

CORRECTION



## Correction to: Coalescence-based species delimitation using genome-wide data reveals hidden diversity in a cosmopolitan group of lichens

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The publisher regrets that the Table 3 of the original published version of this article is incorrect thus the table is now presented correctly.

The original article has been corrected.

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The online version of the original article can be found at <https://doi.org/10.1007/s13127-019-00424-0>

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**Table 3** Species assignment estimations from the two runs of STACEY and tr2-delimitation. For STACEY, only the best five hypotheses are presented. Species level units are color-coded in shades of gray. \* and + denote additional OTUs distinguished as separate species in suboptimal models

count	fraction	number of clusters	Specimens and prior assignment to species																									
			obtusata	odontella																								
26 loci, not accounting for linkage																												
STACEY																												
15295	0.6126	6																										
1887	0.0755	7																										
1605	0.0642	7																										
1241	0.0497	5																										
483	0.0193	7																										
tr2-delimitation																												
NA	NA	6																										
24 loci, accounting for linkage																												
STACEY																												
25794	0.7344	6																										
1870	0.0532	7																										
747	0.0213	7																										
700	0.0199	7																										
394	0.0112	7																										
tr2-delimitation																												
NA	NA	5																										
20 loci, excluding loci under possible environmental selection																												
STACEY																												
1725	0.0453	6																										
1429	0.0375	8																										
820	0.0215	7																										
355	0.0093	9																										
254	0.0067	7																										
tr2-delimitation																												
NA	NA																											