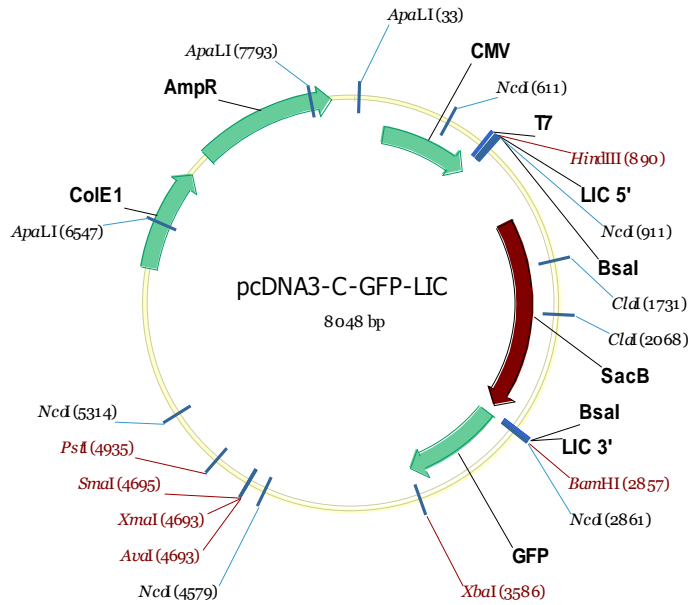


Vector information sheet

Dated: 8th May 2013

Vector Name	pCDNA3-C-GFP-LIC
Source	Grazyna Kochan
Sequence accession/link	(SGC)
Description	pCDNA3 eukaryotic expression vector with GFP tag as C-terminal fusion peptide. Includes sites for LIC cloning, and a "stuffer" fragment that includes the SacB gene, allowing negative selection on 5% sucrose
Antibiotic resistance	Ampicillin, 100 µg/ml
Promoter	CMV
Cloning	LIC. (vector treated with BsaI, then with T4 DNA polymerase in presence of dGTP)
Initiation codon	Supplied in PCR primer
C-terminal fusion – seq.	SKGGYGSMVSKGEELFTGVVPILVELDGDVNGHKFS VSGEGEDATYGKLTCLKFICTTGKLPVPWPTLVTTLT GVQCFSRYPDHMKQHDFFKSAMPEGYVQERTIFFK DGNKTRAEVKFEGDTLVNRIELKGIDFKEDGNILGHK LEYNYNSHNVYIMADKQKNGIKVNFKIRHNIEDGSVQL ADHYQQNTPIGDGPVLLPDNHYLSTQSALSKDPNEKR DHMVLLLEFVTAAGITLGMDELYK
C-terminal fusion – MW	27578.14 Da
Termination codons	Present in vector
Protease cleavage	no
Additional features	
Preferred host	Mammalian cell lines (HEK HeLa, BHK BSC 1 etc.)
5' sequencing primer	pcDNA3-fwd (50), and T7fwd. TCCAAAATGTCGTAACAACCTCC
3' sequencing primer	pcDNA3-rev (48): TTTTATTAGGAAAGGACAGTGG



Polylinker region:

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                                     T7
                                     ~~~~~~
CMV                                     LIC 5'
~~~~~                                     ~
                                     HindIII
                                     ~~~~~~
841 CTGCTTACTG GCTTATCGAA ATTAATACGA CTCACTATAG GGATACCCAA GCTTACCTGT
    GACGAATGAC CGAATAGCTT TAATTATGCT GAGTGATATC CCTATGGGTT CGAATGACA
                                     BsaI
                                     ~~~~~~
LIC 5'
~~~~~
NcoI
~~~~~
901 ACTTCCAATC CATGGAGACC GACGTCCACA TATACCTGCC GTTCACTATT ATTTAGTGAA
    TGAAGGTTAG GTACCTCTGG CTGCAGGTGT ATATGGACGG CAAGTGATAA TAAATCACTT
-----SacB-----
                                     BsaI
                                     ~~~~~~
LIC 3'                                     GFP
~~~~~                                     ~~~~~~
                                     NcoI
                                     ~~~~~~
                                     BamHI
                                     ~~~~~~
2821 TATTGGCATT GACGGTCTCC AGTAAAGGTG GATACGGATC CATGGTGAGC AAGGGCGAGG
     ATAACCGTAA CTGCCAGAGG TCATTTCCAC CTATGCCTAG GTACCACTCG TTCCCGCTCC
                                     GFP
     ~~~~~~
     · L F T G V V P I L V E L D G D V N G H K ·
2881 AGCTGTTCAC CGGGTGGTG CCCATCCTGG TCGAGCTGGA CGGCGACGTA AACGGCCACA
     TCGACAAGTG GCCCACCAC GGGTAGGACC AGCTCGACCT GCCGCTGCAT TTGCCGGTGT
                                     GFP
     ~~~~~~
     · F S V S G E G E G D A T Y G K L T L K F ·
2941 AGTTCAGCGT GTCCGGCGAG GCGAGGGCG ATGCCACCTA CGGCAAGCTG ACCCTGAAGT
     TCAAGTCGCA CAGGCCGCTC CCGTCCCGC TACGGTGGAT GCCGTTGCAC TGGGACTTCA
                                     GFP
     ~~~~~~
     · I C T T G K L P V P W P T L V T T L T Y ·

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3001 TCATCTGCAC CACCGGCAAG CTGCCCGTGC CCTGGCCAC CCTCGTGACC ACCCTGACCT
AGTAGACGTG GTGGCCGTTC GACGGGCACG GGACCGGGTG GGAGCACTGG TGGGACTGGA
      GFP
~~~~~
· G V Q C F S R Y P D H M K Q H D F F K S ·
3061 ACGGCGTGCA GTGCTTCAGC CGTACCCCG ACCACATGAA GCAGCACGAC TTCTTCAAGT
TGCCGCACGT CACGAAGTCG GCGATGGGGC TGGTGTACTT CGTCGTGCTG AAGAAGTTCA
      GFP
~~~~~
· A M P E G Y V Q E R T I F F K D D G N Y ·
3121 CCGCCATGCC CGAAGGCTAC GTCCAGGAGC GCACCATCTT CTTCAAGGAC GACGGCAACT
GGCGGTACGG GCTTCCGATG CAGGTCCTCG CGTGGTAGAA GAAGTTCCTG CTGCCGTTGA
      GFP
~~~~~
· K T R A E V K F E G D T L V N R I E L K ·
3181 ACAAGACCCG CGCCGAGGTG AAGTTCGAGG GCGACACCCT GGTGAACCGC ATCGAGCTGA
TGTTCTGGGC GCGGCTCCAC TTCAAGCTCC CGCTGTGGGA CCACTTGGCG TAGCTGCACT
      GFP
~~~~~
· G I D F K E D G N I L G H K L E Y N Y N ·
3241 AGGGCATCGA CTTCAAGGAG GACGGCAACA TCCTGGGGCA CAAGCTGGAG TACAACCTACA
TCCCGTAGCT GAAGTTCCTC CTGCCGTTGT AGGACCCCGT GTTCGACCTC ATGTTGATGT
      GFP
~~~~~
· S H N V Y I M A D K Q K N G I K V N F K ·
3301 ACAGCCACAA CGTCTATATC ATGGCCGACA AGCAGAAGAA CGGCATCAAG GTGAACTTCA
TGTCGGTGTG GCAGATATAG TACCGGCTGT TCGTCTTCTT GCCGTAGTTC CACTTGAAGT
      GFP
~~~~~
· I R H N I E D G S V Q L A D H Y Q Q N T ·
3361 AGATCCGCCA CAACATCGAG GACGGCAGCG TGCAGCTCGC CGACCACTAC CAGCAGAACA
TCTAGGCGGT GTTGTAGCTC CTGCCGTCGC ACGTCGAGCG GCTGGTGATG GTCGTCTTGT
      GFP
~~~~~
· P I G D G P V L L P D N H Y L S T Q S A ·
3421 CCCCCATCGG CGACGGCCCG GTGCTGCTGC CCGACAACCA CTACCTGAGC ACCCAGTCCG
GGGGGTAGCC GCTGCCGGGG CACGACGACG GGCTGTTGGT GATGGACTCG TGGGTGACGG
      GFP
~~~~~
· L S K D P N E K R D H M V L L E F V T A ·
3481 CCCTGAGCAA AGACCCCAAC GAGAAGCGCG ATCACATGGT CCTGCTGGAG TTCGTGACCG
GGGACTCGTT TCTGGGGTGT CTCTTCGCGC TAGTGTACCA GGACGACCTC AAGCACTGGC
      GFP
~~~~~
                                          XbaI
                                          ~~~~~~
· A G I T L G M D E L Y K * *
3541 CCGCCGGGAT CACTCTCGGC ATGGACGAGC TGTACAAGTA ATAATCTAGA GGGCCCTATT
GGCGGCCCTA GTGAGAGCCG TACCTGCTCG ACATGTTTAT TATTAGATCT CCCGGGATAA

```

Primers for LIC cloning:

Upstream: add TACTTCCAATCCATG to the 5' end (ATG in-frame with the desired coding sequence).

Downstream: add TATCCACCTTTACTGGA to 5' end of downstream primer;

pcDNA3-GFP-LIC sequence:

```

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