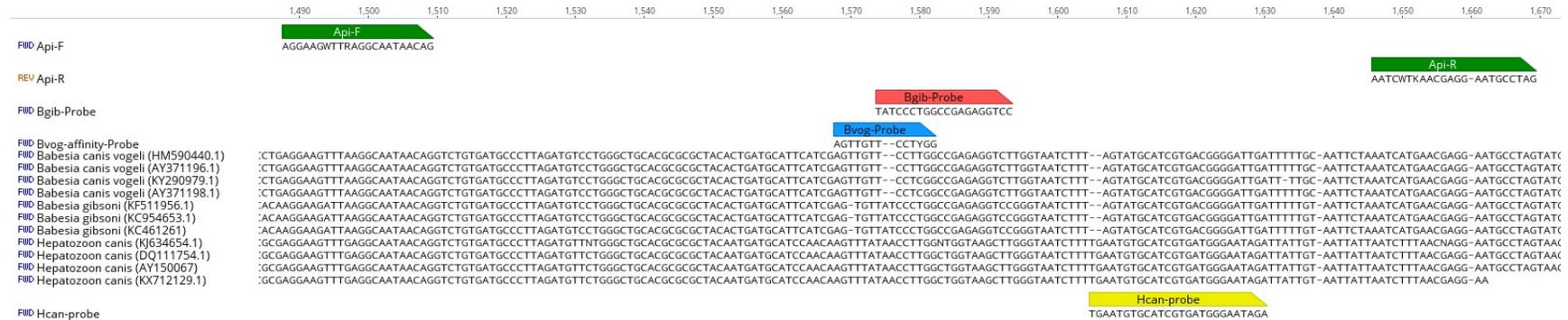
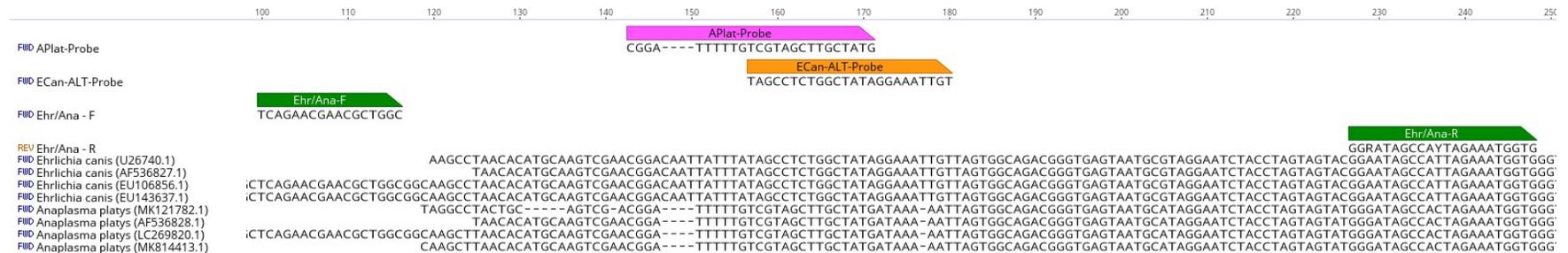


Supplementary file 2. Sequence alignments for qPCR quadruplex designs.

Alignment of *B. vogeli*, *B. gibsoni* and *H. canis* 18S rRNA sequences with relevant primer pair (green) and probes (red, blue, yellow). All alignments made in Geneious Prime v. 2021.0.3 (Biomatters Ltd.).



Alignment of *A. platys* and *E. canis* 16S rRNA sequences with relevant primer pair (green) and probes (pink, orange).



Alignment of canine infecting and non-canine infecting haemotropic *Mycoplasma* species' 16S rRNA sequences with relevant primer pair (green) and probe (purple).

