

Search Details

Job Title/ Assay Name :	Amplicon Sequence: TTCTCGTTCCTCACTGCCTCCCCGCCCTGGACACCCTCTTCGTCGT CAGACCGTCATCCACACCTTATCGTTTTTGTGTATTGGTGCGATGG
Effective Date : 2025-08-19	Forward: TTCTCGTTCCTCACTGCCTCCC
Database : core_nt	Reverse: CCATCGCAACAATACACA
Alignment algorithm : blastN	Probe: TTCGTCGTCAGCACCGTCATCCA
Exclusive Organism:	Inclusive Organism: Human alphaherpesvirus 1
RID: A83GCUW9013	Total Result: 788
App Version: 1.0	

#Accessions	Oligo	Mutation	Impact
Simplexvirus humanalpha1/3050292 Analyzed Accessions: 734			
279	NA	No Mutations	Detected
1	Forward	T1-, T2-	Detected
354	Reverse	C9A	Detected
16	Probe	C12T	Detected
5	Forward	C12T	Detected
2	Probe	C21A	Detected
1	Reverse	A11C	Detected
2	Probe	C3T	Detected
8	Forward	T15A	Detected
	Reverse	C9A	
2	Reverse	C5T, C9A	Detected
4	Reverse	G6A, C9A	Detected
4	Forward	C12T	Detected
	Reverse	C9A	
1	Probe	G4A	Detected
	Reverse	C9A	
1	Forward	C14T	Detected
	Reverse	C9A	
6	Probe	C3T	Detected
	Reverse	C9A	
1	Reverse	A8C, C9A	Detected
1	Forward	C12T	Detected
	Reverse	C5T	
23	Reverse	C5A, C9A	Detected
2	Forward	T15C	Detected

#Accessions	Oligo	Mutation	Impact
	Reverse	C9A	
8	Probe	G7A	Detected
	Reverse	C9A	
1	Forward	T15C	Detected
	Probe	T17G	
	Reverse	C9A	
1	Probe	C6T	Detected
	Reverse	C5A, C9A	
2	Forward	C12T, T15A	Not Detected
	Reverse	C9A	
1	Probe	C18T	Not Detected
	Reverse	C9A, C17A	
1	Forward	C12T, T15A	Not Detected
	Reverse	C17A	
7	Forward	C12T, T15A	Not Detected
	Reverse	C9A, C17A	