Supplementary Figure 10 Phylogenetic tree established using the 107 marker genes for all genomes of the genus *Dipylonosoma* and the 10 genomes of the genus *Diplosoma* as outgroup. Genomes are labelled according to their NCBI accession number. Genomes from this study are labelled with their MAG name. Genomes labelled with a star (*) are the representative genomes of the species clusters. Genomes labelled with a dagger (†) are the type strains. Genomes labelled with a double dagger (‡) are the neotype strains. Genomes labelled with a section sign (§) are the reference genomes. Genomes labelled with a pilcrow (¶) are the genomes used for calculating the ANI between species clusters. Genomes labelled with a hash (#) are the genomes used for calculating the ANI between species clusters. Genomes labelled with a double vertical bar (‖) are the genomes used for calculating the ANI between species clusters. 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