

Search Details

Job Title/ Assay Name : elegans	Amplicon Sequence: CCCTTAACGAGGAACAAYTGGAGGGCAAGTCTGGTGCCAGCAGCC GCGGTAATTCCAGCTCCAATAGCGTATATT
Effective Date : 2025-08-19	Forward: CCCTTAACGAGGAACAATTGGA
Database : core_nt	Reverse: AATATACGCTATTGGAGCTGGAAT
Alignment algorithm : blastN	Probe: AGTCTGGTGCCAGCAGCCGC
Exclusive Organism:	Inclusive Organism: Apophysomyces elegans
RID: A8BCAS9S015	Total Result: 3
App Version: 1.0	

#Accessions	Oligo	Mutation	Impact
Apophysomyces elegans/90269 Analyzed Accessions: 2			
2	Forward	A14T	Detected
1	Forward	Forward Missing	limited coverage (excluded)
	Probe	Probe Missing	
	Reverse	A1-, A2-, T3-, A4-, T5-, A6-, C7-, G8-, C9-, T10-, A11-, T12-, T13-, G14-	