



**Comments for pLenti CMV GFP X2 DEST  
 10463 nucleotides**

- bla promoter: bases 32-131
- ampicillin resistance gene: bases 132-992
- pUC origin: bases 1138-1810
- RSV/5LTR hybrid promoter: bases 2218-2628
- HIV-1 psi ( $\psi$ ) packaging signal: bases 2736-2781
- HIV-1 Rev response element (RRE): bases 3272-3525
- 3 splice acceptor: base 3862
- 3 splice acceptor: base 3901
- Central polypurine tract (cPPT): bases 4011-4073
- CMV promoter: bases 4145-4730
- Green Fluorescent Protein (GFP): bases 4755-5474
- Woodchuck post-transcriptional element (PRE): bases 5489-6083
- Human phosphoglycerate kinase (pGK) promoter: bases 6087-6606
- Enhanced Green Fluorescent Protein (GFP) cDNA: bases 6624-7343
- $\Delta$ U3/3'LTR: bases 7369-9436
- attR2 site: bases 7602-7478
- ccdB gene: bases 7948-7643
- Chloramphenicol resistance gene (Cm): bases 8949-8289
- attR1 site: bases 9183-9059
- SV40 polyadenylation signal: bases 9508-9661

Recommended *E. coli* strain for propagation: DB3.1 or ccDB survival under ampicillin and chloramphenicol selection.

**NOTE: The position of the genetic elements is approximative.**