

## Erratum

In the March 2005 issue of the *Journal*, in the article entitled “Accuracy of Haplotype Reconstruction from Haplotype-Tagging Single-Nucleotide Polymorphisms” by Forton et al. (76:438–448), error-rate simulations in the 5q31 region and the linkage disequilibrium maps in figure 3 are based on genotyping data reported by G. Luoni, J. Forton, M. Jallow, E. Sadighi Akha, F. Sisay-Joof, M.

Pinder, N. Hanchard, M. Herbert, M. Kimber, R. Mott, J. Hull, K. Rockett, and D. Kwiatkowski in “Population-Specific Patterns of Linkage Disequilibrium in the Human 5q31 Region” (*Genes Immun* 6:723–727). Forton et al. thank Elham Sadighi Akha, for her contribution to the genotyping data, and apologize for the inadvertent omission of this acknowledgment in the original publication.