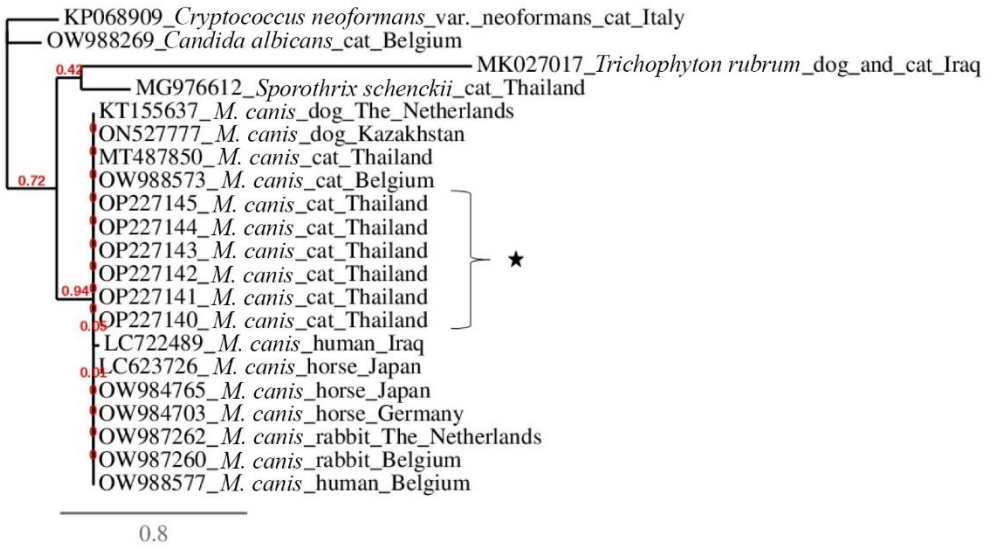

Figure S2: Phylogenetic tree analysis of *Microsporum canis* based on nucleotide sequences from a 720 bp fragment of ITS1-5.8S-ITS2 using the neighbor-joining method. Sequences from this study are marked with stars