

Search Details

Job Title/ Assay Name : test assay dec19th zee	Amplicon Sequence: CCCAGATCGTTTACCGTGAAAAAATTAGAGTGTTCAAAGCAGGCAG ATCCAATTTTCTGCCACCGAATACATTAGCATGGGATAATGGAATAG GACCCTGTCCTCCTATTTTCAGTTGGTTTTGGCAGCGCGAGGACTA GGGTAATGATTAATAGGGATAGTTGGGGGCATTAATATTTA
Effective Date : 2025-12-18	Forward: CCCAGATCGTTTACCGTGAA
Database : core_nt	Reverse: GGATAGTTGGGGGCATTAATATTTA
Alignment algorithm : blastN	Probe: CTGCCACCGAATACATTAGCATGG
Exclusive Organism:	Inclusive Organism: Acanthamoeba byersi (2506114)
RID: M7M0ASBV016	Total Result: 11
App Version: 1.0	

No of Accessions	Accession Details	Oligo Select Oligo ▼	Mutation	BDS Analysis Performance Prediction
Acanthamoeba byersi/2506114 Analyzed Accessions: 1				
1	MN153028	NA	No Mutations	✓✓✓ Detected
4	OR352425 PP767958 OR352420 LC883911	Forward XXXX	Forward Missing	XXXX limited coverage (excluded)
1	OQ247943	Forward XXXX	Forward Missing	XX limited coverage (excluded)
		Probe ✓✓	+9C	
1	PV637010	Forward XXXX	Forward Missing	XXXX limited coverage (excluded)
		Reverse XXXX	Reverse Missing	
1	OQ247944	Forward XXXX	Forward Missing	XX limited coverage (excluded)
		Probe ✓✓	+9G	
3	PV637008 OQ247940 OQ247962	Forward XXXX	Forward Missing	XXXX limited coverage (excluded)
		Probe XXXX	Probe Missing	