

2378 TCGTAACAAC TCCGCCCCAT TGACGCAAAT GGGCGGTAGG CGTGTACGGT GGGAGGTCTA TATAAGCAGA GCTCTCCCTA
 CAAT CMV forward priming site TATA
 Tetracycline operator (TetO₂) Tetracycline operator (TetO₂) Spe I
 2458 TCAGTGATAG AGATCTCCCT ATCAGTGATA GAGATCGTCG ACTAGTCCAG TGTGGTGGAA TTCTGCAGAT ATCAACAAGT
 TAGTTGTTCA
 2539 4222
 2538 TTGTACAAAA AAGCAGGCTN --- ... Pro Ala Phe Leu Tyr Lys Val Val Asp Ile Gln His Ser Gly
 AACATGTTTT TTCGTCCGAN -GENE- NTG GGT CGA AAG AAC ATG TTT CAC CAA CTA TAG GTC GTG TCA CCG
 attB1 attB2
 4256 Gly Arg Ser Ser Leu Glu Gly Pro Arg Phe Glu Gly Lys Pro Ile Pro Asn Pro Leu Leu Gly Leu
 GGC CGC TCG AGT CTA GAG GGC CCG CGG TTC GAA GGT AAG CCT ATC CCT AAC CCT CTC CTC GGT CTC
 CCG GCG AGC TCA GAT CTC CCG GGC GCC AAG CTT CCA TTC GGA TAG GGA TTG GGA GAG GAG CCA GAG
 V5 epitope V5 (C-term) reverse priming site
 4322 Asp Ser Thr Arg Thr Gly *** *** ***
 GAT TCT ACG CGT ACC GGT TAG TAA TGA
 CTA AGA TGC GCA TGG CCA ATC ATT ACT