



Figure S2. Maximum likelihood phylogenies of gene families containing (A) M36, (B) S41, and (C) Asp Pfam domains. Each tip represents a single gene copy and each source species is denoted by color (*Bd.*: blue, *Hp.*: orange, *S. punctatus*: green, *A. macrogynus*: red). Bootstrap values over 80% indicated with asterisks at internal nodes.