

## Correction: A Map of Recent Positive Selection in the Human Genome

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Because of a typesetting error, the symbol " $\sigma$ " is incorrectly displayed as "s" in the legends of Figures 1, 2, and 5.

In the Results section, the sentence "These data consist of ~800,000 polymorphic SNPs in a total of 309 unrelated individuals" should be "These data consist of ~800,000 polymorphic SNPs in a total of 209 unrelated individuals."

In the section "Widespread Signals of Recent Selection," the sentence "For example, in simulations matching aspects of the Yoruba data, only 0.1% of windows of 50 consecutive SNPs had more than 16 SNPs with  $|iHS| > 20$ " should be "For example, in simulations matching aspects of the Yoruba data, only 0.1% of windows of 50 consecutive SNPs had more than 16 SNPs with  $|iHS| > 2.0$ ."

This correction note may be found online at DOI: 10.1371/journal.pbio.0040154.

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