

Comments for pENTR/D-TOPO® 2580 nucleotides

*rrn*B T2 transcription termination sequence: bases 268-295 *rrn*B T1 transcription termination sequence: bases 427-470

M13 forward (-20) priming site: bases 537-552 attL1: bases 569-668 (complementary strand) TOPO® recognition site 1: bases 680-684

Overhang: bases 685-688

TOPO® recognition site 2: bases 689-693

attL2: bases 705-804

M13 reverse priming site: bases 845-861 Kanamycin resistance gene: bases 974-1783

pUC origin: bases 1904-2577





