

Search Details

Job Title/ Assay Name : glabrata(in)	Amplicon Sequence: GCCTGTTTGAGCGTCATTTCTTCTCAAACACGTTGTGTTTGGTAGT GAGTGATACTCTCGTTTTTGAGTTAACTTGAAATTGTAGGCCATATC AGTATGTGGGACACGAGCGCAAGCTTCTCTATTAATCTGCTGCTCGT TTGCGCGAGCGGCGGGGGTTAATACTGTATTAGGTTTTACCAACTC GGTGTTGATCTAGGGAGGGATAAGTGAGTGTTCTGTGCGTGCTGGG CAGACAGACGTCTTTAAGTTTGACCTCAAATCCGGTAGGGTTACCC GCTGAACTTAAGCATATCAATAAGCGGAGG
Effective Date : 2025-10-13	Forward: GCAGACAGACGTCTTTAAGTT
Database : core_nt	Reverse: CCTCCGCTTATTGATATGCTTA
Alignment algorithm : blastN	Probe: CCTCAAATCCGGTAGGGTTACCCGCTG
Exclusive Organism:	Inclusive Organism: Nakaseomyces glabratus (5478)
RID: ET27WZH7014	Total Result: 1573
App Version: 1.0	

SPECIES/TAX-ID DETECTED	# ACCESSIONS RETURNED	# MEETING INCLUSION CRITERIA	# Accessions Analyzed	# ACCESSION DETECTED	% DETECTED
Nakaseomyces glabratus/5478	1573	898	424	305	72%
Total	1573	898	424	305	72%