

<i>Theileria annulata</i> str. Ankara	<i>Theileria equi</i> str. WA	<i>Theileria orientalis</i> str. Shintoku	<i>Theileria orientalis</i> str. Robertson (Ikeda)	<i>Theileria orientalis</i> str. Goon nure (buffeli)	<i>Theileria orientalis</i> str. Fish creek (chitose)	
79.56%	70.44%	69.93%	70.02%	69.88%	69.82%	<i>Theileria parva</i> str. Muguga
	70.42%	69.97%	70.16%	70.00%	69.95%	<i>Theileria annulata</i> str. Ankara
		70.27%	70.06%	70.70%	70.58%	<i>Theileria equi</i> str. WA
			99.13%	81.69%	81.94%	<i>Theileria orientalis</i> str. Shintoku
				81.67%	81.96%	<i>Theileria orientalis</i> str. Robertson (Ikeda)
					85.35%	<i>Theileria orientalis</i> str. Goon nure (buffeli)

Figure S1: Pairwise average nucleotide identities of *Theileria* genomes represented as percentages inferred from pyANI.