

721 AAAATCAACG GGACTTTCCA AAATGTCGTA ACAACTCCGC CCCATTGACG CAAATGGGCG ^{CMV Forward priming site}

781 GTAGGCGTGT ACGGTGGGAG GTCTATATAA GCAGAGCTCT CCCTATCAGT GATAGAGATC
^{TATA box} ^{Tetracycline operator (TetO₂)}

841 TCCCTATCAG TGATAGAGAT CGTCGACGAG CTCGTTTAGT GAACCGTCAG ATCGCCTGGA
^{Tetracycline operator (TetO₂)}

901 GACGCCATCC ACGCTGTTTT GACCTCCATA GAAGACACCG GGACCGATCC AGCCTCCGGA

961 CTCTAGCGTT TAAACTTAAG CTTGGTACCG AGCTCGGATC CACTAGTCCA GTGTGGTGGA
^{Pme I*} ^{Afl II} ^{Hind III} ^{Asp718 I} ^{Kpn I} ^{Bam HI} ^{BstX I*} ^{EcoR I}

1021 ATTCTGCAGA TATCCAGCAC AGTGCGGGCC GCTCGAGTCT AGAGGGCCCG CGGTTC GAA
^{Pst I} ^{EcoR V} ^{BstX I*} ^{Not I} ^{Xho I} ^{Xba I} ^{Apa I} ^{Sac II} ^{Glu}

1080 CAA AAA CTC ATC TCA GAA GAG GAT CTG AAT ATG CAT ACC GGT CAT CAT
^{c-myc epitope} ^{Age I}
 Gln Lys Leu Ile Ser Glu Glu Asp Leu Asn Met His Thr Glu His His

1128 CAC CAT CAC CAT TGA GTTTAAAC CCGCTGATCA GCCTCGACTG TGCCTTCTAG
^{Polyhistidine (6xHis) region} ^{Pme I*} ^{BGH Reverse priming site}
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