

BKV_2.1 (f) 5'-CCCAGGAGGTGCTAATCAAAGA-3'
Polyoma 4.2 (f) 5'-GCTCCTCAATGGATGTTGCCT-3'
BKV-MGB Probe 5'-(6FAM)-CCGTACAGGCCTAGAA-(MGB-NFQ)-3'
BKV_4.1 (r) 5'-TCAIATGCTTCAAGAGCAGGTGT-3'

1 30 60 90 120
NC_007611.1|1347-1589| SA12 TCAGGTGAATTTATAGAAAAAAGTCTAGCACCAGGAGGTGCCAATCAAAGAATTGCTCCTCAATGGATGTTGCCTTTGCTTCTAGGCCTGTACGGGACTGTAACACCTGCTCTTGAAGCATATGAAGATG
NC_001669.1|1378-1620| SV40 TCAGGTGAATTTATTGAAAAATTTGAGGCTCCTGGTGGTGCAAAATCAAAGAAGTCTGCTCCTCAGTGGATGTTGCCTTTACTTCTAGGCCTGTACGGGACTGTAACACCTGCTCTTGAAGCATATGAAGATG
NC_001699.1|1342-1560| JCV TCTGGTGAATTTATAGAAAGAAGTATTGCACCAGGAGGTGCAAAATCAAAGATCTGCTCCTCAATGGATGTTGCCTTTACTTTTAGGGTTGTACGGGACTGTAACACCTGCTCTTGAAGCATATGAAGATG
.....G.CC.....G.A..ACY.A..G.G.....AG.....GG...A.....A.....T.....C.....G.....
NC_001538.1|1437-1679| BKV TCAGGAGAGTTTATAGAAAAAAGTATTGCCAGGAGGTGCTAATCAAAGAAGTCTGCTCCTCAATGGATGTTGCCTTTACTTCTAGGCCTGTACGGGACTGTAACACCTGCTCTTGAAGCATATGAAGATG
.....G.....
.....G.....
.....G.....
.....A.....
.....G.....

BKV_5.1 (r) 5'-CTGCCCTGGACACTCTC-3'

150 180 210 240
NC_007611.1|1347-1589| SA12 CCCCCAGCAAAAAGAAAAGGAGAATGTCCAGGGGCAGCTCCCAAAAAGCCAAAGGACCCCGTGCGAGTTCCTTATAAGAGGAGGCGTAGAAGTACTAGAAGTTAA
NC_001669.1|1378-1620| SV40 GCCCCAACAAAAAGAAAAGGAGTGTCCAGGGGCAGCTCCCAAAAAGCCAAAGGAAACCAGTGCAGTGCCTCAAAAAGCTCGTCATAAAAAGGAGGAATAGAAGTCTAGGAGTTAA
NC_001699.1|1342-1560| JCV GCCCCAACAAAAAGAAAAGGAGAAAAG-----GAAGGACCCCGTCAAGTTCCTTATAAGAGGAGGAGTAGAAGTCTAGGAGTTAA
.....C...G.....C.....-----...C.T...C..G.....A.....GT.....
NC_001538.1|1437-1679| BKV GCCCCAACAAAAAGAAAAGGAGAGTGTCCAGGGGCAGCTCCCAAAAAGCCAAAGGAAACCAGTGCAGTGCCTCAAAAAGTACTAATAAAAAGGAGGAGTAGAAGTCTAGGAGTTAA
..... [56/271]
..... [73/271]
.....C..... [87/271]
.....C..... [7/271]
.....A..... [1/271]
.....A..... [38/271]
.....A.....G..... [6/271]
.....A...A..... [1/271]
.....A.....A..... [1/271]
.....A.....G..... [1/271]
.....A..... [1/271]
.....G..... [1/271]
.....G..... [1/271]
.....G..... [1/271]