Supplementary Figure 2. Further duplication of dcr-1 in Acyrthosiphon pisum LSR1. The maximum likelihood reconstruction from a 615 amino acid alignment (model JTT+I+F) shows that the duplication arose from the dcr-1b copy. An asterisk (*) marks the suggested moment of the duplication of dcr-1a and dcr-1b and a square (■) marks the suggested moment of the duplication of dcr-1b and dcr-1c. The sequence of dcr-1c of A. pisum LSR1 was not included in the phylogeny of Supplementary Figure 1 due to total lack of homologous positions.