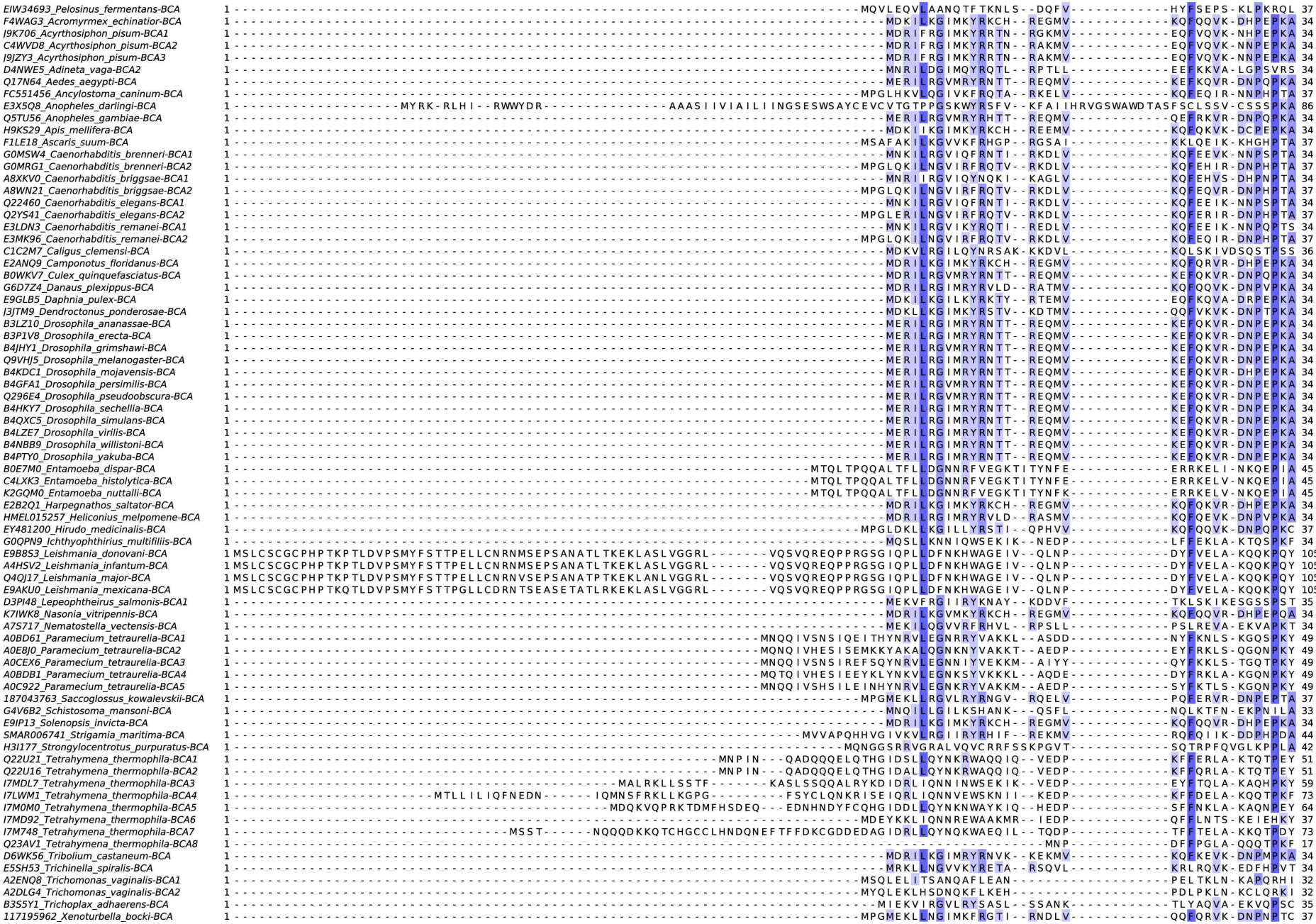
Supplementary Figure 1. Multiple sequence alignment of all 75 β-CAs in metazoan and protozoan species with β-CA of *Pelosinus fermentans* (a bacterial out group). β-CAs contain two highly conserved active site motifs, CxDxR as well as HxxC (C=Cysteine, D=Aspartic acid, R=Arginine, H=Histidine, C=Cysteine) which are indicated by arrows. Alignment was visualized in Jalview [53].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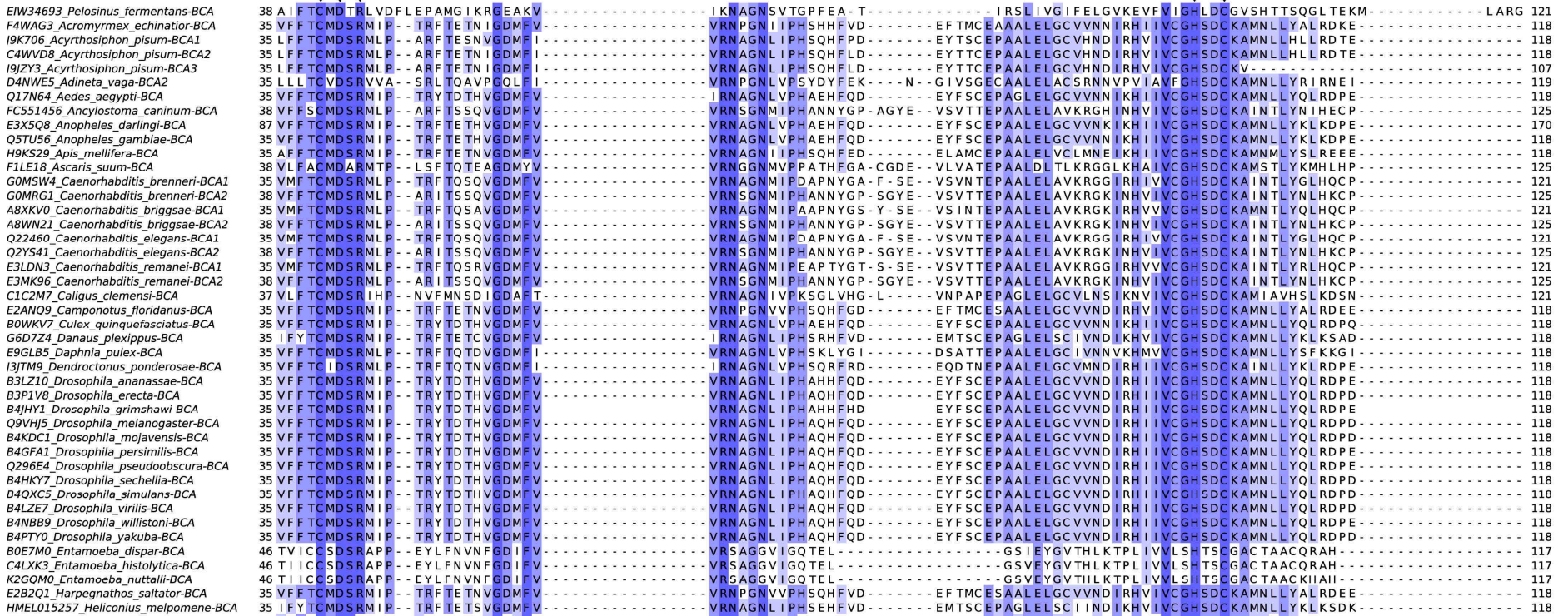
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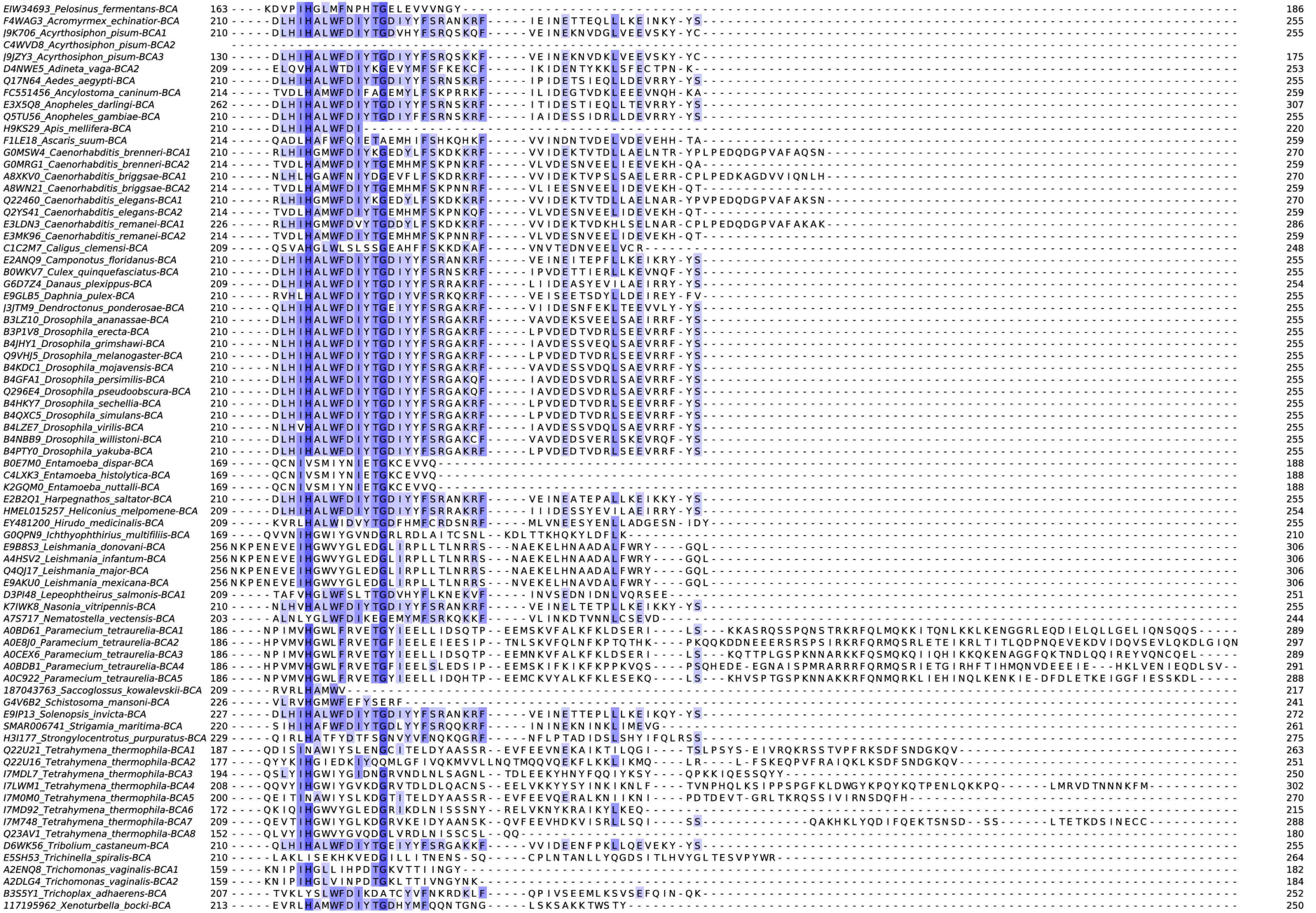
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