

# Gap1 integrative vector

**CURRENT STATUS:** POSTED

Dominique Loqué  
Carnegie Institution

Wolf Frommer  
Carnegie Institution

**DOI:**

10.1038/nprot.2007.128

**SUBJECT AREAS**

*Genetics*    *Molecular Biology*

**KEYWORDS**

*yeast, genomic integration, vector construction, Gap1*

## Introduction

Genes can be integrated into the genome of yeast. This protocol describes how to generate a vector that contains sequences that allow integration of any gene into the GAP1 locus of yeast.

## Procedure

- 1) Amplify a pUC fragment containing the bacterial origin of replication and the kanamycin resistance marker by PCR from PCR-BluntII-TOPO (Invitrogen, Carlsbad, CA) and ligate with a PvuII fragment of pDRf1 containing the f1 origin of replication, the PMA1 promoter and ADH3 terminator producing pDL001.
- 2) Subclone a synthetic oligonucleotide containing 55 bp upstream of the start (-55 to 0) and 55 bp downstream of the stop (+1810 to 1865) of Gap1 ORF with HpaI and AscI restriction sites into pGEM-T-easy.
- 3) Amplify the hphMX3 cassette (for hygromycin B selection in yeast) by PCR from pAG34<sup>1</sup> and clone into the SpeI site located in the 5'-part of the Gap1 cassette.
- 4) Amplify this cassette by PCR and clone into a blunted BglII of pDL001.
- 5) Subclone AtAMT1;1, AtAMT1;1-T460A and AtAMT1;1-eGFP from pDR vectors into the KpnI site of the Gap1 integrative plasmid.
- 6) Use the yeast strain 31019b ( $\Delta::LEU2 ::KanMX2 ura3$ ; see reference 2) to generate DL1 ( $\Delta \Delta$ ) mutant strain and the versions in which AtAMT1;1 or its mutants are integrated.
- 7) Transform yeast strain 31019b using a LiAc protocol<sup>(3)</sup> with an AscI linearized Gap1 integrative vector containing either AtAMT1;1, AtAMT1;1-T460A, AtAMT1;1-eGFP or the empty vector, to generate gap1::WT, gap1::T460A, gap1::AMT1;1-eGFP or  $\Delta gap1$  strains respectively.
- 8) Select transformants on solid YPD medium supplemented with 300  $\mu\text{g}/\text{mL}$  hygromycin.
- 9) Amplify colonies in liquid YPD and reselect on solid YPD supplemented with 300  $\mu\text{g}/\text{mL}$  hygromycin.
- 10) Confirm the insertion at the Gap1 locus by PCR and by complementation of functional proteins.

## References

- (1) Goldstein, A. L. & McCusker, J. M. Three new dominant drug resistance cassettes for gene disruption in *Saccharomyces cerevisiae*. *Yeast* **15**, 1541-1553 (1999).
- (2) Marini, A. M., Soussi-Boudekou, S., Vissers, S. & André, B. A family of ammonium transporters in

*Saccharomyces cerevisiae*. *Mol. Cell. Biol.* **17**, 4282-4293 (1997).

(3) Gietz, R. D. & Woods, R. A. Transformation of yeast by the LiAc/SS carrier DNA/PEG method.

*Methods Enzymol.* **350**, 87-96 (2002).

## A cytosolic trans-activation domain essential for ammonium uptake

by Loqué, D. et al.

Nature (20 December, 2006)