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List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2425077/publications.pdf>

Version: 2024-02-01

22
papers

1,099
citations

623734

14
h-index

713466

21
g-index

25
all docs

25
docs citations

25
times ranked

1513
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome-wide analysis of mRNAs targeted to yeast mitochondria. EMBO Reports, 2002, 3, 159-164.	4.5	285
2	Genome microarray analysis of transcriptional activation in multidrug resistance yeast mutants. FEBS Letters, 2000, 470, 156-160.	2.8	243
3	New Insights into the Pleiotropic Drug Resistance Network from Genome-Wide Characterization of the YRR1 Transcription Factor Regulation System. Molecular and Cellular Biology, 2002, 22, 2642-2649.	2.3	95
4	An artificial transcription activator mimics the genome-wide properties of the yeast Pdr1 transcription factor. EMBO Reports, 2001, 2, 493-498.	4.5	59
5	A General Strategy to Uncover Transcription Factor Properties Identifies a New Regulator of Drug Resistance in Yeast. Journal of Biological Chemistry, 2003, 278, 11427-11432.	3.4	56
6	The eTOX Data-Sharing Project to Advance in Silico Drug-Induced Toxicity Prediction. International Journal of Molecular Sciences, 2014, 15, 21136-21154.	4.1	56
7	Legacy data sharing to improve drug safety assessment: the eTOX project. Nature Reviews Drug Discovery, 2017, 16, 811-812.	46.4	56
8	Robust and tissue-independent gender-specific transcript biomarkers. Biomarkers, 2013, 18, 436-445.	1.9	32
9	Transcriptomes, transcription activators and microarrays. FEBS Letters, 2001, 498, 140-144.	2.8	30
10	yMGV: a database for visualization and data mining of published genome-wide yeast expression data. Nucleic Acids Research, 2001, 29, 63e-63.	14.5	28
11	Computational methods for early predictive safety assessment from biological and chemical data. Expert Opinion on Drug Metabolism and Toxicology, 2011, 7, 1497-1511.	3.3	28
12	yMGV: helping biologists with yeast microarray data mining. Nucleic Acids Research, 2002, 30, 76-79.	14.5	27
13	Transcriptional induction of repair genes during slowing of replication in irradiated <i>Saccharomyces cerevisiae</i> . Mutation Research DNA Repair, 2001, 487, 157-172.	3.7	24
14	Value of shared preclinical safety studies – The eTOX database. Toxicology Reports, 2015, 2, 210-221.	3.3	21
15	OntoBrowser: a collaborative tool for curation of ontologies by subject matter experts. Bioinformatics, 2017, 33, 148-149.	4.1	14
16	Ten simple rules to power drug discovery with data science. PLoS Computational Biology, 2020, 16, e1008126.	3.2	14
17	Arrayplot for visualization and normalization of cDNA microarray data. Bioinformatics, 2002, 18, 888-889.	4.1	8
18	yMGV: a cross-species expression data mining tool. Nucleic Acids Research, 2004, 32, 323D-325.	14.5	6

#	ARTICLE	IF	CITATIONS
19	MiCoViTo: a tool for gene-centric comparison and visualization of yeast transcriptome states. BMC Bioinformatics, 2004, 5, 20.	2.6	4
20	Generating Modeling Data From Repeat-Dose Toxicity Reports. Toxicological Sciences, 2018, 162, 287-300.	3.1	4
21	Computational Methods and Bioinformatic Tools. , 0, , 769-904.		0
22	Coronary Artery Disease Polygenic Risk Score Identifies Patients at Higher Risk for Recurrent Cardiovascular Events in the CANTOS Trial. Circulation Genomic and Precision Medicine, 2021, , CIRCGEN121003440.	3.6	0