

humanparainfluenzavirus1_HN

Mutation Type	Comparison	P-Value
A–G	Syn v CpG	0.6238467
	SynCpG v Non CpG	0.2720918
	NonSynCpG v NonCpG	<0.01
T–C	Syn v CpG	0.01760576
	SynCpG v Non CpG	1.000000
	NonSynCpG v NonCpG	<0.01