## Estimating Odds Ratios in Genome Scans: An Approximate Conditional Likelihood Approach

Arpita Ghosh, Fei Zou, and Fred A. Wright\*

(American Journal of Human Genetics 82, 1064–1074; May 2008)

In the version of this article published online on April 17, 2008, the ratio  $LLR = 2\log(L(\hat{\mu})/L(0))$  incorrectly appeared as  $LLR = -2\log(L(\hat{\mu})/L(0))$  on page 3. In the formula for the specification of *x* under the three genetic models on page 5, under "additive," *x* was incorrectly coded as 0 instead of 1 for the *Aa* genotype. In the Note Added in Proof on page 11, the first author's first initial incorrectly appeared as "E." instead of "H.," and the title of the referenced paper appeared as "Bias-reduced estimators and confidence intervals for logs ratios in genome-wide association studies" instead of "Bias-reduced version of this article now appears online and in print.

\*Correspondence: fwright@bios.unc.edu DOI 10.1016/j.ajhg.2008.04.004. ©2008 by The American Society of Human Genetics. All rights reserved.