

Suggested PCR primers useful for parasitic helminths

Genetic marker	Target helminth group	Primer name	Primer sequence (5'-3')	Reference
18S rRNA	Platyhelminths	WormA	GCGAATGGCTCATTAATCAG	Waeschenbach & Littlewood (2017)
		WormB	CTTGTTACGACTTTTACTTCC	
	Nematodes	988F	CTCAAAGATTAAGCCATGC	Holterman et al. (2006)
		1912R	TTTACGGTCAGAACTAGGG	
		1813F	CTGCGTGAGAGGTGAAAT	
		2646R	GCTACCTTGTTACGACTTTT	
28S rRNA	Platyhelminths	digl2*	AAGCATATCACTAAGCGG	Tkach et al. (2001)
		LO	GCTATCCTGAG(AG)GAAACTTCG	
	Nematodes	D2A**	ACAAGTACCGTGAGGGAAAGTTG	De Ley et al. (2005)
		D3B	TCGGAAGGAACCAGCTACTA	
COI	Helminths	JB3	TTTTTTGGGCATCCTGAGGTTTAT	Bowles et al. (1992)
		JB4.5	TAAAGAAAGAACATAATGAAAATG	
ND1	Platyhelminthes	JB11	AGATTCGTAAGGGGCCTAATA	Bowles & McManus (1993)
		JB12	ACCACTAACTAATTCACTTTC	
12S rRNA	Platyhelminths	12STreF	GTGCCAGCADYYGCGGTTA	Chan et al. (2022)
		12STreR	AGCAGCAYATHGACCTG	
	Nematodes (Trichocephalida)	12S.C1.F	GTGCCAGCTAYCGCGGTTA	Chan et al. (2020)
		12S.C1.R	GRTGACGGGCRATATGTG	
	Nematodes (Ascaridida, Spirurida, Strongylida)	12S.C345.F	GTWCCAGAATAATCGGMTA	
		12S.C345.R	ATTGAYGGATGRTTGTGTRC	
16S rRNA	Platyhelminths	16SCesTreF	GTGYDAAGGTAGSATAAT	Chan et al. (2022)
		16SCesTreR	CCGGTYTYAACTCARCTCAT	
	Nematodes (Trichocephalida)	16S.C1.F	ACGAGAAGACCCTRGRAAYT	Chan et al. (2020)
		16S.C1.R	GRTYTAACTCAAATCACG	
	Nematodes (Ascaridida, Spirurida, Strongylida)	16S.C345.F	AAGATAAGTCTTYGGAARYT	
		16S.C345.R	GAAYTAACTAATATCAMG	

*Primers targeting the D1-D3 region of the 28S rRNA gene

**Primers targeting the D2-D3 region of the 28S rRNA gene

Note:

- The primers provided here are popular for targeting broad species range in helminths. PCR conditions can be found at the respective references.
- Primers for the ITS regions, *COII*, and *cytB* genes were not included as the primers depends on the target helminth species.
- The full-length mitochondrial sequences (*COI*, *COII*, *cytB*, *NAD1*, 12S, 16S) obtained for use in ABlapp were mined from the complete mitochondrial genomes available in GenBank database.
- The nuclear sequences (18S, 28S, ITS1, ITS2) obtained for use in ABlapp were mined from almost complete sequences available in GenBank database.

References:

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