

Search Details

Job Title/ Assay Name : Candida glabrata	Amplicon Sequence: CCCTTAACGAGGAACAAYTGGAGGGCAAGTCTGGTGCCAGCAGCC GCGGTAATTCCAGCTCCAATAGCGTATATT
Effective Date : 2025-08-18	Forward: CCCTTAACGAGGAACAATTGGA
Database : core_nt	Reverse: AATATACGCTATTGGAGCTGGAAT
Alignment algorithm : blastN	Probe: AGTCTGGTGCCAGCAGCCGC
Exclusive Organism:	Inclusive Organism: Nakaseomyces glabratus
RID: A60GZGKU013	Total Result: 38
App Version: 1.0	

#Accessions	Oligo	Mutation	Impact
Nakaseomyces glabratus/5478 Analyzed Accessions: 26			
18	Forward	C1-	Detected
1	Forward	C1-	Detected
	Reverse	A1-, A2-	
1	Forward	C1-, G12A	Detected
2	Forward	C1-	Detected
	Probe	+16C	
1	Forward	C1-	Detected
	Reverse	A11T	
1	Forward	C1-	Detected
	Reverse	A1-, A2-, T3-, A4-, T5-, A6-	
1	Forward	C1-, C2-, C3-, T4-, A17T, Y18-, T19-, G20-	Not Detected
	Probe	+6T	
	Reverse	+17G	
1	Forward	C1-, C2-, A14-, A17T, Y18-, T19-, G20-	Not Detected
	Probe	C17-	
	Reverse	A1-, +7C, T13-, A23-	
1	Forward	C1-, C2-, C3-, T4-, T5-, A6-, A7-, C8-, G9-, A10-, G11-, G12-, A13-, A14-, C15-, A16-	limited coverage (excluded)
1	Forward	C1-, C2-, C3-, T4-, T5-, A6-, A7-, C8-, G9-, A10-, G11-, G12-, A13-, A14-, C15-, A16-, A17-, Y18-, T19-	limited coverage (excluded)
1	Forward	C1-, C2-, C3-, T4-, T5-, A6-, A7-, C8-, G9-, A10-, G11-, G12-, A13-, A16C	limited coverage (excluded)
	Probe	A1T, C4T	
1	Forward	C1-, C2-, C3-, T4-, T5-, A6-, A7-, C8-, G9-, A10-, G11-, G12-, A13-, A16T	limited coverage (excluded)

#Accessions	Oligo	Mutation	Impact
	Probe	A1T	
1	Forward	Forward Missing	limited coverage (excluded)
	Probe	A1-, G2-, T3-, C4-	
2	Forward	Forward Missing	limited coverage (excluded)
	Probe	Probe Missing	
3	Forward	Forward Missing	limited coverage (excluded)
	Probe	A1-, G2-, T3-, C4-, T5-, G6-, C18T	
	Reverse	Reverse Missing	
2	Forward	Forward Missing	limited coverage (excluded)
	Probe	Probe Missing	
	Reverse	A1-, A2-, T3-, A4-, T5-, A6-, C7-, G8-, C9-	