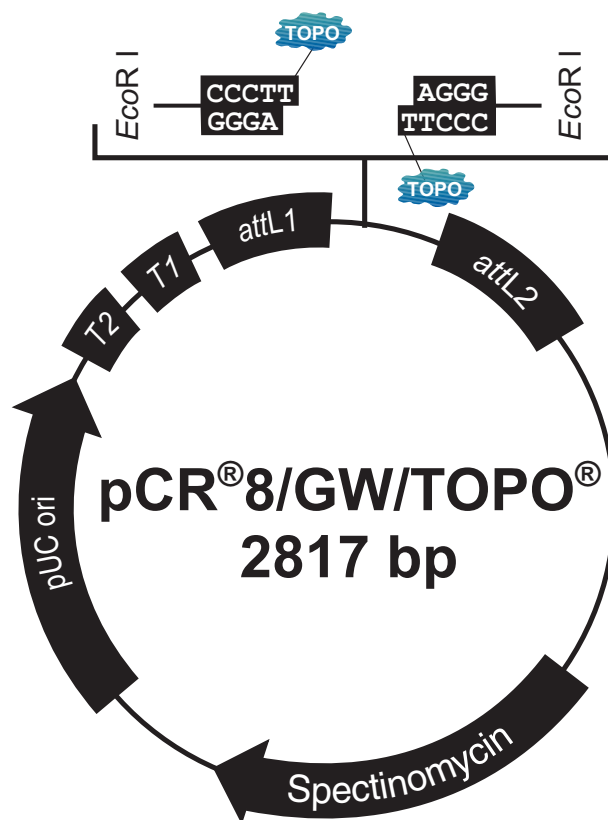




**Comments for pCR®8/GW/TOPO®
2817 nucleotides**

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*^R): 1064-2074

pUC origin: bases 2141-2814

(c) = complementary sequence