



**Comments for pENTR/pSM2(CMV) GFP
3861 nucleotides**

rrrB T1 transcription termination sequence: bases 106-149

rrrB T2 transcription termination sequence: bases 281-308

attL1: bases 358-457 (complementary strand)

CMV promoter: bases 474-1013

Enhanced green fluorescent protein (eGFP): bases 1083-1802

5'miR30 context: 1813-1930

shRNA cloning sites (Xho I/Eco RI): 1931-1948

3'miR30 context: 1949-2051

attL2: bases 2090-2190

Kanamycin resistance gene: bases 2312-3135

pUC origin: bases 3204-3859

Recommended E.coli strain for propagation: TOP10F'