

Diagnostic Plasmid Containing the Small Subunit Ribosomal RNA Gene (18S) from *Plasmodium malariae*

Catalog No. MRA-179

Product Description: The small subunit ribosomal RNA gene (18S rRNA gene; GenBank: [AF145336](#)) from *Plasmodium malariae* (*P. malariae*) was amplified from genomic DNA by nest 1 PCR primers and cloned into vector pCR2.1-TOPO (Invitrogen®). The resulting plasmid (clone 34) may be used in PCR assays for the diagnosis of mixed species malaria infections.

Lot^{1,2}: 2165

Manufacturing Date: MAR2017

TEST	SPECIFICATIONS	RESULTS
Next-Generation DNA Sequencing³ MRA-179 18S rRNA gene pCR2.1-TOPO (Invitrogen®)	~ 5000 base pairs ~ 1100 base pairs ~ 3900 base pairs	5045 base pairs (Figures 1 and 2) 1114 base pairs 3931 base pairs
Genotypic Analysis Sequencing of 18S ribosomal RNA gene (~ 1110 base pairs)	≥ 99% sequence identity to <i>P. malariae</i> 18S rRNA gene (GenBank: AF145336)	100% sequence identity to <i>P. malariae</i> 18S rRNA gene (GenBank: AF145336)
Concentration by PicoGreen® Measurement	Report results	0.5 µg in 50 µL per vial (17.8 µg/mL)
Amount	Report results	~ 500 ng per vial
OD₂₆₀/OD₂₈₀ Ratio	1.7 to 2.1	1.9

¹MRA-179, lot 2165 was extracted from a preparation of MR-MRA-179, lot 58068548.

²Extracted using a QIAprep® Spin Miniprep Kit (QIAGEN® 27104)

³Illumina® MiSeq® sequence was analyzed with CLC Genomics Workbench Version 7.5.

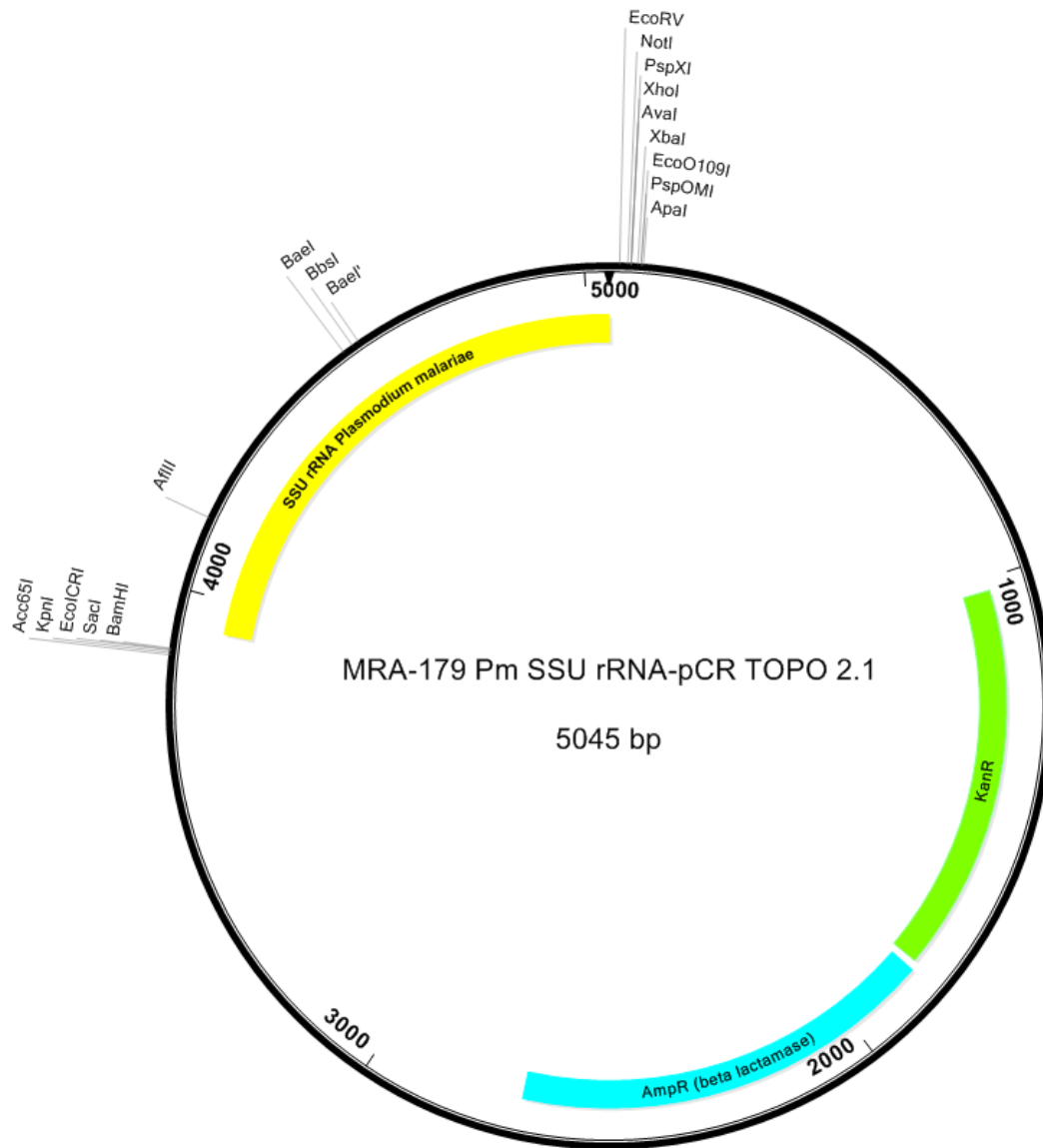
Figure 1: Complete Plasmid Sequence of MRA-179

>MRA-179 [lot_2165] complete plasmid sequence (red text is 18S rRNA gene sequence)

```
GTAAGCCCACTGCAAGTACCTGCTTTCTCTTTGCGCTTGCCTTTTCCCTTGTCCAGATAGCCAGTAGCTGACATTCATCCGGGGTCAGCACCGTT
TCTGCGGACTGGCTTTCTACGTGTTCCGCTTCCTTTAGCAGCCCTTGCGCCCTGAATTTTGTAAATTCGCGTTAAATTTTTGTAAATCAGCTCA
TTTTTTAACCAATAGGCCGAAATCGGCAAAATCCCTTATAAATCAAAGAATAGACCGAGATAGGGTTGAGTGTGTCCAGTTTGAACAAGAGTC
CACTATTAAGAACGTGGACTCCAACGTCAAAGGGCGAAAACCGTCTATCAGGGCGATGGCCACTACGTGAACCATCACCTAATCAAGTTTTTT
GGGTCGAGGTCGCCGTAAGCACTAAATCGGAACCTAAAGGGAGCCCCGATTTAGAGCTTGACGGGGAAAGCCGGCGAACGTGGCGAGAAAGGAA
GGGAAGAAAGCGAAAGGAGCGGGCGCTAGGGCGCTGGCAAGTGTAGCGGTACCGCTGCGCGTAACCACCACACCCGCCGCGCTTAATGCGCCGCTAC
AGGGCGCTCCATTCGCCATTCAGGCTGCGCAACTGTTGGGAAGGGCGATCGGTGCGGGCCCTTTCGCTATTACGCCAGCTGGCGAAAGGGGATGT
GCTGCAAGGCGATTAAGTTGGGTAACGCCAGGGTTTTCCAGTCACGACGTTGTAACACGACGGCCAGTGAATTGTAATACGACTCACTATAGGGCG
AATTGGGCCCTCTAGATGCATGCTCGAGCGGCCAGTGTGATGGATATCTGCAGAATTCGCCCTTTAAAATTTGTCAGTTAAAACGCTCGTAG
TTGAATTTCAAGGAATCAATATTTTTAAGTAATGCTTTGTATATTTATAACAAAGTTGACGTTAAGAATAAACGCCAAGCGTTATATTTTTCTGTT
ACATTTTGTTTTATTAATATATATATGCGTTCTTATATAAAAATGATTTCTTTTTAAAATTTCTTTTGTATAATTTTTATGCATGGGAATTTTGTTA
CTTTGAGTAAATAGAGTGTCAAAGCAAACAGTTAAAACAGTTTCTGTGTTGAATACTACAGCATGGAATAACAAAATGAACAAGTCAGAATTT
GTTCTTTTTTCTTATTTTGGCTTAGTTACGATTAATAGGAGTAGCTTGGGGCAATTTGTATTTCAGATGTCAGAGGTGAAATTTCTAGATTTTCTGG
AGACAAGCAACTGCGAAAGCATTTGCTTAAATACTTCCATTAATCAAAGAACGAAAGTTAAGGGAGTGAAGACGATCAGATACCGTCGTAATCTTAA
CCATAAATATGCCGACTAGGTGTTGGATGATAGAGTAAAAATAAAGAGACATTCATATATATGAGTGTTCCTTTAGATAGCTTCCTTCAGTAC
CTTATGAGAAATCAAAGTCTTTGGGTTCTGGGGCGAGTATTCGCGCAAGCGAGAAAGTTAAAAGAATTGACGGAAGGGCACCACCAGGCGTGGAGCT
TGCGGCTTAATTTGACTCAACACGGGAAACTCACTAGTTTAAAGACAGAGTAGGATGACAGATTAATAGCTCTTCTTGATTTCTTGGATGGTGA
TGCATGGCCGTTTTTAGTTCGTGAATATGATTTGTCTGGTTAATTCGATAACGAACGAGATCTTAACCTGCTAATTAGCGGTAATAACACTATATT
CTTAAGTGAATTAGAATATAGATAAATTTGTGCTAATTTTGATTAATAATAGAAATGTTTTTTTTGATAAAAACGTTCTTTCCCTTTTTTCTTA
ATTATGCATATTTATTTTTTTTCTTTTGCATAAGAATGATTTTGCCTAATTTGTAAGCTTCTTAGAGGAACGATGTGTCTAACACAAGGAAG
TTAAGGCAACAACAGAGGGCGAATTCAGCACACTGGCGCCGTTACTAGTGGATCCGAGCTCGGTACCAAGCTTGGCGTAATCATGGTCATAG
```

CTGTTTCTGTGTGAAATTGTTATCCGCTCACAATTCACACAACATACGAGCCGGAAGCATAAAGTGTAAGCCTGGGGTGCCTAATGAGTGAGCT
AACTCACATTAATTGCGTTGCGCTCACTGCCCGCTTTCAGTCGGGAAACCTGTCTGTCAGCTGCATTAATGAATCGGCCAACGCGCGGGGAGAGG
CGGTTTGGGTATTGGGCGCTCTTCCGCTTCCCTCGCTCACTGACTCGCTGCGCTCGGTCGTTCCGGCTGCGGCGAGCGGTATCAGTCACTCAAAGGCG
GTAATACGGTTATCCACAGAATCAGGGGATAACGCAGGAAAGAACATGTGAGCAAAAAGGCCAGCAAAAAGGCCAGGAACCGTAAAAAGGCCGCTTGC
TGGCGTTTTTCCATAGGCTCCGCCCCCTGACGAGCATCACAAAAATCGACGCTCAAGTCAGAGGTGGCGAAACCCGACAGACTATAAAGATACCA
GGCGTTTTCCCTGGAAGCTCCCTCGTGCCTCTCTGTTCCGACCCTGCCGCTTACCGGATACCTGTCCGCCTTTCTCCCTTCGGGAAGCGTGCGC
CTTTCTCATAGCTCACGCTGTAGGTATCTCAGTTCGGTGTAGGTCGTTCCGCTCCAAGCTGGGCTGTGTGCACGAACCCCGCTTACGCCGACCGCT
GCGCCTTATCCGGTAACTATCGTCTTGAAGTCCAACCCGGTAAGACACGACTTATCGCCACTGGCAGCAGCCACTGGTAAACAGGATTAGCAGAGCGAG
GTATGTAGGCGGTGTACAGAGTCTTGAAGTGGTGGCTAACTACGGCTACACTAGAAGAACAGTATTTGGTATCTGCGCTCTGTGAAGCCAGTT
ACCTTCGGA AAAAGAGTTGGTAGCTCTTGTATCCGGCAAAACAAACCACCGCTGGTAGCGGTGGTTTTTTTGTGTTGCAAGCAGCAGATTACGCGCAGAA
AAAAAGGATCTCAAGAAGATCCTTTGATCTTTTCTACGGGGTCTGACGCTCAGTGGAAACGAAAACACGTTAAGGGATTTGGTTCATGAGATTATC
AAAAAGGATCTTACCTAGATCCTTTTAAATTA AAAATGAAGTTTTTAAATCAATCTAAAAGTATATATGAGTAAACTTGGTCTGACAGTTACCAATGC
TTAATCAGTGAGGCACCTATCTCAGCGATCTGTCTATTTTCGTTTCATCCATAGTTGCCGACTCCCCGTCGTGTAGATAACTACGATACGGGAGGGCT
TACCATCTGGCCCCAGTCTGCAATGATACCGCGAGACCCACGCTCACCGGCTCCAGATTTATCAGCAATAAACCAGCCAGCCGGAAGGGCCGAGCG
CAGAAGTGGTCTGCAACTTTATCCGCTCCATCCAGTCTATTAATGTTGCCGGGAAGCTAGAGTAAGTAGTTCGCCAGTTAATAGTTTGCGCAAC
GTTGTTGCCATTGCTACAGGCATCGTGGTGTACGCTCGTCTGTTGGTATGGCTTCATTCAGCTCCGGTCCCAACGATCAAGGCGAGTTACATGAT
CCCCATGTTGTGCAAAAAGCGGTTAGCTCCTTCGGTCTCCGATCGTGTGAGAAAGTAAAGTTGGCCGAGTGTATCACTCATGGTTATGGCAGC
ACTGCATAATTCTCTTACTGTCTATGCCATCCGTAAGATGCTTTTCTGTGACTGGTGGTACTCAACCAAGTCACTTCTGAGAATAGTGTATGCGGCGA
CCGAGTTGCTCTTGCCCGGCGTCAATACGGGATAATACCGCGCCACATAGCAGAACTTAAAAGTGTCTCATCATTGGAAAACGTTCTTCGGGGCGAA
AACTCTCAAGGATCTTACCCTGTTGAGATCCAGTTCGATGTAACCCACTCGTGCACCCAACTGATCTTCAGCATCTTTACTTTTACCAGCGTTTC
TGGGTGAGCAAAAACAGGAAGCAAAAATGCCGCAAAAAGGGAATAAGGGCGACACGGAAATGTTGAATACTCATACTCTTCTTTTTCAATTCAGA
AGAACTCGTCAAGAAGGCGATAGAAGGCGATGCGCTGCGAATCGGGAGCGGCGATACCGTAAAGCACGAGGAAGCGGTCAGCCCATTCCGCCGCAAG
CTCTTCAGCAATATCACGGGTAGCCAACGCTATGTCTGATAGCGGTCCGCCACACCCAGCCGGCCACAGTCGATGAATCCAGAAAAGCGGCCATTT
TCCACCATGATATTCGGCAAGCAGGCATCGCCATGGGTACGACAGAGATCCTCGCCGTCGGGCATGCGCGCCTTGAGCCTGGCGAACAGTTTCGGCTG
GCGCGAGCCCCCTGATGCTCTTCGTCAGATCATCCTGATCGACAAGACCGGCTTCCATCCGAGTACGTGCTCGCTCGATGCGATGTTTCGCTTGGT
GTGCAATGGGCGAGGTAGCCGATCAAGCGTATGCAGCCGCGCATTCATCAGCCATGATGGATACTTTCTCGGCAGGAGCAAGGTGGGATGACAGG
AGATCCTGCCCGGCACTTCGCCAATAGCAGCCAGTCCCTTCCCGCTTCAGTGACAACGTCGAGCACAGCTGCGCAAGGAACGCCGCTCGTGGCCA
GCCACGATAGCCGCGCTGCCTCGTCTGCAATTCAGGGCACCGGACAGGTCCGCTTTCGCAAAAAGAACCGGGCGCCCTGCGCTGACAGCCG
GAACACGGCGGCATCAGAGCAGCCGATTGTCTGTTGTGCCAGTCA TAGCCGAATAGCCTCTCCACCAAGCGGGCCGGAGAACCTGCGTGAATCCA
TCTTGTTCATCATGCGAAACGATCCTCATCTGTCTTGTGATCAGATCTTGATCCCTGCGCCATCAGATCCTTGGCGGCAAGAAAGCCATCCAGT
TTACTTTGCAGGGCTTCCCAACCTTACCAGAGGGCGCCAGCTGGCAATTCGGGTTGCTGTCCATAAAAACCGCCAGTCTAGCTATCGCCA
T

Figure 2: Plasmid Map of MRA-179



Date: 16 AUG 2017

Signature:

BEI Resources Authentication

ATCC®, on behalf of BEI Resources, hereby represents and warrants that the material provided under this certificate has been subjected to the tests and procedures specified and that the results described, along with any other data provided in this certificate, are true and accurate to the best of ATCC®'s knowledge.

ATCC® is a trademark of the American Type Culture Collection.

You are authorized to use this product for research use only. It is not intended for human use.



BEI Resources

www.beiresources.org

E-mail: contact@beiresources.org

Tel: 800-359-7370

Fax: 703-365-2898