

A

HPyV6-627a	CAATTCTGCTGCCTAATTTTGGCTTCTGGATCCAATAC-TG
HPyV6-607b	CAATTCTGCTGCCTAATTTTGGCTTCTGGATCCAATAC-TG
HPyV6-607a	CAATTCTGCTGCCTAATTTTGGCTTCTGGATCCAATAC-TG
HPyV6-SewagePCR	CAATTCNGCTGCCTAATTTTGGCTTCTGGATCCAATACCTG *****
HPyV6-627a	CAGTATAACCAGTCTCGGGGAGCCCCCTACACCTTTACTGA
HPyV6-607b	CAGTATAACCAGTCTCGGGGAGCCCCCTACACCTTTACTGA
HPyV6-607a	CAGTATAACCAGTCTCGGGGAGCCCCCTACACCTTTACTGA
HPyV6-SewagePCR	CAGTATAACCAGTCTCGGGGAGCCCCCTACACCTTTACTGA *****
HPyV6-627a	CACTTTGGATGCTGGTTCATCT-CTGTGTTATACCC
HPyV6-607b	CACTTTGGATGCTGGTTCATCT-CTGTGTTATACCC
HPyV6-607a	CACTTTGGATGCTGGTTCATCT-CTGTGTTATACCC
HPyV6-SewagePCR	CACTTTGGATGCTGGTTCATCTACTGTGTTATACCC *****

B

SA2/1-152	GCTGGGAATAACCCCGGGCGAACCCTTGTCCATGCCCTA
SA4/1-153	GCTGGGAATAACCCCGGGCGAACCCTTGTCCATGCCCTA
SA3/1-153	GCTGCGGATGACCCTGGGCGAATCCCTGCCCATGCCCTA
SA1-1/1-153	GCTGCGGATGACCCTGGGCGAATCCCTGCCCATGCCCTA
6AIF/129-280	GCTGCGGATAACCCCTGGGCGAACCCTTGTCCATGCCCTA
RatR63/802-954	GCTGCGGTTGACCCTGGGCGAATCCCTGCCCATGCCCTA
GIII/1007-1164	GCCACCCTGGACGATCAGGCGTTTTGTTGTTCCGGCTCAT
GIV/991-1148	GCGACCCTTGATGACCAGGCTTTTTGCTGTTCCAGGCTTAT
GI/1009-1166	GCCACTTTGGATGACCAAGCCTTTTTGCTGCTCCGTTTAAAT
GII/985-1142	GCCACCCTCGACGACCAGGCTTTTTGCTGCTCAGGCTTAT
GV/1000-1157	ACCACACTTGATGATGACGCCTTCTGTTGCTCCGCTGCT * * ** ** *

SA2/1-152	CACCCCTACCCGAAGGCTACCACCATCTATGTGCGCAGCC
SA4/1-153	CACCCCTACCCGAAGGCTACCACCATCTATGTGCGCAGCC
SA3/1-153	CACCCCTACCCGAAAGCCACTACAATCTATGTGCGCAGCC
SA1-1/1-153	CACCCCTACCCGAAAGCCACCACAATTTATGTGCGCAGCC
6AIF/129-280	CACCCCTACCCGAAGGCCACCACCATTTATGTGCGCAGTC
RatR63/802-954	CACCCCTACCCGAAGGCCACCACCATCTATGTGCGCAGTT
GIII/1007-1164	GACTTCTCCGTGGTATTAGCTACAATCACTGTCGGTGCGC
GIV/991-1148	GACATCTCCGTGGTATCAGCTATAATCACGGTTGGCGCTC
GI/1009-1166	GACCTTTTCGCGGCATTAGCTACAATCACTGTTGGCACCC
GII/985-1142	GACGTCTTCGTGGCATTAGCTATAATAACTGTGGGTGCC
GV/1000-1157	AACCTTTTACGTGGGATTTCAACCAATGACAGTAGGTAACA ** * * * ** ** *

SA2/1-152	TGTTTAGTGCCGGTGGAATACCGGTCCGCTAGGCCCCCC
SA4/1-153	TGTTTAGTGCCGGTGGAATACCGGCCCGCTAGGCCCCCC
SA3/1-153	TGTTTAGCGCCGGCGGGAACGCCGGCCCGTTAGGCCCCCC
SA1-1/1-153	TGTTTAGCGCCGGCGGGAACGCCGGCCCGTTAGGCCCCCC
6AIF/129-280	TGTTTAGCGCCGGTGGAATGCCGGCCCATTAGGCCCCCC
RatR63/802-954	TGTTTAGTGCCGGCGGGCAGATATCGGCCCGCTCGGCCCCCC
GIII/1007-1164	TTGTTAATGAGGGATGGAATGCCTCTGAGGACGCCCTTAC
GIV/991-1148	TCGTCAATGAAGGCTGGAATGCCTCCGAAGACGCATTGAC
GI/1009-1166	TTGTTAATGAAGGCTGGAATGCCTCTGAGGACGCCCTCAC
GII/985-1142	TGGTTAATGAAGGCTGGAATGCCACCGAGGATGCGCTCAC
GV/1000-1157	TTGTCAATGAGGGCTGGCAACCGGAGGAGCAGCAGCTTAC * * * * * * *

SA2/1-152	CCCCATGATGGGNGAAAAAGCCCGGTCCAC
SA4/1-153	CCCCATGATGTGGGAAAAAGCCCGGGCCAC
SA3/1-153	TCCCATGATGTGTGAAAAGGCCCGGTCCAC
SA1-1/1-153	TCCTATGATGTGTGAAAAGGCCCGGTCCAC
6AIF/129-280	TCCCTTGATGTGTGAGAAGGCCCGGTCCAC
RatR63/802-954	CCCATTAATGTGTGAGAAGGCCCGGTCTAC
GIII/1007-1164	TGCAGTGATCACTGCGGCTTACCTGACTAT
GIV/991-1148	AGCTGTGATTACCGCGGCTATCTTACTAT
GI/1009-1166	AGCTGTCATCACTGCCGCTTACCTTACCAT
GII/985-1142	TGCAGTTATTACGGCGGCTTACCTCACAAT
GV/1000-1157	GGCTGTCGCTATAGCTGCATACCTTACTGT * * * * *

