

Osamu Ogasawara

List of Publications by Year in descending order

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Version: 2024-02-01

17
papers

1,143
citations

623734

14
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839539

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20
all docs

20
docs citations

20
times ranked

2749
citing authors

#	ARTICLE	IF	CITATIONS
1	DNA Data Bank of Japan (DDBJ) update report 2021. <i>Nucleic Acids Research</i> , 2022, 50, D102-D105.	14.5	22
2	DDBJ update: streamlining submission and access of human data. <i>Nucleic Acids Research</i> , 2021, 49, D71-D75.	14.5	38
3	Practical guide for managing large-scale human genome data in research. <i>Journal of Human Genetics</i> , 2021, 66, 39-52.	2.3	31
4	Accumulating computational resource usage of genomic data analysis workflow to optimize cloud computing instance selection. <i>GigaScience</i> , 2019, 8, .	6.4	8
5	DDBJ Database updates and computational infrastructure enhancement. <i>Nucleic Acids Research</i> , 2019, 48, D45-D50.	14.5	28
6	DDBJ update: the Genomic Expression Archive (GEA) for functional genomics data. <i>Nucleic Acids Research</i> , 2019, 47, D69-D73.	14.5	36
7	MACON: a web tool for computing DNA methylation data obtained by the Illumina Infinium Human DNA methylation BeadArray. <i>Epigenomics</i> , 2018, 10, 249-258.	2.1	21
8	DNA Data Bank of Japan: 30th anniversary. <i>Nucleic Acids Research</i> , 2018, 46, D30-D35.	14.5	51
9	ChIP-Atlas: a data-mining suite powered by full integration of public ChIP-seq data. <i>EMBO Reports</i> , 2018, 19, .	4.5	544
10	RefEx, a reference gene expression dataset as a web tool for the functional analysis of genes. <i>Scientific Data</i> , 2017, 4, 170105.	5.3	55
11	DNA Data Bank of Japan. <i>Nucleic Acids Research</i> , 2017, 45, D25-D31.	14.5	78
12	DNA data bank of Japan (DDBJ) progress report. <i>Nucleic Acids Research</i> , 2016, 44, D51-D57.	14.5	76
13	The DDBJ Japanese Genotype-phenotype Archive for genetic and phenotypic human data. <i>Nucleic Acids Research</i> , 2015, 43, D18-D22.	14.5	57
14	DDBJ new system and service refactoring. <i>Nucleic Acids Research</i> , 2012, 41, D25-D29.	14.5	37
15	On Theoretical Models of Gene Expression Evolution with Random Genetic Drift and Natural Selection. <i>PLoS ONE</i> , 2009, 4, e7943.	2.5	7
16	BodyMap-Xs: anatomical breakdown of 17 million animal ESTs for cross-species comparison of gene expression. <i>Nucleic Acids Research</i> , 2006, 34, D628-D631.	14.5	26
17	Zipf's law and human transcriptomes: an explanation with an evolutionary model. <i>Comptes Rendus - Biologies</i> , 2003, 326, 1097-1101.	0.2	14