

Search Details

Job Title/ Assay Name :	Amplicon Sequence: TTCTCGTTCCTCACTGCCTCCCCGCCCTGGACACCCTCTTCGTCGT CAGCACCGTCATCCACACCTTATCGTTTTTGTGTATTGTTGCGATGG
Effective Date : 2025-08-05	Forward: TTCTCGTTCCTCACTGCCTCCC
Database : core_nt	Reverse: CCATCGCAACAATACACA
Alignment algorithm : blastN	Probe: TTCGTCGTCAGCACCGTCATCCA
Exclusive Organism:	Inclusive Organism: Human alphaherpesvirus 1
RID: 93411A7S013	Total Result: 787
App Version: None	

#Accessions	Oligo	Mutation	Impact
Simplexvirus humanalpha1/3050292 Analyzed Accessions: 733			
353	NA	No Mutations	Detected
279	Reverse	A9C	Detected
1	Reverse	A8C	Detected
4	Forward	C12T	Detected
8	Forward	T15A	Detected
6	Probe	C3T	Detected
2	Reverse	C5T	Detected
1	Probe	G4A	Detected
1	Forward	C14T	Detected
4	Reverse	G6A	Detected
1	Forward	T1-, T2-	Detected
	Reverse	A9C	
23	Reverse	C5A	Detected
16	Probe	C12T	Detected
	Reverse	A9C	
1	Reverse	A9C, A11C	Detected
2	Probe	C21A	Detected
	Reverse	A9C	
8	Probe	G7A	Detected
5	Forward	C12T	Detected
	Reverse	A9C	
2	Forward	T15C	Detected
2	Probe	C3T	Detected
	Reverse	A9C	
1	Forward	T15C	Detected

#Accessions	Oligo	Mutation	Impact
	Probe	T17G	
1	Forward	C12T	Detected
	Reverse	C5T, A9C	
1	Probe	C6T	Detected
	Reverse	C5A	
2	Forward	C12T, T15A	Not Detected
1	Probe	C18T	Not Detected
	Reverse	C17A	
7	Forward	C12T, T15A	Not Detected
	Reverse	C17A	
1	Forward	C12T, T15A	Not Detected
	Reverse	A9C, C17A	