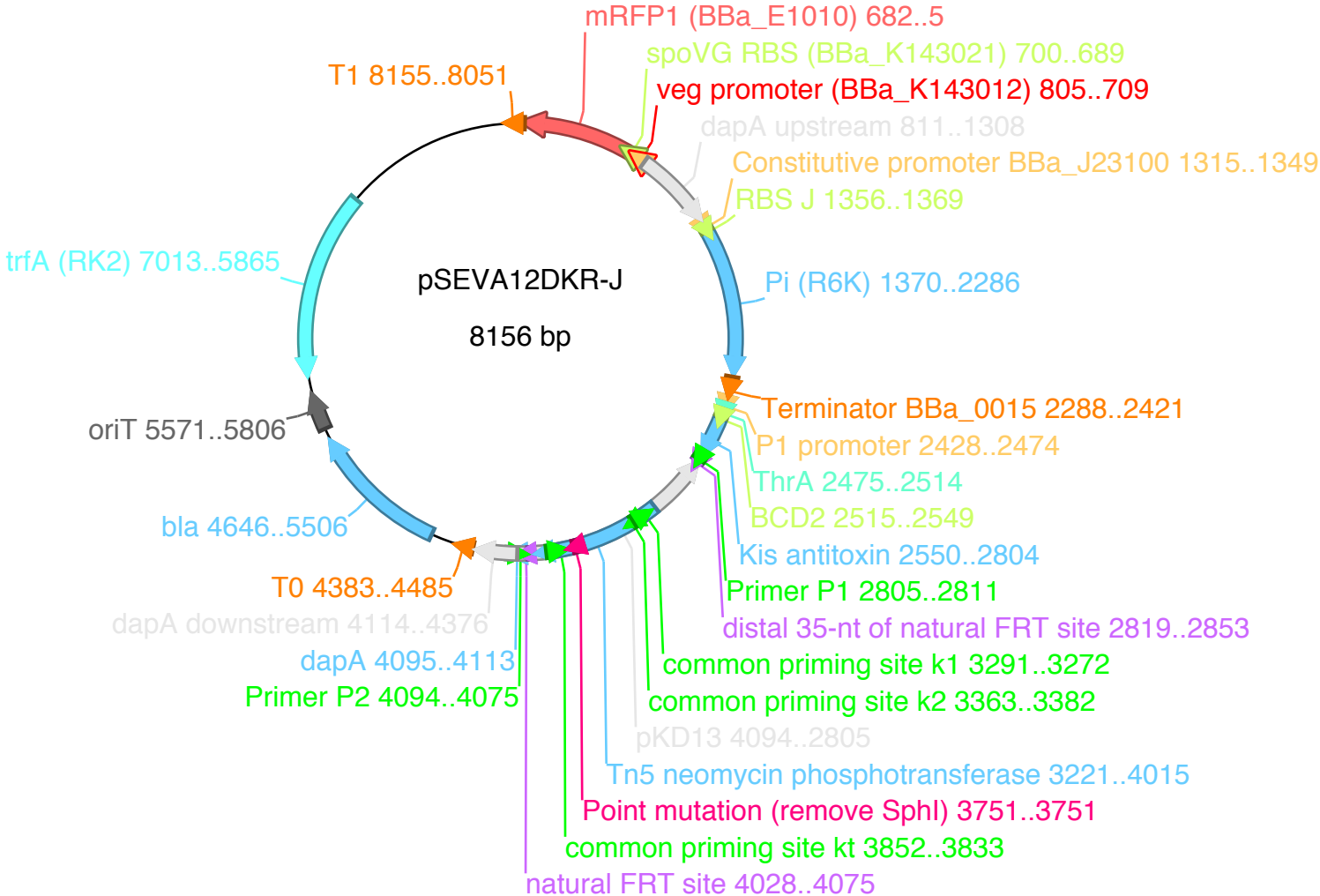
mRFP1 (BBa_E1010) 682..5
spoVG RBS (BBa_K143021) 700..689
veg promoter (BBa_K143012) 805..709
dapA upstream 811..1308
Constitutive promoter BBa_J23100 1315..1349
RBS J 1356..1369
Pi (R6K) 1370..2286
Terminator BBa_0015 2288..2421
P1 promoter 2428..2474
ThrA 2475..2514
BCD2 2515..2549
Kis antitoxin 2550..2804
Primer P1 2805..2811
distal 35-nt of natural FRT site 2819..2853
common priming site k1 3291..3272
common priming site k2 3363..3382
pKD13 4094..2805
Tn5 neomycin phosphotransferase 3221..4015
Point mutation (remove SphI) 3751..3751
common priming site kt 3852..3833
natural FRT site 4028..4075
Primer P2 4094..4075
dapA 4095..4113
dapA downstream 4114..4376
T0 4383..4485
bla 4646..5506
oriT 5571..5806
trfA (RK2) 7013..5865
T1 8155..8051
pSEVA12DKR-J
8156 bp