

Erratum

In the March 2002 issue of the *Journal*, in the report “A Genomewide Linkage Screen for Relative Hand Skill in Sibling Pairs,” by Francks et al. (70:800–805), there is an error regarding the description of the linkage analysis of chromosome X by the program MAPMAKER/SIBS. On page 802, the report states that “MAPMAKER/SIBS uses only male-male pairings for HE [i.e., Haseman-Elston] regression”; however, the results presented in the report were obtained by MAPMAKER/SIBS version 2.0, which, unlike version 1.1, includes all sibling pairs in the regression and assumes that the difference in phenotype is a function of whether 0 or 1 alleles are inherited identical

by descent from the mother. We note that, in an additional analysis of only male-male pairings (79 total pairs), the peak multipoint Haseman-Elston LOD score between markers DXS993 and DXS991 increased to 1.09 (nominal pointwise $P = .012$). This latter analysis is more comparable with that in the 1998 article by Laval et al. (which is listed in the References list of the report by Francks et al.) and provides additional support for the chromosome X linkage that Laval et al. identified. The error does not affect any of the autosomal linkage results presented in the report. The authors regret this error.