



**Comments for pENTR™/D-TOPO®
2580 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 (c)

TOPO® recognition site 1: bases 680-684

Overhang: bases 685-688

TOPO® recognition site 2: bases 689-693

attL2: bases 705-804

T7 Promoter/priming site: bases 821-840 (c)

M13 reverse priming site: bases 845-861

Kanamycin resistance gene: bases 974-1783

pUC origin: bases 1904-2577

(c) = complementary sequence