

CMV promoter
 721 AAAATCAACG GGACTTTCCA AAATGTCGTA ACAACTCCGC CCCATTGACG CAAATGGGCG
 CMV forward priming site TATA 3' end of CMV promoter putative transcriptional start
 781 GTAGGCGTGT ACGGTGGGAG GTCTATATAA GCAGAGCTCT CTGGCTAACT AGAGAACCCA
 T7 promoter/priming site *Nhe* I
 841 CTGCTTACTG GCTTATCGAA ATTAATACGA CTCACTATAG GGAGACCCAA GCTGGCTAGC
Pme I* *Afl* II *Hind* III *Asp* 718 I *Kpn* I *Bam* H I *Bst* X I*
 901 GTTTAAACTT AAGCTTGGTA CCGAGCTCGG ATCCACTAGT CCAGTGTGGT GGAATTCTGC
Eco R V *Bst* X I* *Not* I *Xho* I *Apa* I *Pme* I*
 961 AGATATCCAG CACAGTGGCG GCCGCTCGAG TCTAGAGGGC CCGTTTAAAC CCGCTGATCA
 BGH reverse priming site
 1021 GCCTCGACTG TGCCTTCTAG TTGCCAGCCA TCTGTTGTTT GCCCCTCCCC CGTGCCTTCC