



SUPPLEMENTAL FIGURE 1. DNA gels showing size matching of 19 samples to 120 base pairs from ladder and *Necator* plasmid standards. All samples have bands, but some are very light and difficult to visualize on these images.

SUPPLEMENTAL TABLE 1
Primer, probes, and target regions for all eight parasites. Sequence information for the eight parasites

Parasite	Forward primer sequence Reverse primer sequence probe sequence (FAM)	Target region	Gene ascension number
<i>Ascaris lumbricoides</i>	5'-TGCACATAAGTACTATTTGCGCGTAT-3' 5'-CCGCCGACTGCTATTACATCA-3'	ITS-1	AB571301.1
<i>Cryptosporidium</i> Species	5'-GAGCCACATAGTAAATT-3' 5'-AACTTCACGTGTGTTTGCCAAT-3' 5'-CCAATCACAGAATCATCAGAATCG-3' 5'-CATATGAAGTTATAGGGATACCAG-3'	DNAJ like protein	XM_625506.1
<i>Ancylostoma duodenale</i>	5'-GAATGACAGCAAACCTCGTTGTTG-3' 5'-ATACTAGCCACTGCCGAAACGT-3' 5'-ATCGTTTACCGACTTTAG-3'	ITS-2	EU344797.1
<i>Necator americanus</i>	5'-CTGTTTGTGGAACGGTACTTGC-3' 5'-ATAACAGCGTGACATGTTGC-3' 5'-CTGTACTACGCATTGTATAC-3'	ITS-2	AJ001599.1
<i>Strongyloides stercoralis</i>	5'-GAATTCGAAGTAAACGTAAGTCATTAGC-3' 5'-TGCCTCTGGATATTGCTCAGTTC-3' 5'-ACACACCGGCCGCTCGCTGC-3'	18S rRNA	AF279916.2
<i>Giardia lamblia</i>	5'-CATGCATGCCCGCTCA-3' 5'-AGCGGTGTCCGGCTAGC-3' 5'-AGGACAACGGTTGCAC-3'	16S rRNA	AJ293299.1
<i>Entamoeba histolytica</i>	5'-GTTTGTATTAGTACAAAATGGCCAATTC-3' 5'-TCGTGGCATCCTAACTCACTTAGA-3' 5'-CAATGAATTGAGAAATGACA-3'	18S rRNA	X75434.1