



**Figure S1.** Improvement of detection signals by pre-amplification. Test of primer/probe set sensitivity for a range of dilutions of positive controls by TaqMan real-time PCR using LightCycler 480, before and after pre-amplification. Results of the sensitivity test of the *Leishmania infantum* design using a *Leishmania infantum* culture, before (a) and after (c) pre-amplification; Results of the sensitivity test of the *Rickettsia* spp. design using *Rickettsia conorii*-positive controls (extracted from an infected *Rhipicephalus sanguineus* tick), before (b) and after (d) pre-amplification.

**Table S1.** List of the positive control samples used for the Biomark system development.

Positive control	Sample type	Source
<i>Borrelia burgdorferi</i> s.t.	Culture (Strain B31)	National Reference Center of Borrelia (Strasbourg, France)
<i>Borrelia garinii</i>	Culture (Strain N11)	Lise Gern, University of Neuchâtel (Neuchâtel, Switzerland)
<i>Borrelia afzelii</i>	Culture (Strain NE632)	Lise Gern, University of Neuchâtel (Neuchâtel, Switzerland)
<i>Borrelia lusitaniae</i>	Culture (Strain Poti-B1)	Lise Gern, University of Neuchâtel (Neuchâtel, Switzerland)
<i>Borrelia lonestari</i>	Infected <i>Amblyomma americanum</i> (Tick collected in USA)	CDC (Atlanta, USA)
<i>Borrelia bissettii</i>	Plasmid, rpoB gene & pBluescriptII SK+	GeneCust, September 2012 (Paris, France)
<i>Borrelia anserina</i>	Plasmid, fla gene & pBluescriptII SK+	GeneCust, December 2015 (Paris, France)
<i>Borrelia parkeri</i>	Culture	Cecilia Hizo-Teufel, National Reference Center of Borrelia (Germany)
<i>Borrelia recurrentis</i>	Culture	ANSES (Maisons-Alfort, France)
<i>Anaplasma marginale</i>	Experimentally infected cow blood	Isabel Garcia Fernandez de Mera (Spain)
<i>Anaplasma phagocytophilum</i>	Infected <i>Ixodes</i> spp. (Tick collected in USA)	CDC (Atlanta, USA)
<i>Anaplasma platys</i>	Infected dog blood	H.J. Bouloy, ANSES (Maisons-Alfort, France)
<i>Anaplasma ovis</i>	Plasmid, msp4 gene & pBluescriptII SK+	GeneCust, September 2012 (Paris, France)
<i>Anaplasma bovis</i>	Plasmid, groEL gene & pBluescriptII SK+	GeneCust, January 2016 (Paris, France)
<i>Ehrlichia ewingii</i>	Infected <i>Amblyomma americanum</i> (Tick collected in USA)	CDC (Atlanta, USA)
<i>Ehrlichia chaffeensis</i>	Infected <i>Amblyomma americanum</i> (Tick collected in USA)	CDC (Atlanta, USA)
<i>Ehrlichia ruminantium</i>	Culture	CIRAD (Guadeloupe, France)
Panola Mountain <i>Ehrlichia</i>	Infected <i>Amblyomma americanum</i> (Tick collected in USA)	CDC (Atlanta, USA)
<i>Ehrlichia canis</i>	Culture	ANSES (Maisons-Alfort, France)
<i>Neoehrlichia mikurensis</i>	Infected rodent blood	Lars Raberg, University of Lund (Sweden)
<i>Rickettsia conorii</i>	Infected <i>Rhipicephalus sanguineus</i> (Tick collected in USA)	CDC (Atlanta, USA)
<i>Rickettsia slovaca</i>	Culture	National Reference Center of Rickettsia (Marseille, France)
<i>Rickettsia massiliæ</i>	Culture	National Reference Center of Rickettsia (Marseille, France)
<i>Rickettsia rickettsii</i>	Plasmid, ITS & pBluescriptII SK+	GeneCust, Janvier 2016 (Paris, France)
<i>Rickettsia africae</i>	Culture	National Reference Center of Rickettsia (Marseille, France)
<i>Rickettsia typhi</i>	Culture	ANSES (Maisons-Alfort, France)
<i>Rickettsia prowazekii</i>	Plasmid, gltA gene & pBluescriptII SK+	GeneCust, December 2016 (Paris, France)
<i>Rickettsia felis</i>	Culture	National Reference Center of Rickettsia (Marseille, France)
<i>Bartonella henselae</i>	Culture	ANSES (Maisons-Alfort, France)
<i>Bartonella quintana</i>	Culture	H.J. Bouloy, ANSES (Maisons-Alfort, France)
<i>Bartonella bacilliformis</i>	Culture	ANSES (Maisons-Alfort, France)
<i>Bartonella vinsonii</i> subsp <i>berkhoffii</i>	Culture	Lynn Osikowicz, stacey Bartlett, CDC (Colorado, USA)
<i>Francisella tularensis</i>	Culture	Nora Madani, ANSES (Maisons-Alfort, France)
<i>Coxiella burnetii</i>	Culture	Elodie Rousset, ANSES (Maisons-Alfort, France)
<i>Aegyptianella pullorum</i>	Plasmid, groEL gene & pBluescriptII SK+	GeneCust, December 2015 (Paris, France)
<i>Babesia divergens</i>	Culture (done RFS)	Laurence Malandrin, Oniris (Nantes, France)
<i>Babesia microti</i>	Culture (isolate R1)	Emmanuel Comillot, University of Montpellier (Montpellier, France)
<i>Babesia bovis</i>	Plasmid, CCTeta gene & pBluescriptII SK+	GeneCust, Janvier 2016 (Paris, France)
<i>Babesia canis</i>	Infected dog blood	VetAgroSup (Lyon, France)
<i>Babesia vogeli</i>	Infected dog blood	VetAgroSup (Lyon, France)
<i>Babesia Rossi</i>	Infected dog blood	ANSES (Maisons-Alfort, France)
<i>Babesia caballi</i>	Plasmid, Rap1 gene & pBluescriptII SK+	GeneCust, September 2012 (Paris, France)
<i>Babesia bigemina</i>	Plasmid, 18S rRNA gene & pBluescriptII SK+	GeneCust, September 2012 (Paris, France)
<i>Babesia ovis</i>	Plasmid, 18S rRNA gene & pBluescriptII SK+	GeneCust, September 2012 (Paris, France)
<i>Babesia duncani</i>	Plasmid, ITS2 & pBluescriptII SK+	GeneCust, January 2016 (Paris, France)
<i>Babesia gibsoni</i>	Plasmid, rap gene & pBluescriptII SK+	GeneCust, December 2015 (Paris, France)
<i>Theileria parva</i>	Culture	Dirk Dobbelaera, University of Bern (Bern, Switzerland)
<i>Theileria annulata</i>	Culture	Dirk Dobbelaera, University of Bern (Bern, Switzerland)
<i>Theileria Lestoquari</i>	Culture	Dirk Dobbelaera, University of Bern (Bern, Switzerland)
<i>Cytauxzoon felis</i>	Plasmid, 18S rRNA gene & pBluescriptII SK+	GeneCust, December 2015 (Paris, France)
<i>Hepatozoon caris</i>	Infected dog blood	Domenico Otranto, Alessio Giannelli, University of Bari (Bari, Italy)
<i>Leishmania infantum</i> ( <i>chagasi</i> )	Culture	KASBARI Mohamed, ENVA (Maisons-Alfort, France)
<i>Leishmania Martiniquensis</i>	Culture	KASBARI Mohamed, ENVA (Maisons-Alfort, France)
<i>Rangelia vitalii</i>	Plasmid, 18S rRNA gene & pBluescriptII SK+	GeneCust, December 2015 (Paris, France)

<i>Borrelia theileri</i>	Plasmid, glpQ gene & pBluescriptII SK+	GeneCust, December 2017 (Paris, France)
<i>Theileria mutans</i>	Plasmid, ITS & pBluescriptII SK+	GeneCust, December 2017 (Paris, France)
<i>Theileria velifera</i>	Plasmid, 18S rRNA gene & pBluescriptII SK+	GeneCust, December 2017 (Paris, France)
<i>Theileria equi</i>	Plasmid, ema1 gene & pBluescriptII SK+	GeneCust, December 2017 (Paris, France)
<i>Hepatozoon americanum</i>	Plasmid, 18S rRNA gene & pBluescriptII SK+	GeneCust, December 2017 (Paris, France)
<i>Amblyomma variegatum</i>	Tick DNA (Tick collected in Guadeloupe)	Emmanuel Albina, CIRAD (Guadeloupe, France)
<i>Rhipicephalus microplus</i>	Tick DNA (Tick collected in Gualapagos)	ANSES (Maisons-Alfort, France)
<i>Rhipicephalus sanguineus</i> s.l.	Tick DNA (Tick collected in France)	ANSES (Maisons-Alfort, France)

**Table S2.** GPS coordinates of the tick collection sites and number of ticks collected. A total of 578 adult ticks collected from cattle from Guadeloupe and Martinique were used for the screening of tick-borne pathogens with the newly implemented BioMark™ real-time PCR system.

Location	Collection Sites	Cattle	Animal	Tick species	Tick number
GUADELOUPE					297
				<i>Amblyomma variegatum</i>	132
				<i>Rhipicephalus microplus</i>	165
	Sainte-Anne (N 16° 14'; W 61° 23')	Cattle 1	Animal 1	<i>Amblyomma variegatum</i>	19
		Cattle 2	Animal 2	<i>Amblyomma variegatum</i>	2
				<i>Rhipicephalus microplus</i>	4
			Animal 3	<i>Amblyomma variegatum</i>	7
				<i>Rhipicephalus microplus</i>	9
		Cattle 3	Animal 4	<i>Amblyomma variegatum</i>	1
	Petit Bourg (N 16° 10'; W 61° 36')	Cattle 4	Animal 5	<i>Amblyomma variegatum</i>	48
			Animal 6	<i>Amblyomma variegatum</i>	11
				<i>Rhipicephalus microplus</i>	9
		Cattle 5	Animal 7	<i>Amblyomma variegatum</i>	5
				<i>Rhipicephalus microplus</i>	4
			Animal 8	<i>Amblyomma variegatum</i>	1
		Cattle 6	Animal 9	<i>Amblyomma variegatum</i>	13
			Animal 10	<i>Amblyomma variegatum</i>	1
	Capesterre (N 16° 2'; W 61° 34')	Cattle 7	Animal 11	<i>Amblyomma variegatum</i>	4
			Animal 12	<i>Amblyomma variegatum</i>	9
				<i>Rhipicephalus microplus</i>	1
		Cattle 8	Animal 13	<i>Amblyomma variegatum</i>	2
				<i>Rhipicephalus microplus</i>	1
			Animal 14	<i>Amblyomma variegatum</i>	16
		Cattle 9	Animal 15	<i>Amblyomma variegatum</i>	6
				<i>Rhipicephalus microplus</i>	1
			Animal 16	<i>Amblyomma variegatum</i>	2
				<i>Rhipicephalus microplus</i>	5
				<i>Rhipicephalus microplus</i>	7
	Le Gosier (N 16° 13'; W 61° 28')	Cattle 10	Animal 17	<i>Amblyomma variegatum</i>	27
				<i>Rhipicephalus microplus</i>	11
			Animal 18	<i>Amblyomma variegatum</i>	3
		Cattle 11	Animal 19	<i>Amblyomma variegatum</i>	7
			Animal 20	<i>Amblyomma variegatum</i>	1
		Cattle 12	Animal 21	<i>Amblyomma variegatum</i>	3
				<i>Amblyomma variegatum</i>	2
	Pointe-Noire (N 16° 13'; W 61° 45')	Cattle 13	Animal 22	<i>Amblyomma variegatum</i>	13
			Animal 23	<i>Amblyomma variegatum</i>	2
			Animal 24	<i>Amblyomma variegatum</i>	2
			Animal 25	<i>Amblyomma variegatum</i>	4
		Cattle 14	Animal 26	<i>Amblyomma variegatum</i>	2
			Animal 27	<i>Amblyomma variegatum</i>	2
				<i>Amblyomma variegatum</i>	1
	Les Abymes (N 16° 16'; W 61° 31')	Cattle 15	Animal 28	<i>Amblyomma variegatum</i>	68
		Cattle 16	Animal 29	<i>Rhipicephalus microplus</i>	1
			Animal 30	<i>Rhipicephalus microplus</i>	10
		Cattle 17	Animal 31	<i>Rhipicephalus microplus</i>	6
			Animal 32	<i>Rhipicephalus microplus</i>	18
		Cattle 18	Animal 33	<i>Rhipicephalus microplus</i>	31
	Morne-À-l'Eau (N 16° 19'; W 61° 28')			<i>Rhipicephalus microplus</i>	2
					46

	Cattle 19	Animal 34	<i>Rhipicephalus microplus</i>	6
		Animal 35	<i>Rhipicephalus microplus</i>	2
	Cattle 20	Animal 36	<i>Rhipicephalus microplus</i>	19
		Animal 37	<i>Rhipicephalus microplus</i>	7
	Cattle 21	Animal 38	<i>Amblyomma variegatum</i>	1
			<i>Rhipicephalus microplus</i>	11
Bouillante (N 16° 7'; W 61° 46')				3
	Cattle 22	Animal 39	<i>Amblyomma variegatum</i>	2
MARTINIQUE		Animal 40	<i>Amblyomma variegatum</i>	1
			<i>Rhipicephalus microplus</i>	281
Rivière-Pilote (N 14° 28'; W 60° 54')				281
	Cattle 1	Animal 1	<i>Rhipicephalus microplus</i>	4
Le Diamant (N 14° 29'; W 61° 1')		Cattle 2	<i>Rhipicephalus microplus</i>	4
Le François (N 14° 36'; W 60° 53')		Cattle 3	<i>Rhipicephalus microplus</i>	1
Le Lamentin (N 14° 37'; W 60° 59')		Cattle 4	<i>Rhipicephalus microplus</i>	1
Le Lorrain (N 14° 49'; W 61° 2')		Cattle 5	<i>Rhipicephalus microplus</i>	52
Le Morne-Vert (N 14° 42'; W 61° 8')		Cattle 6	<i>Rhipicephalus microplus</i>	2
Le Robert (N 14° 40'; W 60° 56')		Cattle 7	<i>Rhipicephalus microplus</i>	2
Le Vaudin (N 14° 32'; W 60° 50')		Cattle 8	<i>Rhipicephalus microplus</i>	5
		Cattle 9	<i>Rhipicephalus microplus</i>	3
		Cattle 10	<i>Rhipicephalus microplus</i>	6
		Cattle 11	<i>Rhipicephalus microplus</i>	1
		Cattle 12	<i>Rhipicephalus microplus</i>	57
		Cattle 13	<i>Rhipicephalus microplus</i>	47
		Cattle 14	<i>Rhipicephalus microplus</i>	6
		Cattle 15	<i>Rhipicephalus microplus</i>	3
		Cattle 16	<i>Rhipicephalus microplus</i>	4
		Cattle 17	<i>Rhipicephalus microplus</i>	2
		Cattle 18	<i>Rhipicephalus microplus</i>	5
		Cattle 19	<i>Rhipicephalus microplus</i>	3
		Cattle 20	<i>Rhipicephalus microplus</i>	5
Rivière-Salée (N 14° 31'; W 60° 57')				3
Saint-Esprit (N 14° 33'; W 60° 55')		Cattle 21	<i>Rhipicephalus microplus</i>	3
Saint-Joseph (N 14° 40'; W 61° 2')		Cattle 22	<i>Rhipicephalus microplus</i>	28
Sainte-Anne (N 14° 26'; W 60° 50')		Cattle 23	<i>Rhipicephalus microplus</i>	28
Sainte-Luce (N 14° 29'; W 60° 56')		Cattle 24	<i>Rhipicephalus microplus</i>	13
Sainte-Marie (N 14° 46'; W 60° 59')		Cattle 25	<i>Rhipicephalus microplus</i>	7
		Cattle 26	<i>Rhipicephalus microplus</i>	6
Total		Cattle 27	<i>Rhipicephalus microplus</i>	29
		Cattle 28	<i>Rhipicephalus microplus</i>	3
		Cattle 29	<i>Rhipicephalus microplus</i>	33
				8
				8
				578