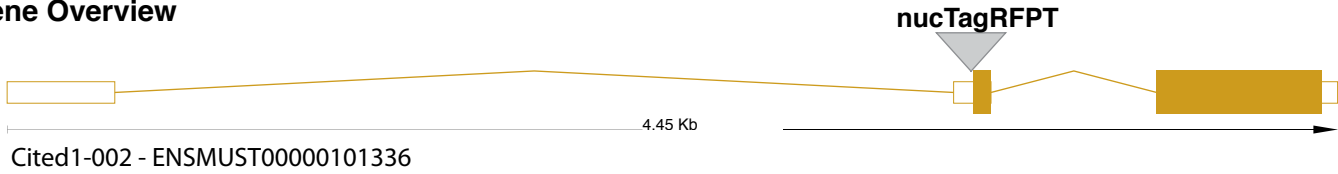


# Cited1-nucTagRFPT Construct Overview

Created 6 December 2010  
Updated na

## Gene Overview



## Design comments

There are three protein encoded transcripts reported for Cited1 that share a start ATG in the second exon. The ATG in Exon 2 was therefore targeted for insertion of the nucTagRFPT reporter cassette.

## Target site in cDNA

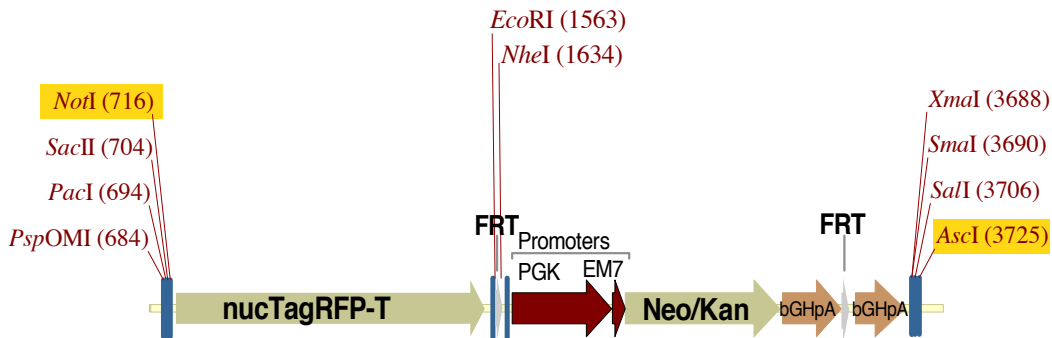
cDNA for Cited1-002  
Transcript length: 1090 bps  
Translation length: 203 residues

```

1  AGGGCTCTACGGGAGGCAGTGGGGCTTAAGAGCCCGGTCCCGCGCCGAGAACCAAGGG
61  TGGTGGCTGTGATCGCCACCCCGGAGTCCGGCCCGGAGCTCGGGCCGGCAAGAG
121  CGCGGGTCGGGGCAGCTCGGCGCTGCCCGGTGCGGTGCGCGCGCGCGGGCCGGCGGGG
181  GTGCAGGCTTTTCCAGCGCCCTGGGGTCGCGTGGCCCGCCGTGGGGCTGGGGACTCTG
241  AAGCAGTGGCTAGCGCGCGCGGGCAGTGTGCGTCCCAAGGGGGCTCGGGCTAGGTC
301  GCTTCGTCCGTACCTCAGCTCCTGTGAGCTTCTCTCCCGGCAAGCAGCTTCGGGAG
361  ACCAACAGGCCAGCTGGCGGCATCACTGCCACCGATTATCGGACTTCTGCCAGGGT
421  CTGAAATGCCAACTATGTCGAGGCCCTGCACCTGATGTCAGGGTGGCACCACCTTGGGA
481  AGGAGGATGCCAACCCAGGAGATGAACCTCTGGCCACTCCAACCTGGAGTGAAGGATC
541  GCAAGGCAGTGACTGTCTGCATACCCCGGGTCAACGCAATGGAGCCAAAGCCAAAG
601  GAGTTCACCTAGCTCCTTGGATCGACATCTCCAATAGGCTCTCTACTGCCACCCCTT
661  CTTCAAACCCCCATCCTTCAACCTGCATCTACCCCTCACCTGATGGCCAGCATGCAGC
721  TTCAGAAGCTTAATAGCCAGTACCAGGGGCTCGGCTACTGCTGCTGCTGCCCTACTG
781  GTGCAGGCTACCAGGGGAGGAAGGCCATGCAAAATGGGTACCAGCCCTCTGGTAG
841  TGGGAGGCTCCGGGATCTGTCTCTCCTCCTGCTGGTCCAGAGCCCTGCTCTCATTTG
901  ATTCAGACCCGGTGGATGAGGAGGTGCTGATGTCCTGGTGGTTGAATGGGGCTAGACC
961  GAGCCAATGAGCTTCCGAGCTGTGGTGGGGCAGAATGAGTTTGAATTTCACTGCAGATT
1021  TTCCCTTGGCTGTGATGGCAAGTGCCCTTAAAAAATGAGGAAAAAAGCCACCGATT
1081  CTGTTGTAAA
    
```

## Reporter Cassette

A 3xNLS modified TagRFP-T reporter was inserted at the ATG site of the Cited1 coding region selected. The Neo/Kan component is used for selection in bacteria and removed with transient expression of Flpe-recombinase prior to microinjection.



Fragment of pCZV-nucTagRFPT-FpNF-v2  
3163 bp (molecule 5358 bp)

# Cited1-nucTagRFPT Target Site Details

Created 4 December 2010

Updated 6 December 2010

## Endogenous Targeting Site

Left homology arm

tgtctctgct cttgggcca ggtgctctct gttttgagtt taactctatc agcatccttg tgtagctgaa tggattcacc cgcacttgga attttctagt  
acagagacga gaacccgggt ccacgagaga caaaactcaa attgagatag tcgtaggaac acatcgactt acctaagtgg gctgaaacct taaaagatca

Left homology arm

tctccctca cctttattat ttcttctctt ttacttgcag **ACCAACAGGC CCAGCTGGCG GCATCAACTG CCACCGATTT ATCGGACTTC TGCCAGGCT**  
agagggaaagt ggaataata aagaagagaa aatgaacgtc **TGGTTGTCCG GGTGCACCGC CGTAGTTGAC GGTGGCTAAA TAGCCTGAAG ACGGGTCCGA**

Left homology arm

Exon 2

**CTGAAATGCC AACTATGTCG AGGCCATGCAC TTGATGTCAA GGGTGGCACC ACCTCTGGGA AGGAGgtgag ttttatcatt ttacagatgg agatactaag**  
**GACTTACCG TTGATACAGC TCCGGACGTG AACTACAGTT CCCACCGTGG TGGAGACCTT TCCTCcactc aaaaatagtaa aatgtctacc tctatgattc**

Right homology arm

gcatggaag atggacttta ggaaggagg tcacgacact ggccttcta ggtgtggcct atagccgac tgcttgacca gagaagatac agcagggcct  
cgtaccttc tactgaaat cccttctcc agtgcgtgga ccggaagat ccacaccgga tatcggcgtg acgaaactggt ctcttctatg tctgcccgaa  
PR-Cited1

Right homology arm

## Targeted Site - 5'

Left homology arm

ttttgagttt aactctatca gcatcctgt gtagctgaat ggattcacc cgcacttgaa ttttctagt ctcccttca ctttattatt tcttctctt  
aaaactcaaa ttgagatagt cgtaggaaca catcgactta cctaagtggg gctgaaacct aaaagatcaa gaggaagtg gaaataataa agaagagaaa

Left homology arm

tacttgcagA **CCAACAGGC CAGCTGGCGG CATCAACTGC CACCGATTTA TCGGACTTCT GCCCAGGCTC TGAAGTTTGG CCGCCACCAT** **G**GTGCACGTG  
atgaacgctT **GGTTGTCCG GTCGACCGCC GTAGTTGACG GTGGCTAAT AGCTGAAGA CGGTCCGAG ACTTCAAACC GCGGTGTA** CCACGTGCAC

3XNLS-Tag-RFP

GATCCAAAAA AGAAGAGAAA GGTAGATCCA AAAAAGAAGA GAAAGGTAGA TCCAAAAAAG AAGAGAAAAG TACACGTGAG CATGGTGTCT AAGGGCGAAG  
CTAGGTTTTT TCTTCTTTT CCATCTAGGT TTTTCTTCT CTTTCCATCT AGGTTTTTTC TTCTCTTTCC ATGTGCACTC GTACCACAGA TTCCCCTTC

## Targeted Site - 3'

bGHpA

GGAAATTGCA TCGCATTGTC TGAGTAGGTG TCATTCCTATT CTGGGGGGTG GGGTGGGGCA GGACAGCAAG GGGGAGGATT GGGAAGACAA TAGCAGGCAT  
CCTTTAACGT AGCGTAACAG ACTCATCCAC AGTAAGATAA GACCCCCAC CCCACCCCGT CCTGTCTGTT CCCCTCCTAA CCCTTCTGTT ATCGTCCGTA

bGHpA

XmaI  
SmaI

XbaI

Sall

BssHII

GCTGGGGATG CCGTGGGCTC TATGGCCCGG GTGATCCTCT AGAGTCGACC TCTAGTGAAG TGGCGCGCGA **AATGCCAACT** ATGTCGAGGC CTGCACTTGA  
CGACCCTAC GCCACCCGAG ATACCGGGCC CACTAGGAGA TCTCAGCTGG AGATCACTCT ACCGCGCGCT **TTACGGTTGA TACAGCTCCG GACGTGAACT**

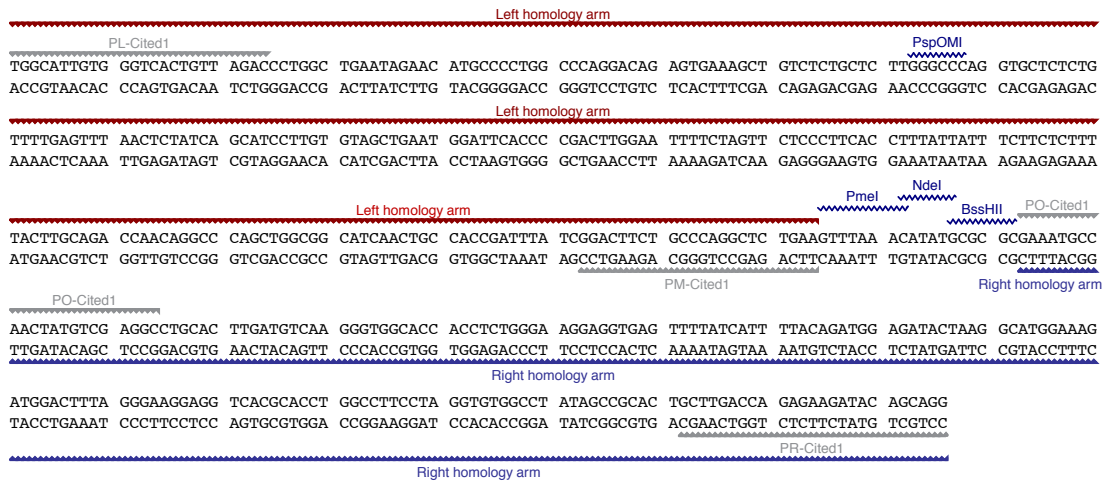
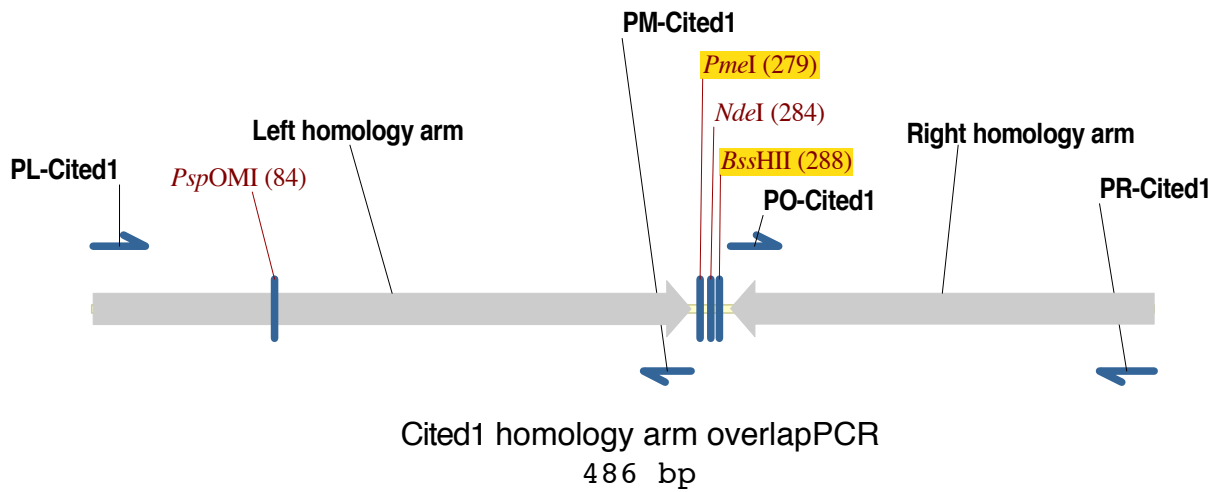
Right homology arm

TGTCAGGGT GGCACCACCT CTGGGAAGGA Ggtgagtttt atcattttac agatggagat actaaggcat ggaaagatgg actttagga aggaggtcac  
ACAGTCCCA CCGTGGTGA GACCCTTCT Ccactcaaaa tagtaaaatg tctacctcta tgattccgta cctttctacc tgaatccct tctccagtg

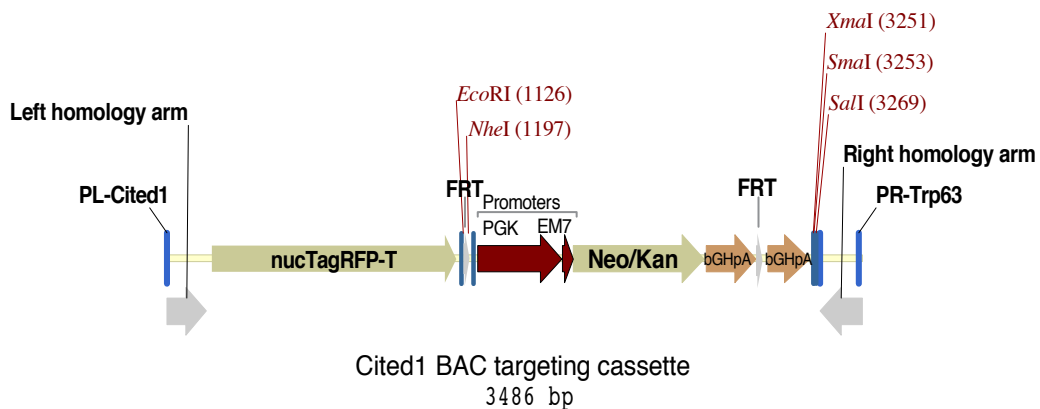
Right homology arm

## BAC targeting cassette for Cited1

The homologous arms for recombineering were created by PCR. The resulting product, cloned into a shuttle vector (not shown), contained the Left and Right homology arms. The nucTagRFPT reporter was cloned into the PmeI and BssHII sites of the homology arm PCT product. To accomplish this, the nucTagRFPT was digested with NotI (then blunted) and AscI (compatible overhang with BssHII).



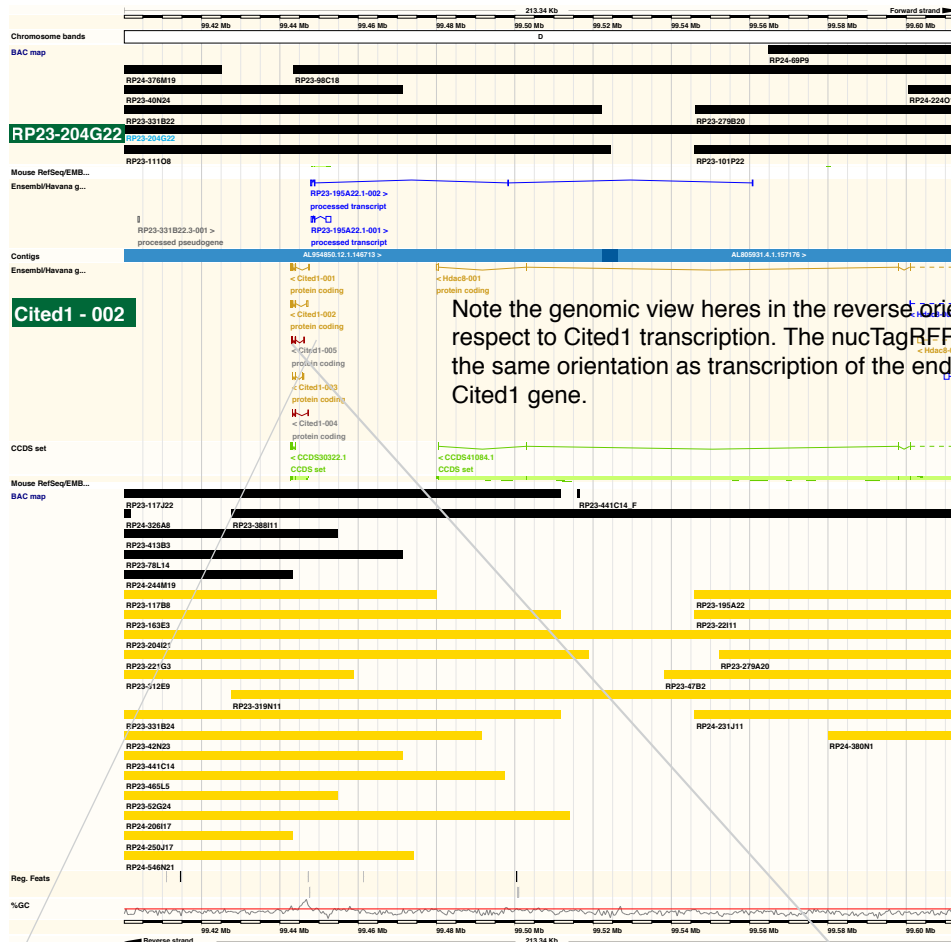
## Reporter + Arms



# Cited1-TagRFPT BAC Transgene

Created 2 December 2010  
 Updated 6 December 2010

BAC clone RP23-204G22 was targeted by recombineering with the Cited1-nucTagRFPT BAC targeting construct. The genomic context of Cited1 and the TagRFPT reporter is shown below. The BAC and the target gene are highlighted in dark green. Flanking primers are highlighted with yellow in the lower schematic.



Cited1-nucTagRFPT BAC construct  
 214565 bp