

Katherine James

List of Publications by Year in descending order

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Version: 2024-02-01

26
papers

564
citations

759055

12
h-index

677027

22
g-index

33
all docs

33
docs citations

33
times ranked

1087
citing authors

#	ARTICLE	IF	CITATIONS
1	OUP accepted manuscript. Briefings in Functional Genomics, 2022, , .	1.3	3
2	Computational Network Inference for Bacterial Interactomics. <i>MSystems</i> , 2022, 7, e0145621.	1.7	5
3	Virtual Parts Repository 2: Model-Driven Design of Genetic Regulatory Circuits. <i>ACS Synthetic Biology</i> , 2021, 10, 3304-3315.	1.9	6
4	High intrinsic hydrolytic activity of cyanobacterial RNA polymerase compensates for the absence of transcription proofreading factors. <i>Nucleic Acids Research</i> , 2020, 48, 1341-1352.	6.5	10
5	Capturing Multicellular System Designs Using Synthetic Biology Open Language (SBOL). <i>ACS Synthetic Biology</i> , 2020, 9, 2410-2417.	1.9	1
6	Complete representation of a tapeworm genome reveals chromosomes capped by centromeres, necessitating a dual role in segregation and protection. <i>BMC Biology</i> , 2020, 18, 165.	1.7	19
7	The tapeworm interactome: inferring confidence scored protein-protein interactions from the proteome of <i>Hymenolepis microstoma</i> . <i>BMC Genomics</i> , 2020, 21, 346.	1.2	4
8	The gene-rich genome of the scallop <i>Pecten maximus</i> . <i>GigaScience</i> , 2020, 9, .	3.3	53
9	Symptom-based stratification of patients with primary Sjögren's syndrome: multi-dimensional characterisation of international observational cohorts and reanalyses of randomised clinical trials. <i>Lancet Rheumatology</i> , The, 2019, 1, e85-e94.	2.2	76
10	The Synarcual of the Little Skate, <i>Leucoraja erinacea</i> : Novel Development Among the Vertebrates. <i>Frontiers in Ecology and Evolution</i> , 2019, 7, .	1.1	12
11	An ancient germ cell-specific RNA-binding protein protects the germline from cryptic splice site poisoning. <i>ELife</i> , 2019, 8, .	2.8	22
12	B-cell activity markers are associated with different disease activity domains in primary Sjögren's syndrome. <i>Rheumatology</i> , 2018, 57, 1222-1227.	0.9	23
13	Genome-wide transcriptome profiling and spatial expression analyses identify signals and switches of development in tapeworms. <i>EvoDevo</i> , 2018, 9, 21.	1.3	30
14	Androgen-dependent alternative mRNA isoform expression in prostate cancer cells. <i>F1000Research</i> , 2018, 7, 1189.	0.8	16
15	A link between transcription fidelity and pausing <i>in vivo</i> . <i>Transcription</i> , 2017, 8, 99-105.	1.7	13
16	Deep sequencing approaches for the analysis of prokaryotic transcriptional boundaries and dynamics. <i>Methods</i> , 2017, 120, 76-84.	1.9	10
17	Single-peptide DNA-dependent RNA polymerase homologous to multi-subunit RNA polymerase. <i>Nature Communications</i> , 2017, 8, 15774.	5.8	22
18	Misincorporation by RNA polymerase is a major source of transcription pausing <i>in vivo</i> . <i>Nucleic Acids Research</i> , 2016, 45, gkw969.	6.5	31

#	ARTICLE	IF	CITATIONS
19	Transformer2 proteins protect breast cancer cells from accumulating replication stress by ensuring productive splicing of checkpoint kinase 1. <i>Frontiers of Chemical Science and Engineering</i> , 2016, 10, 186-195.	2.3	3
20	Glycosylation is an Androgen-Regulated Process Essential for Prostate Cancer Cell Viability. <i>EBioMedicine</i> , 2016, 8, 103-116.	2.7	76
21	Eligibility for clinical trials in primary Sjögren's syndrome: lessons from the UK Primary Sjögren's Syndrome Registry. <i>Rheumatology</i> , 2015, 55, kev373.	0.9	9
22	A Transcriptional Signature of Fatigue Derived from Patients with Primary Sjögren's Syndrome. <i>PLoS ONE</i> , 2015, 10, e0143970.	1.1	45
23	Human Tra2 proteins jointly control a CHEK1 splicing switch among alternative and constitutive target exons. <i>Nature Communications</i> , 2014, 5, 4760.	5.8	47
24	BacillOndex: An Integrated Data Resource for Systems and Synthetic Biology. <i>Journal of Integrative Bioinformatics</i> , 2013, 10, 103-116.	1.0	12
25	Is newer better?â€”evaluating the effects of data curation on integrated analyses in <i>Saccharomyces cerevisiae</i> . <i>Integrative Biology (United Kingdom)</i> , 2012, 4, 715-727.	0.6	2
26	Integration of Full-Coverage Probabilistic Functional Networks with Relevance to Specific Biological Processes. <i>Lecture Notes in Computer Science</i> , 2009, , 31-46.	1.0	8