Craig Knox

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/11930963/publications.pdf

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257450 580821 29,627 25 24 25 h-index citations g-index papers 25 25 25 40060 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	DrugBank 5.0: a major update to the DrugBank database for 2018. Nucleic Acids Research, 2018, 46, D1074-D1082.	14.5	5,428
2	DrugBank: a comprehensive resource for in silico drug discovery and exploration. Nucleic Acids Research, 2006, 34, D668-D672.	14.5	3,070
3	HMDB 4.0: the human metabolome database for 2018. Nucleic Acids Research, 2018, 46, D608-D617.	14.5	2,805
4	HMDB 3.0â€"The Human Metabolome Database in 2013. Nucleic Acids Research, 2012, 41, D801-D807.	14.5	2,564
5	HMDB: the Human Metabolome Database. Nucleic Acids Research, 2007, 35, D521-D526.	14.5	2,563
6	DrugBank: a knowledgebase for drugs, drug actions and drug targets. Nucleic Acids Research, 2008, 36, D901-D906.	14.5	2,336
7	DrugBank 4.0: shedding new light on drug metabolism. Nucleic Acids Research, 2014, 42, D1091-D1097.	14.5	1,884
8	HMDB: a knowledgebase for the human metabolome. Nucleic Acids Research, 2009, 37, D603-D610.	14.5	1,649
9	DrugBank 3.0: a comprehensive resource for 'Omics' research on drugs. Nucleic Acids Research, 2011, 39, D1035-D1041.	14.5	1,566
10	The Human Serum Metabolome. PLoS ONE, 2011, 6, e16957.	2.5	1,378
11	The Human Urine Metabolome. PLoS ONE, 2013, 8, e73076.	2.5	1,125
12	ClassyFire: automated chemical classification with a comprehensive, computable taxonomy. Journal of Cheminformatics, 2016, 8, 61.	6.1	779
13	Phenol-Explorer 3.0: a major update of the Phenol-Explorer database to incorporate data on the effects of food processing on polyphenol content. Database: the Journal of Biological Databases and Curation, 2013, 2013, bat070-bat070.	3.0	590
14	SMPDB 2.0: Big Improvements to the Small Molecule Pathway Database. Nucleic Acids Research, 2014, 42, D478-D484.	14.5	341
15	SMPDB: The Small Molecule Pathway Database. Nucleic Acids Research, 2010, 38, D480-D487.	14.5	290
16	T3DB: the toxic exposome database. Nucleic Acids Research, 2015, 43, D928-D934.	14.5	228
17	PolySearch: a web-based text mining system for extracting relationships between human diseases, genes, mutations, drugs and metabolites. Nucleic Acids Research, 2008, 36, W399-W405.	14.5	217
18	YMDB: the Yeast Metabolome Database. Nucleic Acids Research, 2012, 40, D815-D820.	14.5	162

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#	Article	IF	CITATION
19	Phenol-Explorer 2.0: a major update of the Phenol-Explorer database integrating data on polyphenol metabolism and pharmacokinetics in humans and experimental animals. Database: the Journal of Biological Databases and Curation, 2012, 2012, bas031-bas031.	3.0	135
20	T3DB: a comprehensively annotated database of common toxins and their targets. Nucleic Acids Research, 2010, 38, D781-D786.	14.5	126
21	ECMDB: The E. coli Metabolome Database. Nucleic Acids Research, 2012, 41, D625-D630.	14.5	122
22	ECMDB 2.0: A richer resource for understanding the biochemistry of <i>E. coli </i> /i>. Nucleic Acids Research, 2016, 44, D495-D501.	14.5	121
23	Exposome-Explorer: a manually-curated database on biomarkers of exposure to dietary and environmental factors. Nucleic Acids Research, 2017, 45, D979-D984.	14.5	109
24	Pathways with PathWhiz. Nucleic Acids Research, 2015, 43, W552-W559.	14.5	27
25	BioSpider: a web server for automating metabolome annotations. Pacific Symposium on Biocomputing Pacific Symposium on Biocomputing, 2007, , 145-56.	0.7	12