

## CORRIGENDUM

In the paper by R. S. SINGH and M. B. COULTHART (*Genetics* **102**: 437-453) entitled "Genic variation in abundant soluble proteins of *Drosophila melanogaster* and *Drosophila pseudoobscura*," in Table 1, page 440, misalignment of lines within columns for populations ARG, SWE, UKR, CAS and KOR caused allele frequencies to be erroneously reported for locus 9. This can be corrected simply by shifting the numbers upwards by one line, so that the frequencies listed for allele 2 in these populations are in fact those for allele 1, those for allele 3 are those for allele 2, and those for allele 4 are those for allele 3. The  $F_{ST}$  values plotted in Figure 3B are unaltered, but in reference to Figure 3A, the previously calculated  $F_{ST}$  value of 0.625 for this locus becomes 0.418. This change alters none of our basic conclusions.