



Comments for pENTR/pSM2(CMV)  
3063 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

5'miR30 context: 1014-1132

shRNA cloning sites (Xho I/Eco RI):1133-1150

3'miR30 context: 1151-1253

attL2:bases 1292-1392

Kanamycin resistance gene: bases 1514-2335

pUC origin: bases 2407-3060

Recommended *E.coli* strain for propagation: TOP10F' under kanamycin selection.

NOTE: The position of the genetic elements is approximative.