```
                 CAAT                                  TATA
761   CCCATTGACG CAAATGGGCG GTAGGCGTGT ACGGTGGGAG GTCTATATAA GCAGAGCTCT CTGGCTAACT

                                        T7 Promoter priming site
831   AGAGAACCCA CTGCTTACTG GCTTATCGAA ATTAATACGA CTCACTATAG GGAGACCCAA GCTGGCTAGT
                     918        attB1                                3164   attB2
901   TAAGCTATCA ACAAGTTTGTA CAAAAAAGCA GGCTNN-GENE-NAC CCA GCT TTC TTG TAC AAA GTG GTT
                 TGTTCAAACAT GTTTTTTCGA CCGTNN-GENE-NTG GGT CGA AAG AAC ATG TTT CAC CAA
                                                     Pro Ala Phe Leu Tyr Lys Val Val

                                                V5 reverse priming site
                                                                          V5 epitope
3180  GAT CTA GAG GGC CCG CGG TTC GAA GGT AAG CCT ATC CCT AAC CCT CTC CTC GGT CTC GAT
      CTA GAT CTC CCG GGC GCC AAG CTT CCA TTC GGA TAG GGA TTG GGA GAG GAG CCA GAG CTA
      Asp Leu Glu Gly Pro Arg Phe Glu Gly Lys Pro Ile Pro Asn Pro Leu Leu Gly Leu Asp

3240  TCT ACG CGT ACC GGT TAG TAA TGA GTTTAAACGG
      AGA TGC GCA TGG CCA *** *** ***
      Ser Thr Arg Thr Gly
```

invitrogen™