

Supplementary Table 1. Matches of published primer sequences to sequences from *Mycoplasma* species isolated from small ruminants.

	Reference strain	% identity	GenBank accession	Reference
Forward primer sequence (in 16S rRNA gene)				
5'-ACACCATGGGAGCTGGTAAT				
A <u>A</u> ACCATGGGAGCTGGTAAT	<i>M. adleri</i> strain G145	95	AY770622.1	2
A <u>A</u> ACCATGGGAGCTGGTAAT	<i>M. agalactiae</i> strain PG2	95	CU179680.1	2
ACACCATGGGAGCTGGTAAT	<i>M. arginini</i> strain G230	100	NR_026035.1	2
ACACCATGGGAGCTGGTAAT	<i>M. auris</i> strain UIA	100	AY772217.1	2
A <u>A</u> ACCATGGGAGCTGGTAAT	<i>M. bovirhinalium</i> strain PG11	95	NR_044668.2	3
ACACCATGGGAGCTGGTAAT	<i>M. bovirhinalis</i> strain NBRC 14857	100	NR_113685.1	3
A <u>A</u> ACCATGGGAGCTGGTAAT	<i>M. bovis</i> strain PG45	95	CP002188.1	2
A <u>A</u> ACCATGGGAGCTGGTAAT	<i>M. californicum</i> strain ST-6	95	CP007521.1	2
ACACCATGGGAGCTGGTAAT	<i>M. canadense</i> strain 275c	100	NR_025988.1	5
AAACCATGGGAG <u>T</u> TGGTAAT	<i>M. conjunctivae</i> strain HRC/583	95	FM864216.2	2
A <u>A</u> ACCATGGGAGCTGGTAAT	<i>M. fermentans</i> strain PG18	95	AP009608.1	1
A <u>A</u> ACCATGGGAGCTGGTAAT	<i>M. gallinarum</i> strain ATCC 15319	95	JN935884.1	4
AAACCATGGGAG <u>T</u> TGGTAAT	<i>M. ovipneumoniae</i> strain Y-98	95	AY753216.1	2
Reverse primer sequence (in 23S rRNA gene)				
5'-CTTCATCGACTTTCAGACCCAAGGCAT				

CCTCATCGACTT <u>CC</u> CAGACCCAAGGCAT	<i>M. adleri</i> strain G145	92.6	AY770622.1	2
CTTCATCGACTT <u>CC</u> CAGACCCAAGGCAT	<i>M. agalactiae</i> strain PG2	96.3	CU179680.1	2
CTTCATCGACTT <u>CC</u> CAGACCCAAGGCAT	<i>M. arginini</i> strain G230	96.3	LC158837.1	2
CTTCATCGACTT <u>CC</u> CAGACCCAAGGCAT	<i>M. auris</i> strain UIA	96.3	AY772217.1	2
CTTCATCGACTT <u>CC</u> CAGACCCAAGGCAT	<i>M. bovinegenitalium</i> strain PG11	96.3	LC158838.1	3
CTTCATCGACTT <u>CC</u> CAGACCCAAGGCAT	<i>M. bovirhinis</i> strain NBRC 14857	96.3	LC158839.2	3
CTTCATCGACTT <u>CC</u> CAGACCCAAGGCAT	<i>M. bovis</i> strain PG45	96.3	CP002188.1	2
CTTCATCGACTT <u>CC</u> CAGACCCAAGGCAT	<i>M. californicum</i> strain ST-6	96.3	CP007521.1	2
CTTCATCGACTT <u>CC</u> CAGACCCAAGGCAT	<i>M. canadense</i> strain 275c	96.3	LC158840.1	5
CTTC <u>T</u> TCGACTTTTCAGACCCAAGGCAT	<i>M. conjunctivae</i> strain HRC/583	96.3	FM864216.2	2
CTTCATCGACTTTTCAGACCCAAGGCAT	<i>M. fermentans</i> strain PG18	100	AP009608.1	1
CTTCATCGACTT <u>CC</u> CAGACCCAAGGCAT	<i>M. gallinarum</i> strain ATCC 15319	96.3	JN935884.1	4
CTTCATCGACTTTTCAGACCCAAGGCAT	<i>M. ovipneumoniae</i> strain Y-98	100	AY753216.1	2

Nucleotide(s) differing from the published primer are highlighted in bold font and underlined.

References

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