

humanparainfluenzavirus1_F

Mutation Type	Comparison	P-Value
A–G	Syn v CpG	0.2952929
	SynCpG v Non CpG	0.2280861
	NonSynCpG v NonCpG	<0.01
T–C	Syn v CpG	<0.01
	SynCpG v Non CpG	1.000000
	NonSynCpG v NonCpG	<0.01