Comments for pCR®8/GW/TOPO®
2817 nucleotides

*rrnB* T2 transcription termination sequence: bases 268-295
*rrnB* T1 transcription termination sequence: bases 427-470
M13 forward (-20) priming site: bases 537-552
*attL1*: bases 569-668
GW1 priming site: bases 607-631
TOPO® recognition site 1: bases 678-682
TOPO® recognition site 2: bases 683-687
*attL2*: bases 696-795
GW2 priming site: bases 733-757
T7 Promoter/priming site: 812-831 (c)
M13 reverse priming site: bases 836-852
Spectinomycin promoter: bases 930-1063
Spectinomycin resistance gene (*Spn*<sup>R</sup>): 1064-2074
pUC origin: bases 2141-2814

(c) = complementary sequence