6.5	42
21	Late recurrent cutaneous breast angiosarcoma in an elderly woman: A case report. Oncology Letters, 2016, 11, 3247-3250.	0.8	1
22	The MIntAct Project and Molecular Interaction Databases. Methods in Molecular Biology, 2016, 1415, 55-69.	0.4	16
23	Tools and data services registry: a community effort to document bioinformatics resources. Nucleic Acids Research, 2016, 44, D38-D47.	6.5	113
24	SIGNOR: a database of causal relationships between biological entities. Nucleic Acids Research, 2016, 44, D548-D554.	6.5	243
25	The complex portal - an encyclopaedia of macromolecular complexes. Nucleic Acids Research, 2015, 43, D479-D484.	6.5	100
26	VirusMentha: a new resource for virus-host protein interactions. Nucleic Acids Research, 2015, 43, D588-D592.	6.5	141
27	The MIntAct projectâ€"IntAct as a common curation platform for 11 molecular interaction databases. Nucleic Acids Research, 2014, 42, D358-D363.	6.5	1,634
28	Clear Cell Sarcoma Of Metatarsus. Folia Medica, 2014, 56, 271-274.	0.2	4
29	An overview of the BioCreative 2012 Workshop Track III: interactive text mining task. Database: the Journal of Biological Databases and Curation, 2013, 2013, bas056-bas056.	1.4	68
30	MINT, the molecular interaction database: 2012 update. Nucleic Acids Research, 2012, 40, D857-D861.	6.5	917
31	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.	1.2	121
32	Benchmarking of the 2010 BioCreative Challenge III text-mining competition by the BioGRID and MINT interaction databases. BMC Bioinformatics, 2011, 12, S8.	1.2	11
33	Enriching the viral–host interactomes with interactions mediated by SH3 domains. Amino Acids, 2010, 38, 1541-1547.	1.2	9
34	The FEBS Letters/BioCreative II.5 experiment: making biological information accessible. Nature Biotechnology, 2010, 28, 897-899.	9.4	42
35	MINT, the molecular interaction database: 2009 update. Nucleic Acids Research, 2010, 38, D532-D539.	6.5	458
36	Recurated protein interaction datasets. Nature Methods, 2009, 6, 860-861.	9.0	58

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37	Linking entries in protein interaction database to structured text: The FEBS Letters experiment. FEBS Letters, 2008, 582, 1171-1177.	1.3	62
38	Protein interactions: integration leads to belief. Trends in Biochemical Sciences, 2008, 33, 241-242.	3.7	33
39	MINT and IntAct contribute to the Second BioCreative challenge: serving the text-mining community with high quality molecular interaction data. Genome Biology, 2008, 9, S5.	13.9	24
40	Searching the MINT Database for Protein Interaction Information. Current Protocols in Bioinformatics, 2008, 22, Unit 8.5.	25.8	10
41	Functional feature of a novel model of blood brain barrier: studies on permeation of test compounds. Journal of Controlled Release, 2001, 76, 139-147.	4.8	59
42	Neurons and ECM regulate occludin localization in brain endothelial cells. NeuroReport, 2000, 11, 1081-1084.	0.6	111
43	Synergistic effects of laminin and thyroid hormones on neuron polarity in culture. NeuroReport, 1999, 10, 1269-1272.	0.6	12