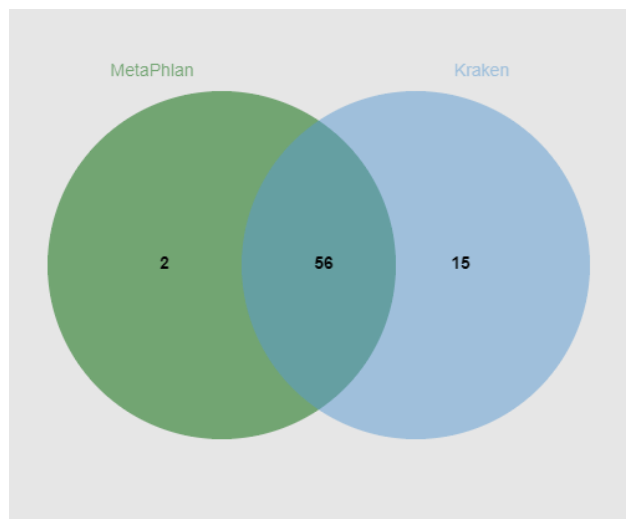
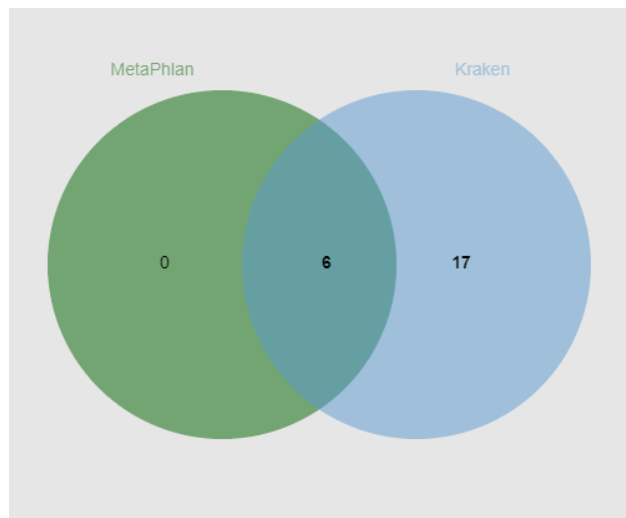


Figure S1: Venn diagram representing the overlap between samples identified as having > 1000 reads associated with either A) *Wolbachia* or B) *Spiroplasma*. MetaPhlan results are represented in Green and Kraken in Blue.

A)



B)



tree scale: 0.1

bootstrap

- 0
- 25
- 50
- 75
- 100

tree scale: 0.1

Time scale: 0.1

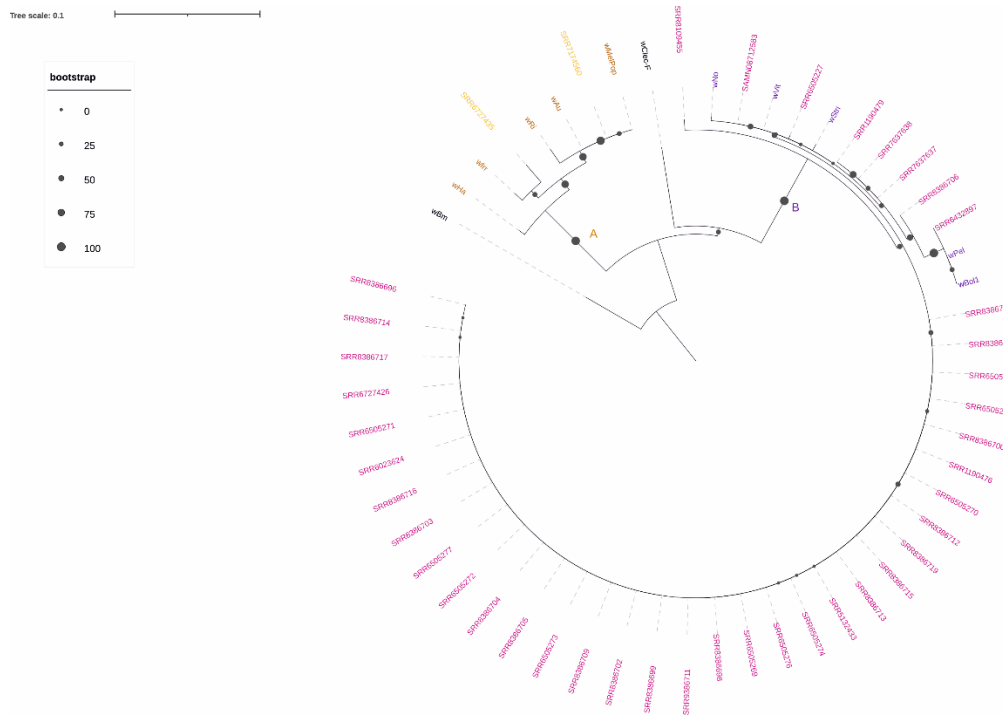
bootstrap

- 0
- 25
- 50
- 75
- 100

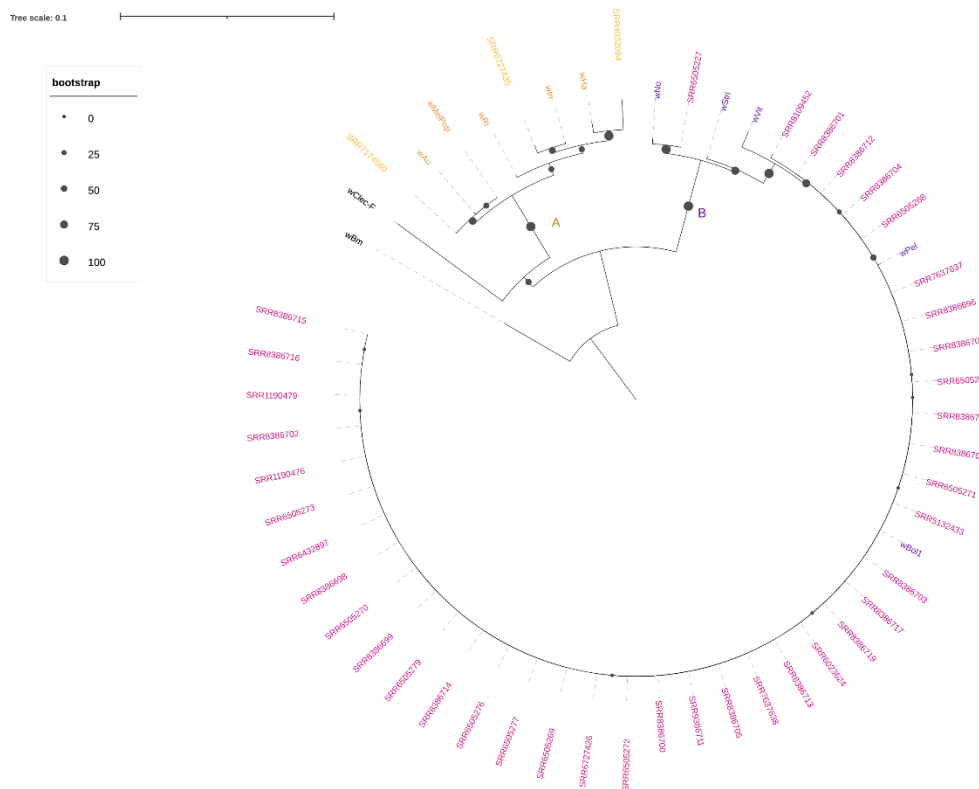
Accession numbers and gene names (clockwise from top):

- SR072745
- SR072746
- SR072747
- SR072748
- SR072749
- SR072750
- SR072751
- SR072752
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- SR072800
- SR072801
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- SR072804
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- SR072996
- SR072997
- SR072998
- SR072999
- SR073000

Tree scale: 0.1



FigureS6: Phylogenetic relationships between Wolbachia positive SRAs and reference sequences for the MLST gene hcpa



FigureS7: Phylogenetic relationships between Wolbachia positive SRAs and reference sequences for the wsp gene

