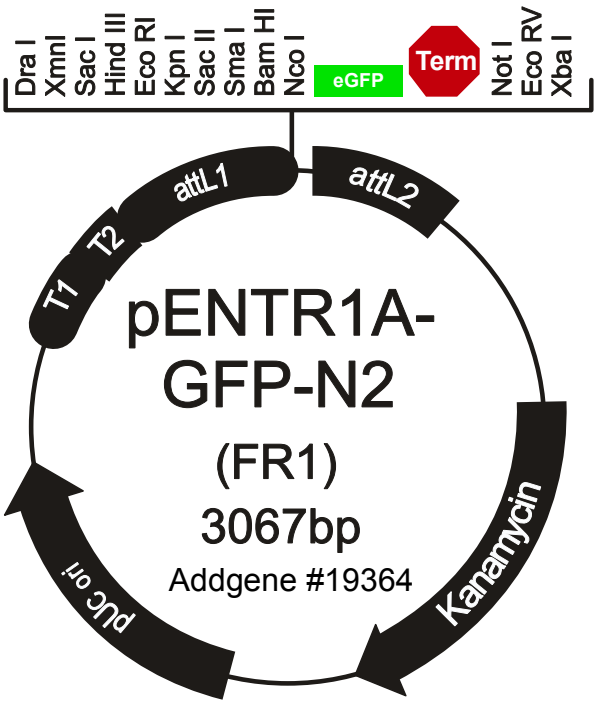
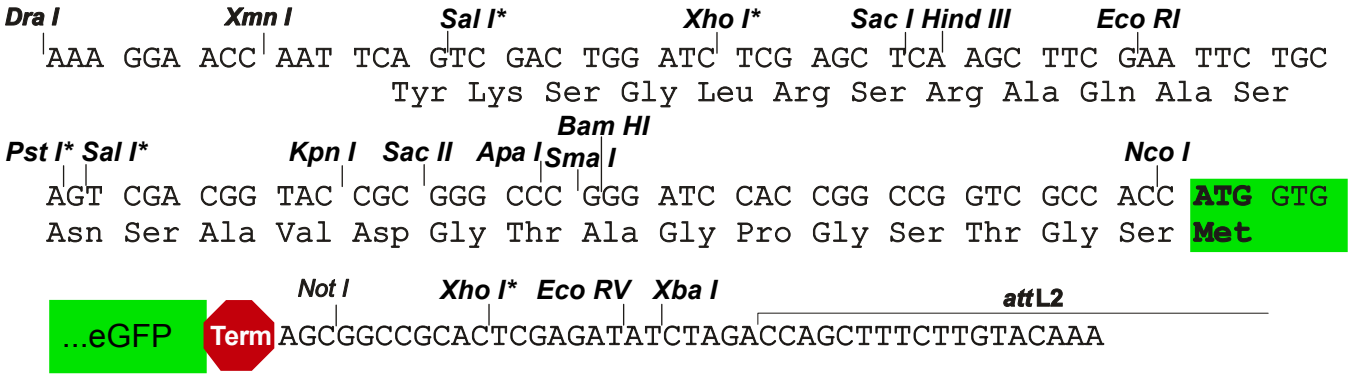


Comments for pENTR1A-GFP-N2
3067 nucleotides

rrnB T1 transcription termination sequence: bases 106-149
rrnB T2 transcription termination sequence: bases 281-308
attL1: bases 358-457 (complementary strand)
enhanced green fluorescent protein (eGFP): bases 552-1271
attL2: bases 1296-1396
Kanamycin resistance gene: bases 1518-2327
pUC origin: bases 2392-3064



Recommended *E.coli* strain for propagation: TOP10F'

The position of the genetic elements is approximative.

Made by Francis Rodier (Lawrence Berkeley National Laboratory)