

Figure S2. Multiple alignment of *Acanthamoeba* full-length mannose-binding proteins (MBP1). The DUF 4114 region, the CXCXC motifs, the transmembrane domain (TMD) and the NPLF motif are highlighted by colored rectangles on the consensus sequence. Note that some sequences present either additional motifs (*A. lenticulata* T5, *A. triangularis* T4F) or extended gaps (*A. culbertsoni* T10, *Acanthamoeba* sp. T22) which seem to be specific to each line.







