ID	DNA Type	Description	Nucleotide Base Change	Sequence (5' → 3')
MEP417	D-DNA	André <i>rpoB</i> MAS-PCR forward primer (301 base-pair amplicon)	N/A	TGGAGTACGTGCCCTCGTC
MEP419	D-DNA	André I491F MAS-PCR forward primer (70 base-pair amplicon)	N/A	GGCCCAACATCGGTCTG <u>A</u>
MEP418	D-DNA	André <i>rpoB</i> MAS-PCR reverse primer (301 or 70 base-pair amplicons)	N/A	GTGGCCACCGACACCATCT
MEP352	D-DNA	rpoB forward primer (66 base-pair amplicon)	N/A	GATGTGCCCGATCGAAACC
MEP353	D-DNA	rpoB reverse primer (66 base-pair amplicon)	N/A	GCGTACACCGACAGCGA
MEP477	D-DNA	Wild-type rifampicin-susceptible melt probe with 3' C3 spacer	N/A	GCCGATCAGACCGATGTTGGGCCCCTCAGG/3SpC3/
24AUG_TB_ rpoBf_I491_ TXR	L-DNA	Wild-type rifampicin-susceptible comparator forward strand with 5' Texas Red-X	N/A	Texas Red / GCCGATCAGACCGATGTTGGGCCCCTCAGG
24AUG_TB_ rpoBf_I491_ Rcomp	L-DNA	Wild-type rifampicin-susceptible comparator reverse complement strand with no modification	N/A	CCTGAGGGGCCCAACATCGGTCTGATCGGC
MEP491-2	D-DNA	Wild-type rifampicin-susceptible target	N/A	GATGTGCCCGATCGAAACCCCTGAGGGGCCCAACATCGGTCTGATCGGCTCGCTGTCGGTGTACGC GCGGGTCAACCCGTTCGGGTTCATCGAAACGCCGTACCGCAAGGTGGTCGACGGCGTGGTTAGCGA CGAGATCGTGTACCTGACCGCCGACGAGGAGGACCGCCACGTGGTGGCACAGGCCAATTCGCCGAT CGATGCGGACGGTCGCTTCGTCGAGCCGCGCGCGTGCTGGTCCGCCCGC
MEP492-2	D-DNA	I491F mutant target	A1471T	GATGTGCCCGATCGAAACCCCTGAGGGGCCCAACATCGGTCTG <u>TTC</u> GGCTCGCTGTCGGTGTACGC GCGGGTCAACCCGTTCGGGTTCATCGAAACGCCGTACCGCAAGGTGGTCGACGGCGTGGTTAGCGA CGAGATCGTGTACCTGACCGCCGACGAGGAGGACCGCCACGTGGTGGCACAGGCCCAATTCGCCGAT CGATGCGGACGGTCGCTTCGTCGAGCCGCGCGTGCTGGTCCGCCGCAAGGCGGGCG
MEP493	D-DNA	I491N mutant target	T1472A	GATGTGCCCGATCGAAACCCCTGAGGGGCCCAACATCGGTCTGAACGGCTCGCTGTCGGTGTACGC GCGGGTCAACCCGTTCGGGTTCATCGAAACGCCGTACCGCAAGGTGGTCGACGGCGTGGTTAGCGA CGAGATCGTGTACCTGACCGCCGAGGAGGAGGACCGCCACGTGGTGGCACAGGCCAATTCGCCGAT CGATGCGGACGGTCGCTTCGTCGAGCCGCGCGCGTGCTGGTCCGCCCGC
MEP494	D-DNA	I491M mutant target	C1473A	GATGTGCCCGATCGAAACCCCTGAGGGGCCCAACATCGGTCTGATAGGCTCGCTGTCGGTGTACGCGCGGGTCAACCCCGTTCGGGTTACGCGCGGGGTCAACCCGTTCGGGTTCATCGAAACGCCGTACCGCAAGGTGGTCGACGGCGTGGTTAGCGACGAGATCGTGTACCTGACCGCCGACGAGGAGGACGCCCACGTGGTGGCACAGGCCAATTCGCCGATCGAT