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Corrigendum

Corrigendum to “Phylogenetic relationships, biogeography and speciation in the avian genus *Saxicola*” [Mol. Phylogenet. Evol. 48 (2008) 1145–1154]

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After designing new primers, and additional sequencing for the stonechats, we have discovered that two sequences belonging to two Spanish common stonechats (*Saxicola torquata*) samples (accession numbers EU421091 and EU421081) published in our paper were erroneous. The published phylogenetic relationships of the incorrect sequences grouped the two individuals with an Asian common stonechat and *Saxicola leucura* from Nepal (node F). However, after reamplifying and resequencing these two individuals both sequences now group unambiguously within the European common stonechats (node I; Fig. 1). We regret this mistake and we would like to apologize for any inconvenience. The sequences have been also corrected in the NCBI gene bank database.

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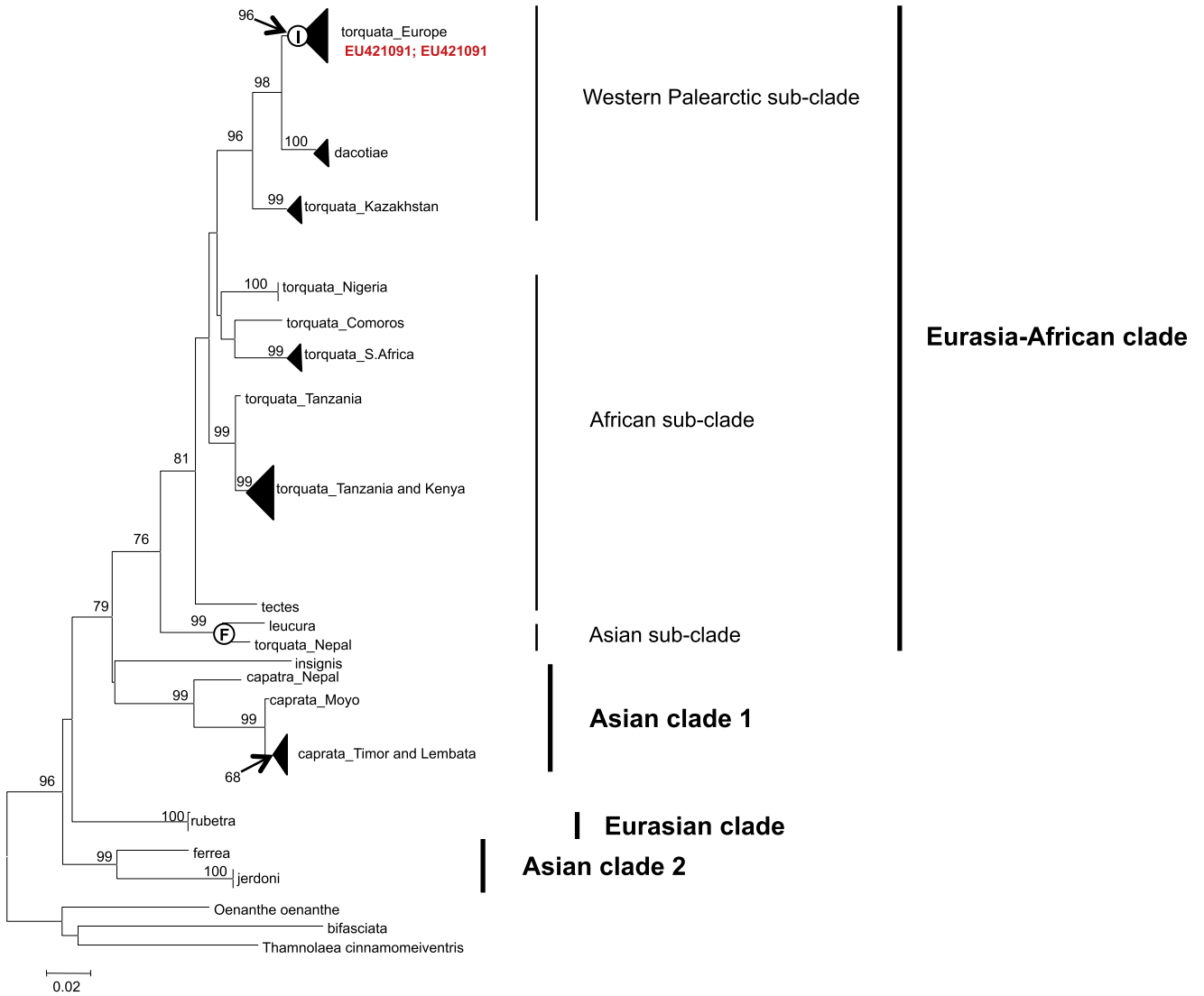


Fig. 1. Maximum likelihood tree (mtDNA cyt b) for *Saxicola* after EU421091 and EU421081 sequences were corrected. Numbers above nodes show the bootstrap support (only > 60% is showed).