

**Table S2.** Oligonucleotides used for the molecular identification and/or characterization of *Cryptosporidium* spp. and *Giardia duodenalis* in the present study.

Target organism	Locus	Oligonucleotide	Sequence (5'–3')	Reference
<i>Cryptosporidium</i> spp.	<i>ssu</i> rRNA	CR-P1	CAGGGAGGTAGTGACAAGAA	[57]
		CR-P2	TCAGCCTTGCGACCATACTC	
		CR-P3	ATTGGAGGGCAAGTCTGGTG	
		CPB-DIAGR	TAAGGTGCTGAAGGAGTAAGG	
<i>Cryptosporidium parvum</i>	<i>gp60</i>	AL 3531	ATAGTCTCCGCTGTATTC	[58]
		AL 3535	GGAAGGAACGATGTATCT	
		AL 3532	TCCGCTGTATTCTCAGCC	
		AL 3534	GCAGAGGAACCAGCATC	
<i>Giardia duodenalis</i>	<i>ssu</i> rRNA	Probe	FAM–CCCGCGGCGGTCCCTGCTAG–BHQ1	[59]
		Gd-80F	GACGGCTCAGGACAACGGTT	
		Gd-127R	TTGCCAGCGGTGTCCG	
	<i>ssu</i> rRNA	Gia2029	AAGTGTGGTGCAGACGGA CT C	[60]
		Gia2150c	CTGCTGCCGTCCTTGGATGT	
		RH11	CATCCGGTCGATCCTGCC	
	<i>gdh</i>	RH4	AGTCGAACCTGATTCTCCGCCAGG	[61]
		GDHeF	TCAACGYAAYCGYGGYTTCCGT	
		GDHiF	CAGTACACCTCYGCTCTCGG	
	<i>bg</i>	GDHiR	GTTRTCCTTGACATCTCC	[62]
		G7_F	AAGCCCGACGACCTCACCCGCAGTGC	
		G759_R	GAGGCCGCCCTGGATCTTCGAGACGAC	
	<i>tpi</i>	G99_F	GAACGAACGAGATCGAGGTCCG	[63]
		G609_R	CTCGACGAGCTTCGTGTT	
		AL3543	AAATATGCCTGCTCGTCG	
	<i>tpi</i>	AL3546	CAAACCTTITCCGCAAACC	[64]
		AL3544	CCCTTCATCGGIGGTA ACT T	
		AL3545	GTGGCCACCACICCCGTGCC	

*bg*:  $\beta$ -giardin; *gdh*: Glutamate dehydrogenase; *ssu* rRNA: Small subunit ribosomal RNA; *tpi*: Triose phosphate isomerase.