Supplementary Table 1. Oligonucleotides used for the molecular identification and/or characterization of the intestinal protists investigated in the present study.

Target organism	Locus	Oligonucleotide	Sequence (5´–3´)	Reference
Giardia duodenalis	*ssu* rRNA	Probe	FAM–CCCGCGGCGGTCCCTGCTAG–BHQ1	[44]
		Gd-80F	GACGGCTCAGGACAACGGTT	
		Gd-127R	TTGCCAGCGGTGTCCG	
	ssu rRNA	Gia2029	AAGTGTGGTGCAGACGGACTC	[45]
		Gia2150c	CTGCTGCCGTCCTTGGATGT	
		RH11	CATCCGGTCGATCCTGCC	[46]
		RH4	AGTCGAACCCTGATTCTCCGCCAGG	
	gdh	GDHeF	TCAACGTYAAYCGYGGYTTCCGT	[47]
		GDHiF	CAGTACACCTCYGCTCTCGG	
		GDHiR	GTTRTCCTTGCACATCTCC	
	bg	G7_F	AAGCCCGACGACCTCACCCGCAGTGC	[48]
		G759_R	GAGGCCGCCCTGGATCTTCGAGACGAC	
		G99_F	GAACGAACGAGATCGAGGTCCG	[49]
		G609_R	CTCGACGAGCTTCGTGTT	
	tpi	AL3543	AAATIATGCCTGCTCGTCG	[50]
		AL3546	CAAACCTTITCCGCAAACC	

		AL3544	CCCTTCATCGGIGGTAACTT	
		AL3545	GTGGCCACCACICCCGTGCC	
Cryptosporidium spp.	*ssu* rRNA	CR-P1	CAGGGAGGTAGTGACAAGAA	[51]
		CR-P2	TCAGCCTTGCGACCATACTC	
		CR-P3	ATTGGAGGGCAAGTCTGGTG	
		CPB-DIAGR	TAAGGTGCTGAAGGAGTAAGG	
	gp60	AL-3531	ATAGTCTCCGCTGTATTC	[52]
		AL-3535	GGAAGGAACGATGTATCT	
		AL-3532	TCCGCTGTATTCTCAGCC	
		AL-3534	GCAGAGGAACCAGCATC	
Entamoeba histolytica	*ssu* rRNA	Probe	FAM–TCATTGAATGAATTGGCCATTT–MGB	[54]
Entamoeba dispar		Probe	VIC–TTACTTACATAAATTGGCCACTTTG–MGB	
Entamoeba histolytica/*dispar*		Ehd-239F	ATTGTCGTGGCATCCTAACTCA	
		Ehd-88R	GCGGACGGCTCATTATAACA	
Blastocystis sp.	*ssu* rRNA	BhRDr	GAGCTTTTTAACTGCAACAACG	[55]
		RD5	ATCTGGTTGATCCTGCCAGT	
	ssu rRNA	ILMN_Blast505_532F	TCGTCGGCAGCGTCAGATGTGTATAAGAGACAG GGAGGTAGTGACAATAAATC[a]	[56]
		ILMN_Blast998_1017R	GTCTCGTGGGCTCGGAGATGTGTATAAGAGACA GTGCTTTCGCACTTGTTCATC[a]	

[a]Primer used in next-generation amplicon sequencing. Adapter sequences are shown underlined.

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