

Ian M Donaldson

List of Publications by Year in descending order

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Version: 2024-02-01

22
papers

6,059
citations

516710

16
h-index

713466

21
g-index

22
all docs

22
docs citations

22
times ranked

8017
citing authors

#	ARTICLE	IF	CITATIONS
1	Systematic identification of protein complexes in <i>Saccharomyces cerevisiae</i> by mass spectrometry. <i>Nature</i> , 2002, 415, 180-183.	27.8	3,445
2	BIND--The Biomolecular Interaction Network Database. <i>Nucleic Acids Research</i> , 2001, 29, 242-245.	14.5	842
3	iRefIndex: A consolidated protein interaction database with provenance. <i>BMC Bioinformatics</i> , 2008, 9, 405.	2.6	488
4	PSICQUIC and PSIScore: accessing and scoring molecular interactions. <i>Nature Methods</i> , 2011, 8, 528-529.	19.0	274
5	Inflammation driven by tumour-specific Th1 cells protects against B-cell cancer. <i>Nature Communications</i> , 2011, 2, 240.	12.8	251
6	PreBIND and Textomy--mining the biomedical literature for protein-protein interactions using a support vector machine. <i>BMC Bioinformatics</i> , 2003, 4, 11.	2.6	239
7	iRefWeb: interactive analysis of consolidated protein interaction data and their supporting evidence. <i>Database: the Journal of Biological Databases and Curation</i> , 2010, 2010, baq023-baq023.	3.0	194
8	iRefR: an R package to manipulate the iRefIndex consolidated protein interaction database. <i>BMC Bioinformatics</i> , 2011, 12, 455.	2.6	65
9	Literature curation of protein interactions: measuring agreement across major public databases. <i>Database: the Journal of Biological Databases and Curation</i> , 2010, 2010, baq026-baq026.	3.0	54
10	Interaction databases on the same page. <i>Nature Biotechnology</i> , 2011, 29, 391-393.	17.5	36
11	Zinc Stoichiometry of Yeast RNA Polymerase II and Characterization of Mutations in the Zinc-binding Domain of the Largest Subunit. <i>Journal of Biological Chemistry</i> , 2000, 275, 13780-13788.	3.4	29
12	Adaptive from Innate: Human IFN- γ +CD4+ T Cells Can Arise Directly from CXCL8-Producing Recent Thymic Emigrants in Babies and Adults. <i>Journal of Immunology</i> , 2017, 199, 1696-1705.	0.8	27
13	Effects of protein interaction data integration, representation and reliability on the use of network properties for drug target prediction. <i>BMC Bioinformatics</i> , 2012, 13, 294.	2.6	24
14	Navigating the Global Protein-Protein Interaction Landscape Using iRefWeb. <i>Methods in Molecular Biology</i> , 2014, 1091, 315-331.	0.9	19
15	Postmortem Genetic Testing for Cardiac Ion Channelopathies in Stillbirths. <i>Circulation Genomic and Precision Medicine</i> , 2018, 11, e001817.	3.6	18
16	Proteome analysis of microtubule-associated proteins and their interacting partners from mammalian brain. <i>Amino Acids</i> , 2011, 41, 363-385.	2.7	16
17	iRefScape. A Cytoscape plug-in for visualization and data mining of protein interaction data from iRefIndex. <i>BMC Bioinformatics</i> , 2011, 12, 388.	2.6	13
18	Automatic annotation of BIND molecular interactions from three-dimensional structures. <i>Biopolymers</i> , 2001, 61, 111-120.	2.4	11

#	ARTICLE	IF	CITATIONS
19	The Contribution of DNA Base Damage to Human Cancer Is Modulated by the Base Excision Repair Interaction Network. <i>Critical Reviews in Oncogenesis</i> , 2008, 14, 217-273.	0.4	7
20	A survey of protein interaction data and multigenic inherited disorders. <i>BMC Bioinformatics</i> , 2013, 14, 47.	2.6	4
21	Navigating the Global Protein-Protein Interaction Landscape Using iRefWeb. <i>Methods in Molecular Biology</i> , 2021, 2199, 191-207.	0.9	2
22	Protein Interaction Data Resources. , 2010, , 1375-1385.		1