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 4,314 17 26 papers citations h-index g-index

times ranked

citing authors

docs citations

all docs

#	Article	IF	CITATIONS
1	Three Genomes from the Phylum <i>Acidobacteria</i> Provide Insight into the Lifestyles of These Microorganisms in Soils. Applied and Environmental Microbiology, 2009, 75, 2046-2056.	3.1	804
2	Comparative genomics of the neglected human malaria parasite Plasmodium vivax. Nature, 2008, 455, 757-763.	27.8	756
3	The genome sequence of the anaerobic, sulfate-reducing bacterium Desulfovibrio vulgaris Hildenborough. Nature Biotechnology, 2004, 22, 554-559.	17.5	559
4	The psychrophilic lifestyle as revealed by the genome sequence of Colwellia psychrerythraea 34H through genomic and proteomic analyses. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 10913-10918.	7.1	529
5	Major Structural Differences and Novel Potential Virulence Mechanisms from the Genomes of Multiple Campylobacter Species. PLoS Biology, 2005, 3, e15.	5. 6	483
6	Genome sequence of Silicibacter pomeroyi reveals adaptations to the marine environment. Nature, 2004, 432, 910-913.	27.8	415
7	Plasmodium cynomolgi genome sequences provide insight into Plasmodium vivax and the monkey malaria clade. Nature Genetics, 2012, 44, 1051-1055.	21.4	172
8	Malaria in India: The Center for the Study of Complex Malaria in India. Acta Tropica, 2012, 121, 267-273.	2.0	115
9	foxD5a, a Xenopus Winged Helix Gene, Maintains an Immature Neural Ectoderm via Transcriptional Repression That Is Dependent on the C-Terminal Domain. Developmental Biology, 2001, 232, 439-457.	2.0	64
10	Comparative Genomic Evidence for a Close Relationship between the Dimorphic Prosthecate Bacteria Hyphomonas neptunium and Caulobacter crescentus. Journal of Bacteriology, 2006, 188, 6841-6850.	2.2	57
11	What is the value of reactive case detection in malaria control? A case-study in India and a systematic review. Malaria Journal, 2016, 15, 67.	2.3	54
12	Microsatellite polymorphism in the sexually transmitted human pathogen Trichomonas vaginalis indicates a genetically diverse parasite. Molecular and Biochemical Parasitology, 2011, 175, 30-38.	1.1	48
13	Genetic Indicators of Drug Resistance in the Highly Repetitive Genome of Trichomonas vaginalis. Genome Biology and Evolution, 2017, 9, 1658-1672.	2.5	44
14	The burden of submicroscopic and asymptomatic malaria in India revealed from epidemiology studies at three varied transmission sites in India. Scientific Reports, 2019, 9, 17095.	3.3	44
15	The use of mosquito repellents at three sites in India with declining malaria transmission: surveys in the community and clinic. Parasites and Vectors, 2016, 9, 418.	2.5	27
16	The biology of <i>Plasmodium vivax</i> explored through genomics. Annals of the New York Academy of Sciences, 2015, 1342, 53-61.	3.8	25
17	Characterizing the genetic diversity of the monkey malaria parasite Plasmodium cynomolgi. Infection, Genetics and Evolution, 2016, 40, 243-252.	2.3	23
18	Multiple maternal influences on dorsal-ventral fate of Xenopus animal blastomeres. Developmental Dynamics, 2002, 225, 581-587.	1.8	18

#	Article	IF	CITATIONS
19	Characterization of sequence variability in nucleosome core histone folds. Proteins: Structure, Function and Bioinformatics, 2003, 52, 454-465.	2.6	18
20	Early Events in Frog Blastomere Fate Determination. , 1999, , 297-321.		12
21	Household and individual level risk factors associated with declining malaria incidence in Meghalaya, India: implications for malaria elimination in low-endemic settings. Malaria Journal, 2021, 20, 460.	2.3	12
22	Malaria in Sundargarh district, Odisha, India: Epidemiological and behavioral aspects from surveys. Acta Tropica, 2020, 211, 105647.	2.0	9
23	Defining symptoms of malaria in India in an era of asymptomatic infections. Malaria Journal, 2020, 19, 237.	2.3	7
24	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. Global Health Action, 2021, 14, 1886458.	1.9	7
25	Noggin signaling fromXenopus animal blastomere lineages promotes a neural fate in neighboring vegetal blastomere lineages. Developmental Dynamics, 2007, 236, 171-183.	1.8	4
26	Small RNAs Are Implicated in Regulation of Gene and Transposable Element Expression in the Protist Trichomonas vaginalis. MSphere, $2021, 6, \ldots$	2.9	4
27	Mining Core Histone Sequences from Public Protein Databases. Methods in Enzymology, 2003, 375, 3-20.	1.0	3