

Steven A Sullivan

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10973135/publications.pdf>

Version: 2024-02-01

27
papers

4,314
citations

471509

17
h-index

552781

26
g-index

28
all docs

28
docs citations

28
times ranked

6373
citing authors

#	ARTICLE	IF	CITATIONS
1	Small RNAs Are Implicated in Regulation of Gene and Transposable Element Expression in the Protist <i>Trichomonas vaginalis</i> . <i>MSphere</i> , 2021, 6, .	2.9	4
2	The effectiveness of malaria camps as part of the Durgama Anchalare Malaria Nirakaran (DAMaN) program in Odisha, India: study protocol for a cluster-assigned quasi-experimental study. <i>Global Health Action</i> , 2021, 14, 1886458.	1.9	7
3	Household and individual level risk factors associated with declining malaria incidence in Meghalaya, India: implications for malaria elimination in low-endemic settings. <i>Malaria Journal</i> , 2021, 20, 460.	2.3	12
4	Malaria in Sundargarh district, Odisha, India: Epidemiological and behavioral aspects from surveys. <i>Acta Tropica</i> , 2020, 211, 105647.	2.0	9
5	Defining symptoms of malaria in India in an era of asymptomatic infections. <i>Malaria Journal</i> , 2020, 19, 237.	2.3	7
6	The burden of submicroscopic and asymptomatic malaria in India revealed from epidemiology studies at three varied transmission sites in India. <i>Scientific Reports</i> , 2019, 9, 17095.	3.3	44
7	Genetic Indicators of Drug Resistance in the Highly Repetitive Genome of <i>Trichomonas vaginalis</i> . <i>Genome Biology and Evolution</i> , 2017, 9, 1658-1672.	2.5	44
8	The use of mosquito repellents at three sites in India with declining malaria transmission: surveys in the community and clinic. <i>Parasites and Vectors</i> , 2016, 9, 418.	2.5	27
9	What is the value of reactive case detection in malaria control? A case-study in India and a systematic review. <i>Malaria Journal</i> , 2016, 15, 67.	2.3	54
10	Characterizing the genetic diversity of the monkey malaria parasite <i>Plasmodium cynomolgi</i> . <i>Infection, Genetics and Evolution</i> , 2016, 40, 243-252.	2.3	23
11	The biology of <i>Plasmodium vivax</i> explored through genomics. <i>Annals of the New York Academy of Sciences</i> , 2015, 1342, 53-61.	3.8	25
12	<i>Plasmodium cynomolgi</i> genome sequences provide insight into <i>Plasmodium vivax</i> and the monkey malaria clade. <i>Nature Genetics</i> , 2012, 44, 1051-1055.	21.4	172
13	Malaria in India: The Center for the Study of Complex Malaria in India. <i>Acta Tropica</i> , 2012, 121, 267-273.	2.0	115
14	Microsatellite polymorphism in the sexually transmitted human pathogen <i>Trichomonas vaginalis</i> indicates a genetically diverse parasite. <i>Molecular and Biochemical Parasitology</i> , 2011, 175, 30-38.	1.1	48
15	Three Genomes from the Phylum <i>Acidobacteria</i> Provide Insight into the Lifestyles of These Microorganisms in Soils. <i>Applied and Environmental Microbiology</i> , 2009, 75, 2046-2056.	3.1	804
16	Comparative genomics of the neglected human malaria parasite <i>Plasmodium vivax</i> . <i>Nature</i> , 2008, 455, 757-763.	27.8	756
17	Noggin signaling from <i>Xenopus</i> animal blastomere lineages promotes a neural fate in neighboring vegetal blastomere lineages. <i>Developmental Dynamics</i> , 2007, 236, 171-183.	1.8	4
18	Comparative Genomic Evidence for a Close Relationship between the Dimorphic Prosthecate Bacteria <i>Hyphomonas neptunium</i> and <i>Caulobacter crescentus</i> . <i>Journal of Bacteriology</i> , 2006, 188, 6841-6850.	2.2	57

#	ARTICLE	IF	CITATIONS
19	The psychrophilic lifestyle as revealed by the genome sequence of <i>Colwellia psychrerythraea</i> 34H through genomic and proteomic analyses. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 10913-10918.	7.1	529
20	Major Structural Differences and Novel Potential Virulence Mechanisms from the Genomes of Multiple <i>Campylobacter</i> Species. <i>PLoS Biology</i> , 2005, 3, e15.	5.6	483
21	The genome sequence of the anaerobic, sulfate-reducing bacterium <i>Desulfovibrio vulgaris</i> Hildenborough. <i>Nature Biotechnology</i> , 2004, 22, 554-559.	17.5	559
22	Genome sequence of <i>Silicibacter pomeroyi</i> reveals adaptations to the marine environment. <i>Nature</i> , 2004, 432, 910-913.	27.8	415
23	Characterization of sequence variability in nucleosome core histone folds. <i>Proteins: Structure, Function and Bioinformatics</i> , 2003, 52, 454-465.	2.6	18
24	Mining Core Histone Sequences from Public Protein Databases. <i>Methods in Enzymology</i> , 2003, 375, 3-20.	1.0	3
25	Multiple maternal influences on dorsal-ventral fate of <i>Xenopus</i> animal blastomeres. <i>Developmental Dynamics</i> , 2002, 225, 581-587.	1.8	18
26	<i>foxD5a</i> , a <i>Xenopus</i> Winged Helix Gene, Maintains an Immature Neural Ectoderm via Transcriptional Repression That Is Dependent on the C-Terminal Domain. <i>Developmental Biology</i> , 2001, 232, 439-457.	2.0	64
27	Early Events in Frog Blastomere Fate Determination. , 1999, , 297-321.		12