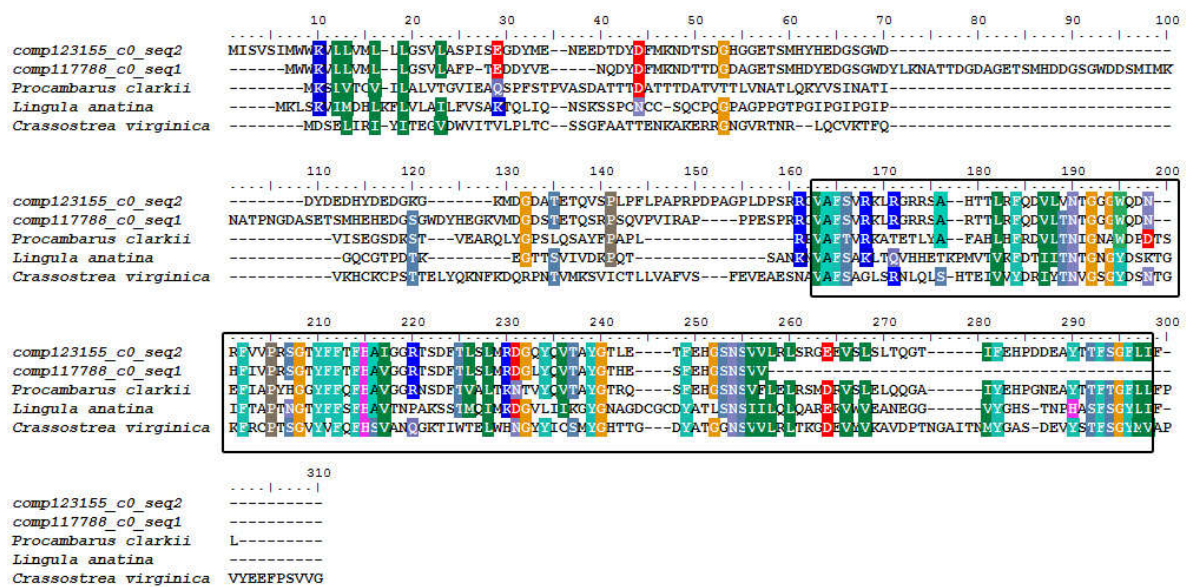




Additional file 8: Figure S5 Deduced amino acid sequence alignment of comp123155_c0_seq2, comp117788_c0_seq1 with complement component 1 q (C1q) from *Procamburus clarkia* (ASC55672), *Lingula anatine* (XP_013406649), and *Crassostrea virginica* (XP_022323368). Conserved amino acid residues were in colors. The highly conserved C1q domain was highlighted in a black box.