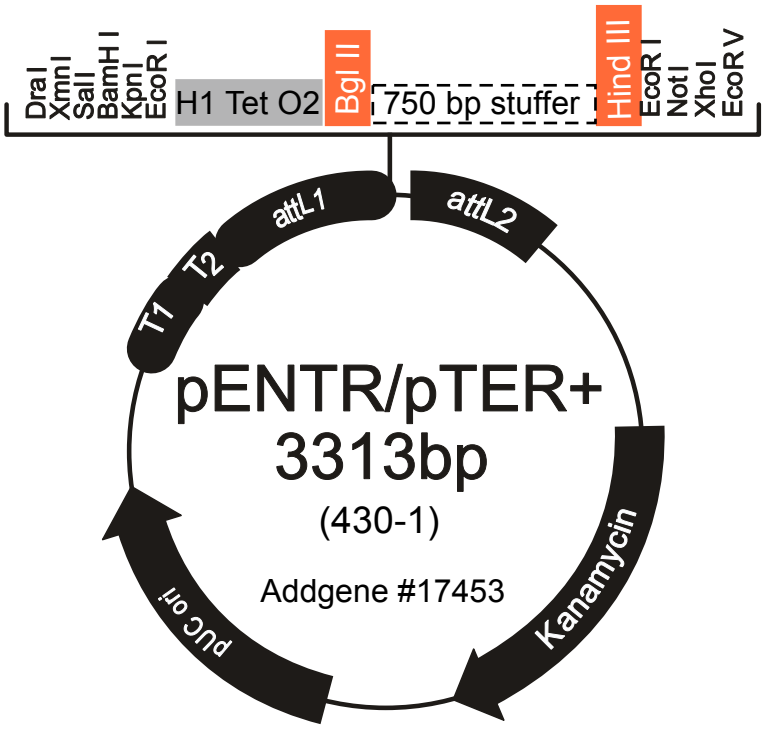


Comment s for pENTR/pTER+
3313 nucleotides

rrnB T1 transcription termination sequence: bases 106-149
rrnB T2 transcription termination sequence: bases 281-308
attL1: bases 358-457 (complementary strand)
H1 RNA promoter: bases 494-715
750 bp stuffer: bases 716-1472
attL2: bases 1542-1640
Kanamycin resistance gene: bases 1764-2573
pUC origin: bases 2635-3308

Recommended *E.coli* strain for propagation: TOP10F' under kanamycin selection.

NOTE: The position of the genetic elements is approximative.

Eric Campeau March 2004