



**Comments for pENTR™/TEV/D-TOPO®
2610 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)

TEV recognition site: bases 689-709

TOPO® recognition site: bases 710-714

Overhang: bases 715-718

TOPO® recognition site: bases 719-723

attL2: bases 735-834

T7 Promoter/priming site: bases 851-870 (c)

M13 reverse priming site: bases 875-891

Kanamycin resistance gene: bases 1004-1813

pUC origin: bases 1934-2607

(c) = complementary sequence