Adam L Bazinet

List of Publications by Year
in descending order

14 papers	823 citations	759190 12 h-index	1125717 13 g-index
17 all docs	17 docs citations	17 times ranked	1604 citing authors

#	Article	IF	Citations
1	A Gateway for Phylogenetic Analysis Powered by Grid Computing Featuring GARLI 2.0. Systematic Biology, 2014, 63, 812-818.	5.6	195
2	A comparative evaluation of sequence classification programs. BMC Bioinformatics, 2012, 13, 92.	2.6	99
3	Pan-genome and phylogeny of Bacillus cereus sensu lato. BMC Evolutionary Biology, 2017, 17, 176.	3.2	92
4	Can RNA-Seq Resolve the Rapid Radiation of Advanced Moths and Butterflies (Hexapoda: Lepidoptera:) Tj ETQq0 0 0 rgBT /Overlock 10 T	2.5	89
5	Further progress on the phylogeny of <scp>N</scp>octuoidea (<scp>I</scp>nsecta:) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 582	3.9	54
6	Prey preference follows phylogeny: evolutionary dietary patterns within the marine gastropod group Cladobranchia (Gastropoda: Heterobranchia: Nudibranchia). BMC Evolutionary Biology, 2017, 17, 221.	3.2	53
7	Relationships within Cladobranchia (Gastropoda: Nudibranchia) based on RNA-Seq data: an initial investigation. Royal Society Open Science, 2015, 2, 150196.	2.4	44
8	Biological Sexing of a 4000-Year-Old Egyptian Mummy Head to Assess the Potential of Nuclear DNA Recovery from the Most Damaged and Limited Forensic Specimens. Genes, 2018, 9, 135.	2.4	39
9	Phylotranscriptomics resolves ancient divergences in the Lepidoptera. Systematic Entomology, 2017, 42, 305-316.	3.9	38
10	Fast and simple protein-alignment-guided assembly of orthologous gene families from microbiome sequencing reads. Microbiome, 2017, 5, 11.	11.1	31
11	Computing the Tree of Life: Leveraging the Power of Desktop and Service Grids. , 2011, , .		29
12	Historical introgression drives pervasive mitochondrial admixture between two species of pelagic sharks. Molecular Phylogenetics and Evolution, 2017, 110, 122-126.	2.7	24
13	BLAST-based validation of metagenomic sequence assignments. PeerJ, 2018, 6, e4892.	2.0	16
14	Draft Genome Sequences of Five Historical Bacillus anthracis Strains. Microbiology Resource Announcements, 2020, 9, .	0.6	0