

Search Details

Job Title/ Assay Name :	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: Aspergillus flavus
RID: A8N7013P015	Total Result: 76
App Version: 1.0	

#Accessions	Oligo	Mutation	Impact
Aspergillus flavus/5059 Analyzed Accessions: 67			
57	NA	No Mutations	Detected
1	Forward	C1-	Detected
1	Forward	G11A	Detected
1	Probe	G9C	Detected
1	Reverse	T13G	Detected
1	Forward	A13T	Detected
1	Forward	+12G	Detected
1	Reverse	C7A, +8G	Detected
1	Forward	G11A	Detected
	Probe	G19C	
	Reverse	C9T	
1	Forward	C8G	Not Detected
	Probe	G2T, G9C	
1	Forward	G11A	Not Detected
	Probe	A12C, G16A	