Corrigendum

In the paper by J. Karotam, A. C. Delves and J. Oakeshott (*Genetica* 88: 11-28) entitled 'Conservation and change in structural and 5' flanking sequences of esterase 6 in sibling *Drosophila* species' the authors wish to point out several errors in Figure 2. A corrected version of this figure is presented overleaf.

Fig. 2. Comparison of the inferred amino acid sequences of EST6 in the three species, with the complete *D. melanogaster* (mel) sequence as a reference and differences in *D. simulans* (sim) and *D. mauritiana* (mau) shown below. Gaps inserted in any of the three sequences to improve the alignment are represented by dots. Three possible catalytic residues and four N-linked glycosylation sites (*) are indicated by arrows. The signal peptide and the three disulfide bridges (C-C) are shown above the sequence. Sequences are numbered with the first residue in the mature peptide as +1.