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**Supplemental Data**

**Whole-Genome-Sequence-Based Haplotypes  
Reveal Single Origin of the Sickle Allele  
during the Holocene Wet Phase**

**Daniel Shriner and Charles N. Rotimi**

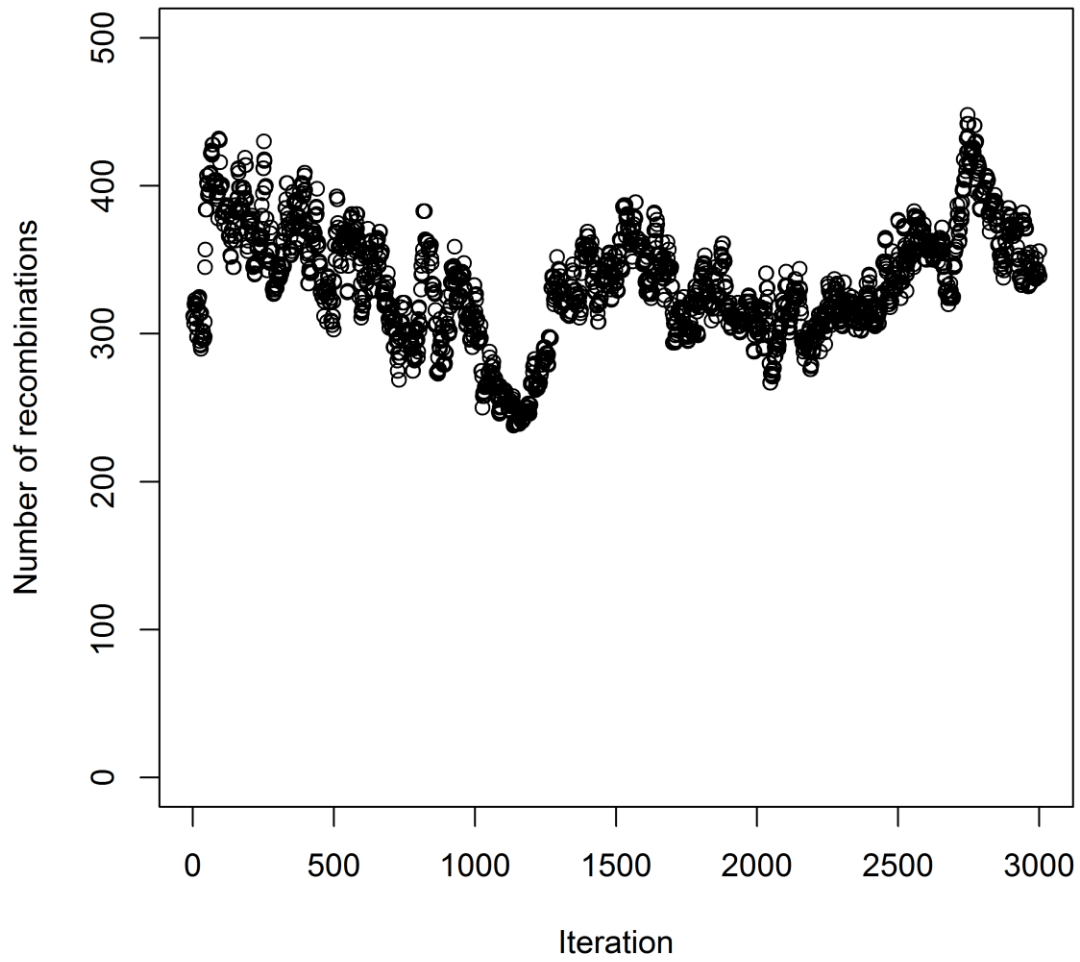


Figure S1. Trace of the number of recombinations. If the target stationary distribution has been reached, then the trace should depict white noise around a horizontal line.