Figure S3. Alignment of Classic OBPs. Clustal Omega alignment of the 32 Classic OBPs identified in the *Ae. albopictus* transcriptome: sub-families or clusters as identified in Manoharan et al. (2013) are indicated on the left side. Conserved Cysteins are in bold and highlighted in light blue with the numbering reported below. Selected signatures typical of each cluster/subfamily are shown with distinctive colors.