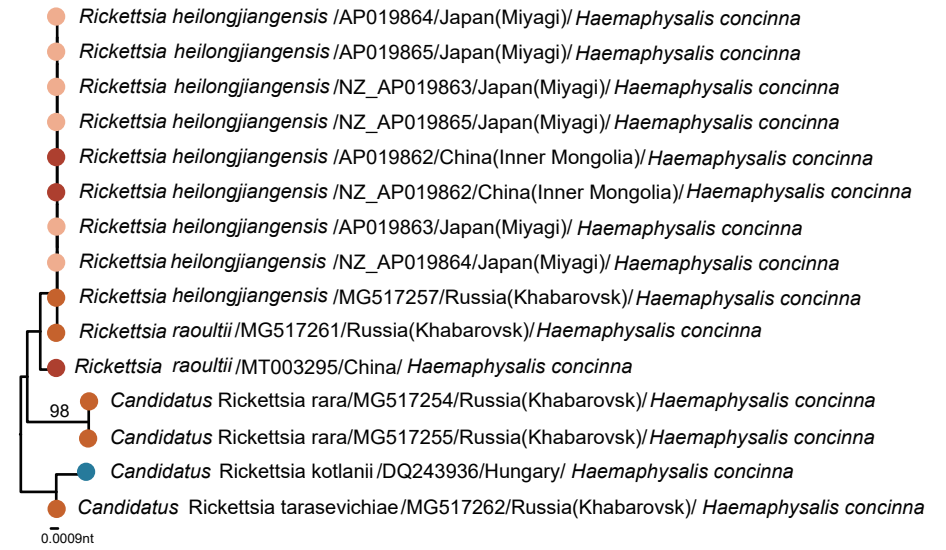


Figure S3: Phylogenetic analysis of *Haemaphysalis concinna*-associated microbes

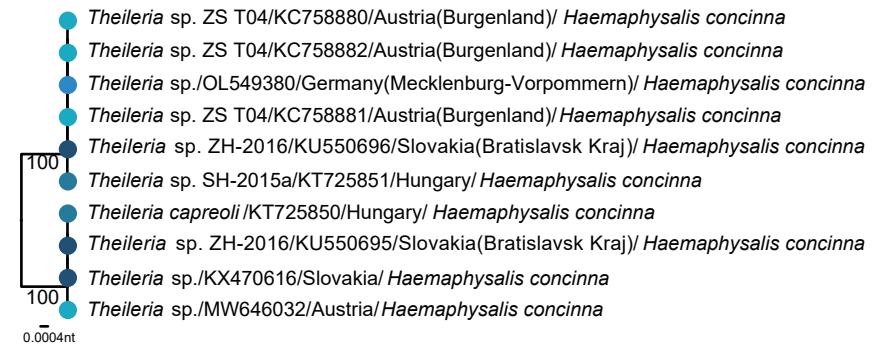
Babesia(18S rRNA)



Rickettsia(16S rRNA)



Theileria(18S rRNA)



Country of sequence

