

Search Details

Job Title/ Assay Name : test sequence	Amplicon Sequence: GGGTAGCAGACCTCACCTATGTGTCGACCTGGGCAGGGTTCGCCTACGTGGCCTTTGTCACCGACGCCTACGCT
Effective Date : 2025-11-11	Forward: GGGTAGCAGACCTCACCTATG
Database : core_nt	Reverse: AGCGTAGGCGTCGGTGA
Alignment algorithm : blastN	Probe: CGTAGGCGAACCCTGCCCAGGTCTGA
Exclusive Organism:	Inclusive Organism: Mycobacterium tuberculosis (1773)
RID: H60NM393014	Total Result: 871
App Version: 1.0	

Species/tax-id Detected	# Accessions Returned	# Meeting Inclusion Criteria	# Accessions Analyzed	# Accession Detected	% Detected
Mycobacterium tuberculosis/1773	805	793	788	781	99%
Mycobacterium tuberculosis variant bovis/1765	53	52	52	52	100%
Mycobacterium tuberculosis variant microti/1806	8	8	8	8	100%
Mycobacterium tuberculosis variant africanum/33894	3	3	3	3	100%
Mycobacterium tuberculosis variant caprae/115862	2	2	2	2	100%
Total	871	858	853	846	99%