

Supplementary Data

SUPPLEMENTARY TABLE S1. PRIMERS AND T_AQMAN PROBES USED FOR PATHOGEN DETECTION AND PHYLOGENETIC ANALYSIS

Pathogen	Target		Primers/probe	Product size (bp)	C/ <i>hrn</i> (nM)	Conditions	References
<i>Anaplasma phagocytophilum</i>	<i>msp2/p44</i>	p44-F: ATG GAA GGT AGT GTT GGT TAT GGT ATT p44-R: TTG GTC TTG AAG CGC TCG TA p44-probe: HEX-TGG TGC CAG GGT GCT TGA GAT TG-BHQ1		77	300 300 200	3 min 95°C, (10 s 95°C + 45 s 60°C) × 40 cycles	Graham et al. (2018)
	<i>msp4</i>	msp4-F: TAT ATC CAA CTT CCA CTC msp4-R: CAT TCA AGT TCG CTA AGA GTT TAC msp4-probe: HEX-CTC CGC CAA TAG CAT AGC CAG TTC-BHQ1		93	300 300 200		
<i>Borrelia</i> spp.	<i>16S rDNA</i>	16S-F: AGC YTT TAA AGC TTC GCT TGT AG 16S-R: GCC TCC CGT AGG AGT CTG G 16S-probe: FAM-CCG GCG TGA GAG GGT GAW CGG-BHQ1		148	600 600 200		
	<i>fliD</i>	fliD-F: TGG TGA CAG AGT GTA TAA TGG AA fliD-R: ACT CCT CCG GAA GCC ACA A fliD-probe: FAM-TGC TAA AAT GCT AGG AGA TTG TCT GTC GCC-BHQ1		78	400 400 200		
<i>Borrelia</i> spp.	<i>16S-18S rRNA</i>	IGS-F: CCTGAGGTGGATGTTCAACTC IGS-R: GCAAGGCCGAGGTCAAGG					Hojgaard et al. (2014)
	<i>16S-23S</i>	Round 1: 80: GTA TGT TTA GTG AGG GGG GTG 81: GGA TCA TAG CTC AGG TGG TTA G Round 2: 82: AGG GGG GTG AAG TCG TAA CAA G 83: GTC TGA TAA ACC TGA GGT CGG A	Bbss: 1030 Bbss: 988	500 500 500	Rd 1: 2 min 95°C, (94°C 30 s + 63°C 30 s) × 34 cycles; Rd 2: 2 min 95°C (94°C 30 s + 63°C 30 s) × 39 cycles		Bunikis et al. (2004)
<i>Bartonella</i> spp.	<i>tmRNA</i> (<i>ssrA</i>)	tmRNA-F: GCTATGGTAATAAATGGACAATGAAATAA tmRNA-R: GCTCTGTTGCCAGGTG tmRNA-probe: FAM-ACCCCGCTTAACCTGGCACG-BHQ1		253	1000 1000 200	2 min 95°C, (95°C 15 s + 60°C 60 s) × 45 cycles	Diaz et al. (2012)
<i>Brucella</i> spp.	<i>IS711</i>	IS711-F: GCTTGAAAGCTTGGGACAGT IS711-R: GGCCCTACCGCTGGAAT IS711-probe: HEX-AAGCCAACACCCGGCCATTATGGTBHQ1		63	1000 1000 200		Hinic et al. (2008)

(continued)

SUPPLEMENTARY TABLE S1. (CONTINUED)

Pathogen	Target	Primers/probe	Product size (bp)	C/ <i>xn</i> (nM)	Conditions	References
<i>Yersinia</i> spp.	<i>pal</i>	pal-F: CGC AAA TAA TGA CCA ATC TGG pal-R: CGT GGC CTT CAA CAA CAA C pal-probe: Quasar 670 - CGG TTC TGA CTT CGC TCA AAT GCT GG-BHQ2	1000 1000 200			Bai et al. (2017)
<i>Bartonella</i> spp.	ITS	325f: CTTTCAGATGATGATCCCCAAGGCCCTCTGGCG 1100r: GAACCCGACGACCCTGTGCAAAAGC 896f: GGC TAA TGA AGC AGT GAT AA 1233r: GCG ACG GTA TAC CCA TAG C Round 1: 443f: GCTATGTTGCAATTCTATCA 1210r: GATCYTCAATCATTTCTTCCA Round 2: 781f: GGGGACCAAGCTCATGGTGG 1137r: AATGCAAAAAGAACAGTAACACA	var	400 400 400	3 min 95°C, (95°C 30 s + 66°C 30 s + 72°C 30 s)×55 cycles;	Diniz et al. (2007)
<i>Bartonella</i> spp.	<i>gltA</i>	770	400	Rd 1: 2 min 95°C, (95°C 30 s + 48°C 30 s + 72°C 2 min)×40 cycles;	Birtles and Raoult (1996), Norman et al. (1995)	
<i>Yersinia pestis</i>	<i>pla</i>	Ypl: ATCTTACTTTCCGTGAGAAAG Yp2: CTGGATGTTGAGCTTCCTA	480	400	Rd 2: 2 min 95°C, (95°C 30 s + 55°C 30 s + 72°C 2 min)×40 cycles).	Hinnebusch and Schwan (1993)
<i>Rickettsia</i> spp.	<i>gltA</i>	CS-5: GAGAGAAAATTATATCCAAATGTTGAT CS-6: AGGGTCTTCGTGCATTCTT CS-5/6-FAM-CATTGTGCCATCCAGCTACGGT-BHQ1 Round 1: 877f: GGG GGC CTG CTC ACG GCG G 1258r: ATT GC AAA AAG TAC AGT GAA CA Round 2: 896f: 5'- GGC TAA TGA AGC AGT GAT AA-3' 1233r: 5'- GCG ACG GTA TAC CCA TAG C-3'	147 381 338	225 225 400 400	5 min 95°C, (1 min 95°C + 1 min 72°C)×35 cycles, 10 min 72°C, 4°C	Labruna et al. (2004)
<i>Rickettsia</i> spp.	<i>gltA</i>				Rd 1: 2 min 95°C, (95°C 20 s + 48°C 30 s + 60°C 2 min)×35 cycles);	Lee et al. (2014), Regnery et al. (1991)
					Rd 2: 10 min 95°C, (94°C 30 s + 55°C 30 s + 72°C 1 min)×30 cycles).	

Supplementary References

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