Supplementary Table 2. Oligonucleotides used for the molecular identification and/or characterization of the protist species investigated in this study.

Target organism	Locus	Oligonucleotide	Sequence (5′–3′)	Reference
Cryptosporidium spp.	*ssu* rRNA	CR-P1	CAGGGAGGTAGTGACAAGAA	40
		CR-P2	TCAGCCTTGCGACCATACTC	
		CR-P3	ATTGGAGGGCAAGTCTGGTG	
		CPB-DIAGR	TAAGGTGCTGAAGGAGTAAGG	
	gp60	AL3531	ATAGTCTCCGCTGTATTC	41
		AL3535	GGAAGGAACGATGTATCT	
		AL3532	TCCGCTGTATTCTCAGCC	
		AL3534	GCAGAGGAACCAGCATC	
	gp60	GP60 Ccanis_F1	ATACTCTGGTCTCCCGTTT	42
		GP60 Ccanis_R1	GTACTCGGAAGCGGTGTA	
		GP60 Ccanis_F2	AAGGCGCCTCACTCATT	
		GP60 Ccanis_R2	TCAGTTAGATATCACCCATTAA	
	gp60	GP60 CF_F1	TTTCCGTTATTGTTGCAGTTGCA	43
		GP60 CF_R1	ATCGGAATCCCACCATCGAAC	
		GP60 CF_F2	GGGCGTTCTGAAGGATGTAA	
		GP60 CF_R2	CGGTGGTCTCCTCAGTCTTC	
Giardia duodenalis	*ssu* rRNA	Probe	FAM–CCCGCGGCGGTCCCTGCTAG–BHQ1	44
		Gd-80F	GACGGCTCAGGACAACGGTT	
		Gd-127R	TTGCCAGCGGTGTCCG	
	gdh	GDHeF	TCAACGTYAAYCGYGGYTTCCGT	45
		GDHiF	CAGTACACCTCYGCTCTCGG	
		GDHiR	GTTRTCCTTGCACATCTCC	
	bg	G7_F	AAGCCCGACGACCTCACCCGCAGTGC	46
		G759_R	GAGGCCGCCCTGGATCTTCGAGACGAC	
		G99_F	GAACGAACGAGATCGAGGTCCG	
		G609_R	CTCGACGAGCTTCGTGTT	
	tpi	AL3543	AAATIATGCCTGCTCGTCG	47
		AL3546	CAAACCTTITCCGCAAACC	
		AL3544	CCCTTCATCGGIGGTAACTT	
		AL3545	GTGGCCACCACICCCGTGCC	

Enterocytozoon bieneusi	ITS	EBITS3	GGTCATAGGGATGAAGAG	48
		EBITS4	TTCGAGTTCTTTCGCGCTC	
		EBITS1	GCTCTGAATATCTATGGCT	
		EBITS2.4	ATCGCCGACGGATCCAAGTG	
Blastocystis spp.	*ssu* rRNA	BhRDr	GAGCTTTTTAACTGCAACAACG	49
		RD5	ATCTGGTTGATCCTGCCAGT	

bg: β-giardin; *gdh*: Glutamate dehydrogenase; *gp60*: 60-kDa glycoprotein; ITS: Internal transcribed spacer; *ssu* rRNA: Small subunit ribosomal RNA; *tpi*: Triose phosphate isomerase.