

Table S3. GenBank accession numbers of *Bartonella* spp. 16S–23S rRNA ITS, *Babesia microti* and *Hepatozoon* sp. 18S rRNA gene sequences identified in fleas collected from rodents

Name of the isolate	Source	bp	GenBank accession number	
			16S–23S rRNA ITS	18S rRNA
Uncultured <i>Bartonella</i> sp. clone FU184	<i>Ctenophthalmus agyrtes</i> from <i>Myodes glareolus</i>	460	MK239960	
Uncultured <i>Bartonella</i> sp. clone BA37	<i>Ctenophthalmus solutus</i> from <i>Apodemus flavicollis</i>	497	MK239961	
<i>Hepatozoon</i> sp. SK1 isolate F73	<i>Ctenophthalmus agyrtes</i> from <i>Myodes glareolus</i>	502		MH784529
<i>Hepatozoon</i> sp. SK1 isolate F103	<i>Ctenophthalmus agyrtes</i> from <i>Apodemus flavicollis</i>	502		MH784530
<i>Hepatozoon</i> sp. SK1 isolate F114	<i>Ctenophthalmus agyrtes</i> from <i>Apodemus flavicollis</i>	502		MH784531
<i>Hepatozoon</i> sp. SK1 isolate F185	<i>Ctenophthalmus agyrtes</i> from <i>Myodes glareolus</i>	502		MH784532
<i>Babesia microti</i> isolate F85	<i>Ctenophthalmus agyrtes</i> from <i>Myodes glareolus</i>	474		MH784533
<i>Babesia microti</i> isolate F92	<i>Ctenophthalmus congener</i> from <i>Myodes glareolus</i>	474		MH784534
<i>Babesia microti</i> isolate F110	<i>Palaeopsylla similis</i> from <i>Apodemus flavicollis</i>	474		MH784535