

**Restriction Map and Multiple Cloning Site (MCS) of pEGFP-C2.** (Unique restriction sites are in bold). The *Eag I* site is not unique. The *Xba I* site cannot be used for fusions since it contains an in-frame stop codon. The *Xba I* and *BcI I* sites (\*) are methylated in the DNA provided by BD Biosciences Clontech. If you wish to digest the vector with these enzymes, you will need to transform the vector into a *dam*<sup>-</sup> host and make fresh DNA.

### **Description:**

pEGFP-C2 encodes a red-shifted variant of wild-type GFP (1-3) which has been optimized for brighter fluorescence and higher expression in mammalian cells. (Excitation maximum = 488 nm; emission maximum = 507 nm.) pEGFP-C2 encodes the GFPmut1 variant (4) which contains the double-amino-acid substitution of Phe-64 to Leu and Ser-65 to Thr. The coding sequence of the EGFP gene contains more than 190 silent base changes which correspond to human codon-usage preferences (5). Sequences flanking EGFP have been converted to a Kozak consensus translation initiation site (6) to further increase the translation efficiency in eukaryotic cells. The MCS in pEGFP-C2 is between the EGFP coding sequences and the SV40 poly A. Genes cloned into the MCS will be expressed as fusions to the C terminus of EGFP if they are in the same reading frame as EGFP and there are no intervening stop codons. SV40 polyadenylation signals downstream of the EGFP gene direct proper processing of the 3' end of the EGFP mRNA. The vector backbone also contains an SV40 origin for replication in mammalian cells expressing the SV40 T-antigen. A neomycin-resistance cassette (Neo'), consisting of the SV40 early promoter, the neomycin/ kanamycin resistance gene of Tn5, and polyadenylation signals from the Herpes simplex virus thymidine kinase (HSV TK) gene, allows stably transfected eukaryotic cells to be selected using G418. A bacterial promoter upstream of this cassette expresses kanamycin resistance in E. coli. The pEGFP-C2 backbone also provides a pUC origin of replication for propagation in E. coli and an f1 origin for single-stranded DNA production.

pEGFP-C2 Vector Information

#### Use:

Fusions to the C terminus of EGFP retain the fluorescent properties of the native protein allowing the localization of the fusion protein *in vivo*. The target gene should be cloned into pEGFP-C2 so that it is in frame with the EGFP coding sequences, with no intervening in-frame stop codons. The recombinant EGFP vector can be transfected into mammalian cells using any standard transfection method. If required, stable transformants can be selected using G418 (7). pEGFP-C2 can also be used simply to express EGFP in a cell line of interest (e.g., as a transfection marker).

### **Location of Features:**

• Human cytomegalovirus (CMV) immediate early promoter: 1-589

Enhancer region: 59-465; TATA box: 554-560

Transcription start point: 583

C→G mutation to remove *Sac* I site: 569
• Enhanced green fluorescent protein gene

Kozak consensus translation initiation site: 606–616 Start codon (ATG): 613–615; Stop codon: 1408–1410

Insertion of Val at position 2: 616-618

GFPmut1 chromophore mutations (Phe-64 to Leu; Ser-65 to Thr): 805-810

His-231 to Leu mutation (A→T): 1307 Last amino acid in wild-type GFP: 1327–1329

MCS: 1343–1421

SV40 early mRNA polyadenylation signal

Polyadenylation signals: 1554-1559 & 1583-1588; mRNA 3' ends: 1592 & 1604

• f1 single-strand DNA origin: 1651–2106 (Packages the noncoding strand of EGFP)

Bacterial promoter for expression of Kan<sup>r</sup> gene

-35 region: 2168-2173; -10 region: 2191-2196

Transcription start point: 2203

• SV40 origin of replication: 2447–2582

• SV40 early promoter

Enhancer (72-bp tandem repeats): 2280–2351 & 2352–2423

21-bp repeats: 2427-2447, 2448-2468 & 2470-2490

Early promoter element: 2503-2509

Major transcription start points: 2499, 2537, 2543 & 2548

• Kanamycin/neomycin resistance gene

Neomycin phosphotransferase coding sequences: Start codon (ATG): 2631–2633; stop codon: 3423–3425

G→A mutation to remove *Pst* I site: 2813

C→A (Arg to Ser) mutation to remove *Bss*H II site: 3159

Herpes simplex virus (HSV) thymidine kinase (TK) polyadenylation signal

Polyadenylation signals: 3661-3666 & 3674-3679

• pUC plasmid replication origin: 4010-4653

## **Primer Locations:**

- EGFP-N Sequencing Primer (#6479-1): 679-658
- EGFP-C Sequencing Primer (#6478-1): 1266–1287

# Propagation in E. coli:

- Suitable host strains: DH5α, HB101, and other general purpose strains. Single-stranded DNA production requires
  a host containing an F plasmid such as JM109 or XL1-Blue.
- Selectable marker: plasmid confers resistance to kanamycin (30 μg/ml) to E. coli hosts.
- E. coli replication origin: pUC
- Copy number: ≈500
- Plasmid incompatibility group: pMB1/ColE1

### References:

- 1. Prasher, D. C., et al. (1992) Gene 111:229–233.
- 2. Chalfie, M., et al. (1994) Science 263:802-805.
- 3. Inouye, S. & Tsuji, F. I. (1994) FEBS Letters **341**:277–280.
- 4. Cormack, B., et al. (1996) Gene **173**:33–38.
- 5. Haas, J., et al. (1996) Curr. Biol. 6:315-324.
- 6. Kozak, M. (1987) Nucleic Acids Res. 15:8125-8148.
- 7. Gorman, C. (1985) In DNA Cloning: A Practical Approach, Vol. II, Ed. Glover, D. M. (IRL Press, Oxford, UK) pp. 143-190.

pEGFP-C2 Vector Information

**Note:** The attached sequence file has been compiled from information in the sequence databases, published literature, and other sources, together with partial sequences obtained by BD Biosciences Clontech. This vector has not been completely sequenced.

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