



**Comments for pENTR™/SD/D-TOPO®  
2601 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)  
T7 gene 10 translational enhancer: bases 684-692  
Ribosome binding site: bases 694-700  
TOPO® recognition site 1: bases 701-705  
Overhang: bases 706-709  
TOPO® recognition site 2: bases 710-714  
*attL2*: bases 726-825  
T7 Promoter/priming site: bases 842-861 (c)  
M13 reverse priming site: bases 866-882  
Kanamycin resistance gene: bases 995-1804  
pUC origin: bases 1925-2598

(c) = complementary sequence