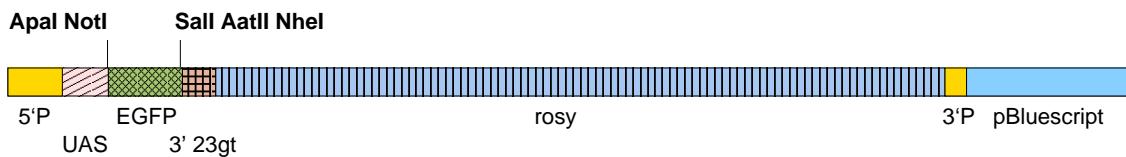


pP{GS[ry⁺, UASEGFP3']}

12434 bp



5' multicloning site:

3' multicloning site:

NheI
AatII
AvrII

1650 aaggcccgggcaagtgcgacgacgtcgctagcatacctaggtagtttaaaatttattgttgc 1709

^ * ^ * ^ * ^ * ^ * ^ * ^ * ^ * ^ * ^ *

K A R A S R R R R *

23gt 3'

unique sites are indicated in **bold**

Location of features:

1 –	541	5' P-repeat
542 –	1004	hsp70 promoter
1005 –	1007	5' multicloning site
1008 –	1748	EBFP
1749 –	1763	3' multicloning site
1764 –	2074	3' 23gt
2075 –	9366	rosy
9367 –	9571	3'P-repeat
9572 –	12434	pBluescriptII KS+