

## **Erratum to: Multilocus population analysis of *Gavia immer* (Aves: Gaviidae) mtDNA reveals low genetic diversity and lack of differentiation across the species breeding range**

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The original version of this article unfortunately contained a mistake. The correct information is given below.

The last two sentences of page 313 should read:

Considering that the rate of nucleotide evolution for avian mitochondrial loci is of the order of 0.9 (95 % CI: 0.4–1.4) substitutions per site per million years (Lambert et al. 2002), and that we have analysed 780 sites, it can be estimated that the derived haplotypes originated about 1,068 years ago (95 % CI: 687–2,404). Although this figure should be taken with care, as substitution rates might vary across loci and taxons, it suggests that the extant variation at the *cytb* locus is relatively recent in this species, and that most variants are likely to date well after the Last Glacial Maximum, between 26,500 and 19,000 years ago (Clark et al. 2009).

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