

Jessica W Leigh

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/12057083/publications.pdf>

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9
papers

5,553
citations

1039880

9
h-index

1474057

9
g-index

9
all docs

9
docs citations

9
times ranked

9770
citing authors

#	ARTICLE	IF	CITATIONS
1	<scp>popart</scp>: fullâ€feature software for haplotype network construction. <i>Methods in Ecology and Evolution</i> , 2015, 6, 1110-1116.	2.2	4,199
2	Monte Carlo Strategies for Selecting Parameter Values in Simulation Experiments. <i>Systematic Biology</i> , 2015, 64, 741-751.	2.7	21
3	Hybridization may facilitate in situ survival of endemic species through periods of climate change. <i>Nature Climate Change</i> , 2013, 3, 1039-1043.	8.1	94
4	Evaluating Phylogenetic Congruence in the Post-Genomic Era. <i>Genome Biology and Evolution</i> , 2011, 3, 571-587.	1.1	46
5	Let Them Fall Where They May: Congruence Analysis in Massive Phylogenetically Messy Data Sets. <i>Molecular Biology and Evolution</i> , 2011, 28, 2773-2785.	3.5	43
6	A multi-locus time-calibrated phylogeny of the brown algae (Heterokonta, Ochrophyta, Phaeophyceae): Investigating the evolutionary nature of the â€brown algal crown radiationâ€. <i>Molecular Phylogenetics and Evolution</i> , 2010, 56, 659-674.	1.2	261
7	Network analyses structure genetic diversity in independent genetic worlds. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 127-132.	3.3	265
8	Phylogenomic analyses support the monophyly of Excavata and resolve relationships among eukaryotic â€supergroupsâ€. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 3859-3864.	3.3	444
9	Testing Congruence in Phylogenomic Analysis. <i>Systematic Biology</i> , 2008, 57, 104-115.	2.7	180