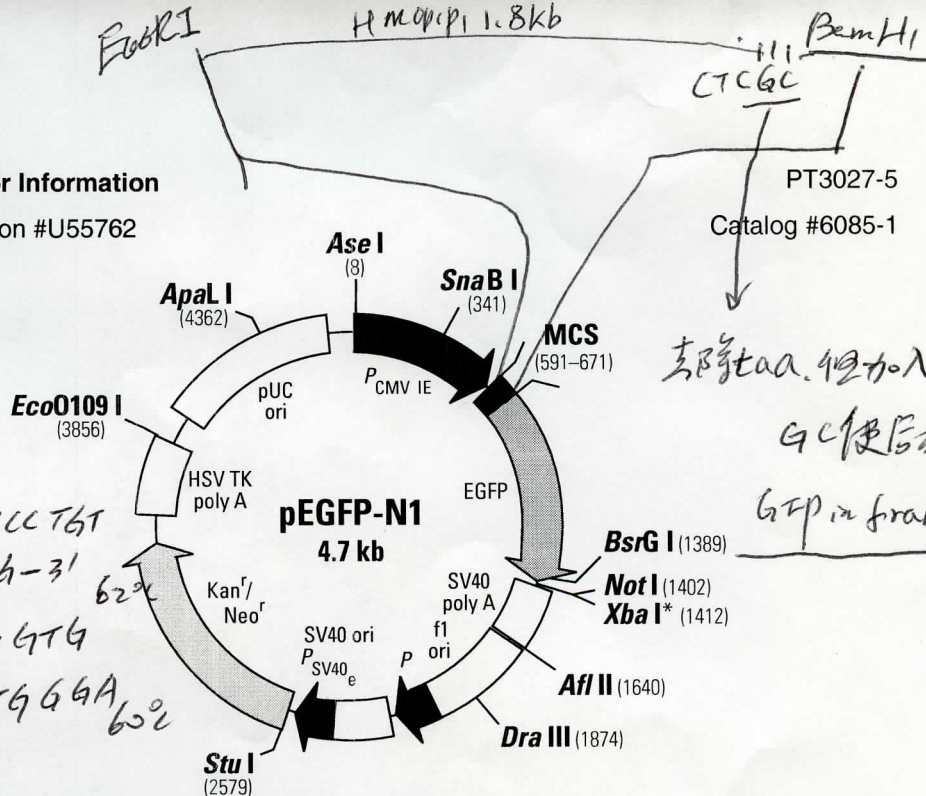


hMCP1-GFP

# pEGFP-N1 Vector Information GenBank Accession #U55762



5': 5'-CG **GAATTC** AA ATG AAT GGC CCG TGT  
3': 5'-CG **GAATTC** G C C T C A C T G G G G T G  
CTG GGA

591 601 611 621 631 641 651 661 671 EGFP  
G CTA GCG CTA CCG GAC TCA GAT CTC GAG CTC AAG CTT CGA ATT CTG CAG TCG ACG GTA CCG GCG GCC CCG GAT CCA CCG GTC GCC ACC ATG GTG  
NheI Eco47III BglII XhoI HindIII EcoRI PstI SalI AccI KpnI Asp718I SacII ApaI Bsp120I XmaI SmaI AgeI

**Restriction Map and Multiple Cloning Site (MCS) of pEGFP-N1 Vector.** (Unique restriction sites are in bold.) The *Not* I site follows the EGFP stop codon. The *Xba* I site (\*) is methylated in the DNA provided by CLONTECH. If you wish to digest the vector with this enzyme, you will need to transform the vector into a *dam*<sup>-</sup> and make fresh DNA.

## Description:

pEGFP-N1 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-N1 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-N1 is between the immediate early promoter of CMV ( $P_{CMV IE}$ ) and the EGFP coding sequences. Genes cloned into the MCS will be expressed as fusions to the N-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<sup>r</sup>), 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-N1 backbone also provides a pUC origin of replication for propagation in *E. coli* and an f1 origin for single-stranded DNA production.



Cla I                      Mlu I                      Xba I                      EcoR V                      Kpn I                      BamH I                      Bgl II  
 TCG ATA GAT ACG CGT CAT CTA GAT ATC GGT AOC GGA TCC AGA TCT  
 AGC TAT CTA TGC GGA GTA GAT CTA TAG OCA TGG CCT AGG TCT AGA

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