

**Table S9** The GenBank accession numbers of the ten *Bartonella* partial 16S-23S ITS sequence types detected in fleas only in this study. Sequence types are grouped into *Bartonella* species groups based on their closest match to existing named *Bartonella* species within GenBank.

<i>Bartonella</i> species	pITS variant	BLAST accession number
<i>B. grahamii</i>	grahamii-6	KU589247
	grahamii-7	KU589248
	grahamii-8	KU589249
<i>B. taylorii</i>	taylorii-11	KU589250
	taylorii-12	KU589251
<i>B. birtlesii</i>	birtlesii-8	KU589252
	birtlesii-9	KU589253
<i>B. doshiae</i>	doshiae-2	KU589254
Unknown	unknown-1	KU589255
<i>B. tribocorum</i>	tribocorum-1	KU589256