



*rrnB* T2 transcription terminator: bases 268-295 (c)  
*rrnB* T1 transcription terminator: bases 427-470 (c)  
 M13 forward (-20) priming site: bases 537-552  
*attL4*: bases 592-688  
 GW1 priming site: bases 630-654  
 TOPO® recognition site 1: bases 701-705  
 TOPO® recognition site 2: bases 706-710  
*attR1*: bases 721-845  
 GW3 priming site: bases 752-781  
 M13 reverse priming site: bases 945-961  
 Kanamycin resistance gene: 1074-1883  
 pUC origin: bases 2004-2677

(c) = complementary strand