James H Degnan

List of Publications by Year in descending order

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IAMES H DECNAN

#	Article	IF	CITATIONS
1	Trying out a million genes to find the perfect pair with <i>RTIST</i> . Bioinformatics, 2022, 38, 3565-3573.	4.1	0
2	Statistical inconsistency of the unrooted minimize deep coalescence criterion. PLoS ONE, 2021, 16, e0251107.	2.5	1
3	Heuristics for unrooted, unranked, and ranked anomaly zones under birth-death models. Molecular Phylogenetics and Evolution, 2021, 161, 107162.	2.7	0
4	Meng and Kubatko (2009): Modeling hybridization with coalescence. Theoretical Population Biology, 2020, 133, 36-37.	1.1	1
5	Probabilities of Unranked and Ranked Anomaly Zones under Birth–Death Models. Molecular Biology and Evolution, 2020, 37, 1480-1494.	8.9	3
6	PRANC: ML species tree estimation from the ranked gene trees under coalescence. Bioinformatics, 2020, 36, 4819-4821.	4.1	7
7	An initiative to implement immediate postpartum long-acting reversible contraception in rural New Mexico. American Journal of Obstetrics and Gynecology, 2020, 222, S911.e1-S911.e7.	1.3	18
8	Species Tree Inference from Gene Splits by Unrooted STAR Methods. IEEE/ACM Transactions on Computational Biology and Bioinformatics, 2018, 15, 337-342.	3.0	32
9	Split Probabilities and Species Tree Inference Under the Multispecies Coalescent Model. Bulletin of Mathematical Biology, 2018, 80, 64-103.	1.9	4
10	Modeling Hybridization Under the Network Multispecies Coalescent. Systematic Biology, 2018, 67, 786-799.	5.6	97
11	Displayed Trees Do Not Determine Distinguishability Under the Network Multispecies Coalescent. Systematic Biology, 2017, 66, syw097.	5.6	34
12	Inferring rooted species trees from unrooted gene trees using approximate Bayesian computation. Molecular Phylogenetics and Evolution, 2017, 116, 13-24.	2.7	12
13	Does Gene Tree Discordance Explain the Mismatch between Macroevolutionary Models and Empirical Patterns of Tree Shape and Branching Times?. Systematic Biology, 2016, 65, 628-639.	5.6	18
14	There are no caterpillars in a wicked forest. Theoretical Population Biology, 2015, 105, 17-23.	1.1	10
15	Hybrid-Lambda: simulation of multiple merger and Kingman gene genealogies in species networks and species trees. BMC Bioinformatics, 2015, 16, 292.	2.6	30
16	Anomalous Unrooted Gene Trees. Systematic Biology, 2013, 62, 574-590.	5.6	64
17	Evaluating variations on the STAR algorithm for relative efficiency and sample sizes needed to reconstruct species trees. Pacific Symposium on Biocomputing Pacific Symposium on Biocomputing, 2013, , 262-72.	0.7	2
18	A Characterization of the Set of Species Trees that Produce Anomalous Ranked Gene Trees. IEEE/ACM Transactions on Computational Biology and Bioinformatics, 2012, 9, 1558-1568.	3.0	15

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19	The probability distribution of ranked gene trees on a species tree. Mathematical Biosciences, 2012, 235, 45-55.	1.9	44
20	EVALUATING VARIATIONS ON THE STAR ALGORITHM FOR RELATIVE EFFICIENCY AND SAMPLE SIZES NEEDED TO RECONSTRUCT SPECIES TREES. , 2012, , .		1
21	Identifying the rooted species tree from the distribution of unrooted gene trees under the coalescent. Journal of Mathematical Biology, 2011, 62, 833-862.	1.9	124
22	Performance of Matrix Representation with Parsimony for Inferring Species from Gene Trees. Statistical Applications in Genetics and Molecular Biology, 2011, 10, .	0.6	8
23	Properties of Consensus Methods for Inferring Species Trees from Gene Trees. Systematic Biology, 2009, 58, 35-54.	5.6	135
24	Gene tree discordance, phylogenetic inference and the multispecies coalescent. Trends in Ecology and Evolution, 2009, 24, 332-340.	8.7	1,500
25	Genomics and genome-wide association studies: An integrative approach to expression QTL mapping. Genomics, 2008, 92, 129-133.	2.9	18
26	Discordance of Species Trees with Their Most Likely Gene Trees. PLoS Genetics, 2006, 2, e68.	3.5	761
27	GENE TREE DISTRIBUTIONS UNDER THE COALESCENT PROCESS. Evolution; International Journal of Organic Evolution, 2005, 59, 24-37.	2.3	296
28	GENE TREE DISTRIBUTIONS UNDER THE COALESCENT PROCESS. Evolution; International Journal of Organic Evolution, 2005, 59, 24.	2.3	91
29	Gene tree distributions under the coalescent process. Evolution; International Journal of Organic Evolution, 2005, 59, 24-37.	2.3	106