

Kathryn Crouch

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2064827/publications.pdf>

Version: 2024-02-01

23
papers

1,264
citations

687363

13
h-index

794594

19
g-index

36
all docs

36
docs citations

36
times ranked

1953
citing authors

#	ARTICLE	IF	CITATIONS
1	FungiDB: An Integrated Bioinformatic Resource for Fungi and Oomycetes. Journal of Fungi (Basel,) Tj ETQq1 1 0.784314 rgBT/Overlook	3.5	309
2	VEuPathDB: the eukaryotic pathogen, vector and host bioinformatics resource center. Nucleic Acids Research, 2022, 50, D898-D911.	14.5	277
3	EuPathDB: the eukaryotic pathogen genomics database resource. Nucleic Acids Research, 2017, 45, D581-D591.	14.5	191
4	EuPathDB: The Eukaryotic Pathogen Genomics Database Resource. Methods in Molecular Biology, 2018, 1757, 69-113.	0.9	80
5	A Leishmania infantum genetic marker associated with miltefosine treatment failure for visceral leishmaniasis. EBioMedicine, 2018, 36, 83-91.	6.1	56
6	Validation of the protein kinase CLK3 as a multistage cross-species malarial drug target. Science, 2019, 365, .	12.6	51
7	Ribonuclease H1-targeted R-loops in surface antigen gene expression sites can direct trypanosome immune evasion. PLoS Genetics, 2018, 14, e1007729.	3.5	40
8	Characterization of the immunoglobulin repertoire of the spiny dogfish (Squalus acanthias). Developmental and Comparative Immunology, 2012, 36, 665-679.	2.3	38
9	Humoral immune response of the small-spotted catshark, Scyliorhinus canicula. Fish and Shellfish Immunology, 2013, 34, 1158-1169.	3.6	38
10	TrypanoCyc: a community-led biochemical pathways database for Trypanosoma brucei. Nucleic Acids Research, 2015, 43, D637-D644.	14.5	35
11	Trypanosoma brucei ribonuclease H2A is an essential R-loop processing enzyme whose loss causes DNA damage during transcription initiation and antigenic variation. Nucleic Acids Research, 2019, 47, 9180-9197.	14.5	32
12	Genome-wide mapping reveals conserved and diverged R-loop activities in the unusual genetic landscape of the African trypanosome genome. Nucleic Acids Research, 2018, 46, 11789-11805.	14.5	27
13	Trypanosoma brucei ATR Links DNA Damage Signaling during Antigenic Variation with Regulation of RNA Polymerase I-Transcribed Surface Antigens. Cell Reports, 2020, 30, 836-851.e5.	6.4	24
14	Conditional knockout of RAD51-related genes in Leishmania major reveals a critical role for homologous recombination during genome replication. PLoS Genetics, 2020, 16, e1008828.	3.5	21
15	Genome duplication in Leishmania major relies on persistent subtelomeric DNA replication. ELife, 2020, 9, .	6.0	17
16	Divergent metabolism between Trypanosoma congolense and Trypanosoma brucei results in differential sensitivity to metabolic inhibition. PLoS Pathogens, 2021, 17, e1009734.	4.7	11
17	Next-Generation Analysis of Trypanosomatid Genome Stability and Instability. Methods in Molecular Biology, 2020, 2116, 225-262.	0.9	2
18	Transcriptional differentiation of Trypanosoma brucei during in vitro acquisition of resistance to acoziborole. PLoS Neglected Tropical Diseases, 2021, 15, e0009939.	3.0	2

#	ARTICLE	IF	CITATIONS
19	Bringing bioinformatics to schools with the 4273pi project. PLoS Computational Biology, 2022, 18, e1009705.	3.2	2
20	Title is missing!. , 2020, 16, e1008828.		0
21	Title is missing!. , 2020, 16, e1008828.		0
22	Title is missing!. , 2020, 16, e1008828.		0
23	Title is missing!. , 2020, 16, e1008828.		0