

Search Details

Job Title/ Assay Name : L. ramosa	Amplicon Sequence: CCCTTAACGAGGAACAAYTGGAGGGCAAGTCTGGTGCCAGCAGCC GCGGTAATTCCAGCTCCAATAGCGTATATT
Effective Date : 2025-08-20	Forward: CCCTTAACGAGGAACAATTGGA
Database : core_nt	Reverse: AATATACGCTATTGGAGCTGGAAT
Alignment algorithm : blastN	Probe: AGTCTGGTGCCAGCAGCCGC
Exclusive Organism:	Inclusive Organism: Candidozyma auris
RID: AB4751GX013	Total Result: 44
App Version: 1.0	

#Accessions	#Accession Details	Oligo	Mutation	Impact
Candidozyma auris/498019 Analyzed Accessions: 28				
28	28	Forward	C1-	Detected
		Reverse	A11C	
8	MK645720 MK880273 MK645721 MK880276 MK880275 MK880278 MK880274 MK880279	Forward	C1-, C2-, C3-, T4-, T5-, A6-, A7-, C8-, G9-, A10-, G11-, G12-, A13-, A14-, C15-, A16-	limited coverage (excluded)
		Reverse	A11C	
1	MK651238	Forward	C1-, C2-, C3-, T4-, T5-, A6-, A7-, C8-, G9-, A10-, G11-, G12-, A13-, A14-, C15-	limited coverage (excluded)
		Reverse	A11C	
1	LT968778	Forward	Forward Missing	limited coverage (excluded)
		Reverse	A11C	
1	MK880272	Forward	C1-, C2-, C3-, T4-, T5-, A6-, A7-, C8-, G9-, A10-, G11-, G12-, A13-, A14-, C15-, A16-	limited coverage (excluded)
		Probe	C10-, C11-, A12-, G13-, C14-, A15-, G16-, C17-, C18-, G19-, C20-	
		Reverse	Reverse Missing	
2	CP157515 CP157508	Forward	Y18-, T19-, G20-, G21-, A22-	limited coverage (excluded)
		Probe	Probe Missing	
		Reverse	Reverse Missing	
3	CP041140 CP157513 CP157520	Forward	Forward Missing	limited coverage (excluded)
		Probe	A1-, G2-, T3-, C4-, T5-, G6-	
		Reverse	Reverse Missing	