



**Restriction Map and Multiple Cloning Site (MCS) of pAcGFP1-Hyg-N1 Vector.** Unique restriction sites are shown in bold. NOTE: The *Xba* I and *Bcl* I sites are methylated in the DNA provided by Clontech Laboratories, Inc. 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

pAcGFP1-Hyg-N1 encodes a green fluorescent protein (GFP) from *Aequorea coerulea*. (Excitation maximum = 475 nm; emission maximum = 505 nm). The coding sequence of the AcGFP1 gene contains silent base changes, which correspond to human codon-usage preferences (1). The MCS in pAcGFP1-Hyg-N1 is between the immediate early promoter of CMV ( $P_{CMVIE}$ ) and the AcGFP1 coding sequences. Genes cloned into the MCS will be expressed as fusions to the N-terminus of AcGFP1 if they are in the same reading frame as AcGFP1 and there are no intervening stop codons. SV40 polyadenylation signals downstream of the AcGFP1 gene direct proper processing of the 3' end of the AcGFP1 mRNA. The vector backbone also contains an SV40 origin for replication in mammalian cells expressing the SV40T antigen. A hygromycin-resistance cassette ( $Hyg^r$ ), consisting of the SV40 early promoter, the hygromycin resistance gene, and SV40 polyadenylation signals, allows stably transfected eukaryotic cells to be selected using hygromycin. A bacterial promoter upstream of the ampicillin gene expresses ampicillin resistance in *E. coli*. The pAcGFP1-Hyg-N1 backbone also provides a pUC origin of replication for propagation in *E. coli* and an f1 origin for single-stranded DNA production.

### Use

Fusions to the N-terminus of AcGFP1 retain the fluorescent properties of the native protein, allowing the localization of the fusion protein *in vivo*. The target gene should be cloned into pAcGFP1-Hyg-N1 so that it is in frame with the AcGFP1 coding sequences, with no intervening in-frame stop codons. The inserted gene should include the initiating ATG codon. The recombinant AcGFP1 vector can be transfected into mammalian cells using any standard transfection method. If required, stable clones can be selected using hygromycin. This vector can be used in conjunction with other Living Colors mammalian expression vectors containing a neomycin resistance gene to establish cell lines that simultaneously express two different fluorescent proteins. pAcGFP1-Hyg-N1 can also be used simply to express AcGFP1 in a cell line of interest (e.g., as a transfection marker).

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**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
- Multiple Cloning Site (MCS): 591–671
- *Aequorea coerulescens* Green Fluorescent Protein (AcGFP1): 673–1389  
Start codon (ATG): 673–675; Stop codon: 1390–1392  
Insertion of Val at position 2: 676–678  
Last amino acid: 1387–1389
- SV40 early mRNA polyadenylation signal  
Polyadenylation signals: 1544–1549 & 1573–1578; mRNA 3' ends: 1582 & 1594
- f1 single-strand DNA origin: 1642–2097 (Packages the noncoding strand of AcGFP1.)
- SV40 origin of replication: 2438–2573
- SV40 early promoter  
Enhancer (72-bp tandem repeats): 2271–2342 & 2343–2414  
21-bp repeats: 2418–2438, 2439–2459 & 2467–2481  
Early promoter element: 2494–2500  
Major transcription start points: 2490, 2528, 2534 & 2539
- Hygromycin resistance gene:  
Start codon (ATG): 2595–2597; stop codon: 3618–3620
- SV40 early mRNA polyadenylation signal: 3767–3772 & 3796–3801; mRNA 3' ends: 3805 & 3817
- Bacterial promoter for expression of Amp<sup>r</sup> gene:  
–35 region: 3967–3972; –10 region: 3990–3995
- Ampicillin resistance gene:  
Start codon (ATG): 4037–4039; stop codon: 4895–4897
- pUC plasmid replication origin: 5060–5703

**Propagation in *E. coli***

- Suitable host strains: DH5 $\alpha$  and other general purpose strains.
- Selectable marker: plasmid confers resistance to ampicillin (100  $\mu$ g/ml) in *E. coli* hosts.
- *E. coli* replication origin: pUC

**Reference**

1. Haas, J., et al. (1996) *Curr. Biol.* **6**:315–324.

**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 Clontech Laboratories, Inc. This vector has not been completely sequenced.

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