Supplementary Figure 10. Amino acid sequence alignments of portions of GTPase activating protein-homologous sequences from *Reticulitermes* (present study), human (AAC35881), and *Drosophila* (NP_609544). Shaded residues are identical to the *Reticulitermes* sequence, and dashes (-) indicate gaps in the alignments. In the alignment, residues shown as “X” are undetermined.