

Supplemental Table 2: Helminth target regions, primer sequences, and probe sequences helminths for DNA amplification.

Parasite	Target region	Forward primer sequence (5' to 3')
		Reverse primer sequence (5' to 3')
		Probe sequence (5'FAM to 3')
<i>Acanthamoeba</i> species	18S rRNA	CCCAGATCGTTTACCGTGAA TAAATATTAATGCCCCCAACTATCC CTGCCACCGAATACATTAGCATGG
<i>Ancylostoma</i> species	ITS-2	GAATGACAGCAAACCTCGTTGTTG ATACTAGCCACTGCCGAAACGT ATCGTTTACCGACTTTAG
<i>Ascaris lumbricoides</i>	ITS-1	TGCACATAAGTACTATTTGCGCGTAT CCGCCGACTGCTATTACATCA GAGCCACATAGTAAATT
<i>Cryptosporidium</i> species	DNA-J like protein	AACTTCACGTGTGTTTGCCAAT CCAATCACAGAATCATCAGAATCG CATATGAAGTTATAGGGATACCAG

<i>Blastocystis</i> species.	16s rRNA	AGTAGTCATACGCTCGTCTCAAA TCTTCGTTACCCGTTACTGC CGTGTAATCTTACCATTAGAGGA
<i>Entamoeba histolytica</i>	18S rRNA	GTTTGTATTAGTACAAAATGGCCAAT TC TCGTGGCATCCTAACTCACTTAGA CAATGAATTGAGAAATGACA
<i>Giardia intestinalis</i>	16S rRNA	CATGCATGCCCCGCTCA AGCGGTGTCCGGCTAGC AGGACAACGGTTGCAC
<i>Necator americanus</i>	ITS-2	CTGTTTGTGGAACGGTACTTGC ATAACAGCGTGACATGTTGC CTGTACTACGCATTGTATAC
<i>Strongyloides stercoralis</i>	18s rRNA	GAATTCCAAGTAAACGTAAGTCATTA GC

		TGCCTCTGGATATTGCTCAGTTC ACACACCGGCCGTCGCTGC
<i>Toxocara canis</i>	ITS-2	GCGCCAATTTATGGAATGTGAT GAGCAAACGACAGCSATTTCTT CCATTACCACACCAGCATAGCTCAC CGA
<i>Toxocara cati</i>	ITS-2	ACGCGTACGTATGGAATGTGCT GAGCAAACGACAGCSATTTCTT TCTTTCGCAACGTGCATTCCGGTGA
<i>Trichuris trichiura</i>	ITS-1	TCCGAACGGCGGATCA CTCGAGTGTCACGTCGTCCTT TTGGCTCGTAGGTCGTT
<i>Internal Amplification Control (IAC)</i>	Synthetic sequence	CTAACCTTCGTGATGAGCAATCG GATCAGCTACGTGAGGTCCTAC TCGATGCACTCCAGTCCTCCT

*ITS = internal transcribed spacer; rRNA =
ribosomal RNA