

Female-to-Male Breeding Ratio in Modern Humans—an Analysis Based on Historical Recombinations

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Lohmueller et al. have recently brought to our attention in a Letter to the Editor that we made an error in the way we corrected for unequal male-female population sizes in our original manuscript. This error affected many of our equations, figures, and tables. We are thankful to Lohmueller et al. for identifying this mistake, and we have supplied corrected versions of our equations, tables and figures with our Response to the Letter to the Editor.

The authors regret these errors.

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Pooled Association Tests for Rare Variants in Exon-Resequencing Studies

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As a result of a coding error affecting permutation randomization, Tables 1–3 and Tables S1–S4 in the version of this article published online on May 12 contained slight inaccuracies, which have been corrected in the current version both online and in print. In addition, the text “p value increases from 0.002 to 0.008” on the bottom of p. 5 has been changed to “p value increases from 0.002 to 0.009.” The authors regret this error, and are grateful to Adam Kiezun for correcting it.

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