

# Benchmarks

## BENCHMARKS

Benchmarks are brief communications that describe helpful hints, shortcuts, techniques or substantive modifications of existing methods.

### Shuttle-Vector System for *Saccharomyces cerevisiae* Designed to Produce C-Terminal-Myc-Tagged Fusion Proteins

*BioTechniques* 25:936-938 (December 1998)

While the completion of whole genome sequence analysis of *Saccharomyces cerevisiae* has been achieved, the function of its 2300 genes are still unknown, and the genetic and biochemical analysis on these genes are ongoing (5). Determining the subcellular localization, co-immunoprecipitation analysis and the disruption or overexpression of unknown gene products provides us the basic information to understand the function of these genes. The addition of a commonly used epitope such as the decapeptide of the *myc* oncogene product, -EQKLISEEDL (Myc-tag) recognized by mouse monoclonal antibody (MAb) 9E10 (Oncogene Science, Cambridge, MA, USA), facilitates this process (2).

Here, we describe the new shuttle-vector system that produces a C-terminal-Myc-tagged fusion protein from a polymerase chain reaction (PCR)-amplified gene of interest. Figure 1A shows a schematic of vectors pAMT20 and pGMT10. The starting plasmid is pGBT9 (CLONTECH Laboratories, Palo Alto, CA, USA), a 2- $\mu$ m ori-type shuttle vector with a *TRP1* selection marker. After disrupting the *SphI* site at the hind part of the multiple cloning site (MCS), the *GAL4* open reading frame (ORF) of pGBT9 was removed, and the alternative MCS and the Myc-tag sequence (Figure 1B) from pPADHmyc-Ch, a derivative of pKT10 (7), were added to generate pAMT20. To generate pAMT10, the *ScaI-SphI* DNA fragment containing the 2- $\mu$ m ori of pAMT20 was replaced with the *ScaI-SphI* fragment of the *CEN6/ARSH4* cassette from pCEN6-118 containing a *ScaI-AluI* fragment of pRS413 (6). The *ScaI-EcoRI* (blunt-end) fragment of the *GAL1* promoter from pYX223 (R&D Systems Europe, Oxon, England, UK) was cloned into the *HincII* site of pUC118 (pGAL1-118). The *SphI-EcoRI* fragment of the *ADH1* promoter

Table 1. Characteristics of the Vectors

Marker	Name <sup>a</sup>	Size (kb)	Usable Sites <sup>b</sup>
<i>TRP1</i>	pAMT10	4.3	5
	pAMT20	5.1	5
	pGMT10	4.6	5
	pGMT20	5.4	5
<i>HIS3</i>	pAMH10	4.5	6
	pAMH20	5.3	6
	pGMH10	4.8	6
	pGMH20	5.6	6
<i>LEU2</i>	pAML10	5.1	6
	pAML20	5.9	6
	pGML10	5.4	6
	pGML20	6.2	6
<i>URA3</i>	pAMU10	4.6	6
	pAMU20	5.4	6
	pGMU10	4.9	6
	pGMU20	5.7	6

<sup>a</sup>All plasmids are obtainable from Riken DNA Bank (<http://www.rtc.riken.go.jp/DNA/HTML/engsearch.html>).

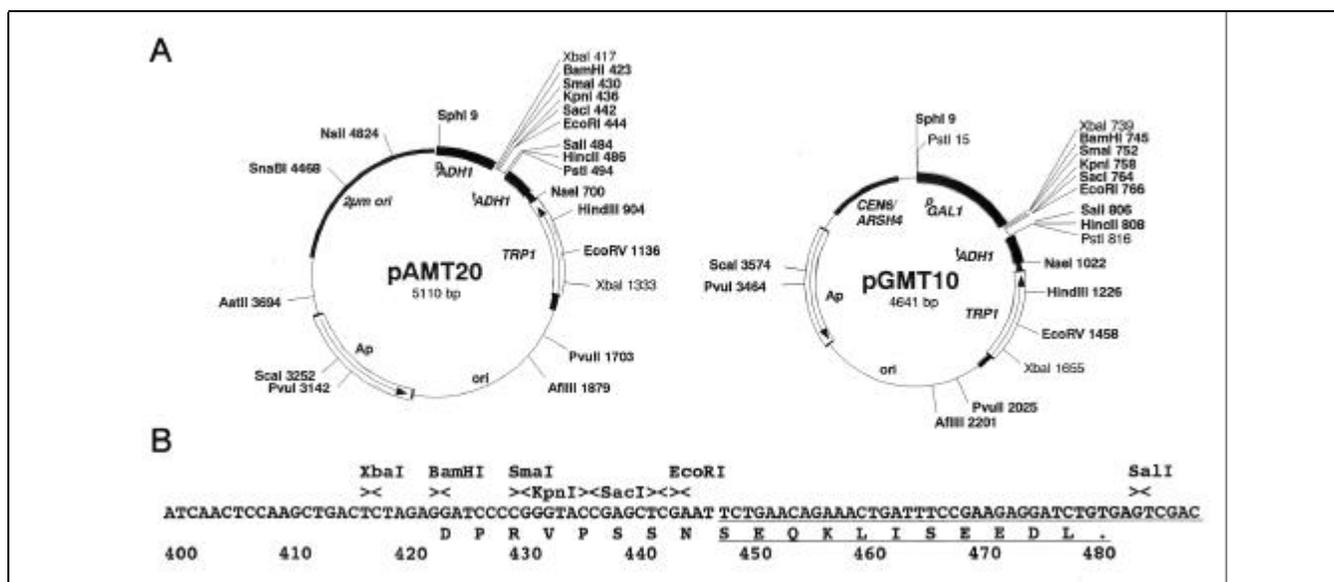
<sup>b</sup>Unique restriction site in the MCS in each vector was indicated.

**Table 2. Protocol for Immunoprecipitates' Preparation from Cell Lysates with Rabbit Polyclonal Anti-Myc-Tag Antibody**

1. After a 6–9-h induction in SG (0.67% Bacto-yeast nitrogen base without aa, 20–400 mg/mL aa mixture and adenine sulfate and uracil, 2% galactose), 5 mL culture of each strain carrying a Myc-tagging vector were washed once with distilled water.
2. An equal volume of zirconium beads were added to the cell pellets and then resuspended with 500  $\mu$ L of ice-cold sample buffer (10 mM Tris-HCl, pH 7.5, 150 mM NaCl, 1 mM EDTA, 1% Nonidet<sup>®</sup> P-40 (NP40), 0.1% sodium dodecyl sulfate (SDS), 0.1% deoxycholic acid, 1 mM benzamidine).
3. Cell suspension was crushed with Mini-BeadBeater<sup>®</sup> (Biospec Products, Bartlesville, OK, USA) for 5 min at 4°C.
4. The crushed-cell mixture was centrifuged at 12 000 $\times$  g for 10 min at 4°C, and then the supernatant (cell lysate) was transferred to a newly prepared tube.
5. Fifty micrograms of rabbit polyclonal antibody against Myc-tag (MBL) conjugated with protein-A Sepharose<sup>®</sup> (Sigma-Aldrich Chemical, Tokyo, Japan), were added to each cell lysate and incubated for overnight at 4°C.
6. The Sepharose beads were washed 3 $\times$  with ice-cold sample buffer.
7. The precipitates were resuspended with 10  $\mu$ L of 1 $\times$  SDS-polyacrylamide gel electrophoresis (PAGE) loading solution [65 mM Tris-HCl, pH 6.8, 2% SDS, 10% glycerol, 0.01% phenol red, 42 mM dithiothreitol (DTT)] and boiled for 3 min.
8. The prepared samples were loaded on 12% SDS-PAGE gel, run for the appropriate separation time and then transferred to Hybond<sup>™</sup> ECL nitrocellulose membrane (Amersham Pharmacia Biotech, Little Chalfont, Bucks, England, UK).
9. Detection of the immunoprecipitated protein was done with ECL<sup>™</sup> Western Blotting Detection Reagents (Amersham Pharmacia Biotech) using mouse MAb 9E10 or mouse anti-Gts1p polyclonal antibody as the primary antibody and peroxidase-conjugated rabbit anti-mouse IgG as the secondary antibody, respectively.
10. The photo images of X-ray films in Figure 2 were processed with Adobe<sup>®</sup> Photoshop<sup>®</sup> Version 3.0 (Adobe Systems, San Jose, CA, USA).

and the MCS of pAMT10 or pAMT20 were replaced with the *SphI-EcoRI* fragment of the *GAL1* promoter and the MCS of pGAL1-118 to generate pGMT10 or pGMT20. Five restriction sites of the MCS are usable in the pA/GMT series, and the *XbaI* site is added in other series (Figure 1B). For construction of other series having *HIS3*, *LEU2* and *URA3*, DNA fragments with selection markers were cloned at the *PvuII-HindIII* site of the pA/GMT vectors. Table 1 summarizes the characteristics of each vector. Numbers 10 and 20 at the end of each vector's name indicates the type of replication origin as *CEN6/ARS4* and 2- $\mu$ m ori, respectively.

Using this vector system, the structural analysis was performed on the two gene products involved in flocculation machinery in *S. cerevisiae*. The primary structure of the *GTS1/LSR1* (1,4) gene product has been controversial, because it has been reported that the ORF of *LSR1* is equal to that of *GTS1*, except for the single T deletion 990 bases from the start codon of its ORF (1). Therefore, we assessed the primary structure of Gts1p (or Lsr1p) again. GDU2168, BJ2168 (9) with *gts1::URA3*, was transformed with pGML20-GTS1 and pGML20-LSR1. Both were designed to attach the Myc-tag in their C terminus if the DNA sequence of the ORF is correct, and the whole-cell lysate from each



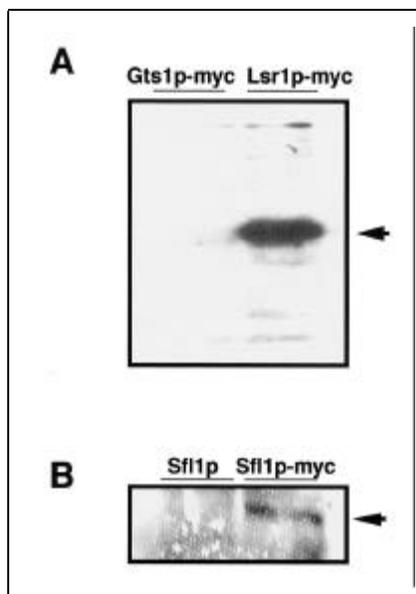
**Figure 1.** (A) Schematic representation of the pAMT20 and pGMT10 vectors. Bold characters indicate unique restriction sites. (B) The relevant DNA and aa sequence around MCS of pAMT20 is shown. The Myc-tag sequences are indicated with underline.

# Benchmarks

transformant was applied to immunoprecipitation and Western blot analysis. The anti-Gts1p antibody against both immunoprecipitates recognized the Lsr1p-type only (Figure 2A). Amino acid (aa) composition analysis, on the purified Gts1p expressed in *E. coli*, also indicated that the Lsr1p-type composition is more likely than that of the Gts1p-type (8). Thus, we concluded that the primary structure of the product from *GTS1/LSR1* is the Lsr1p-type. Another protein involved in flocculation machinery, Sfl1p (3), was also tagged with Myc and detected by Myc-tag-specific 9E10 MAb (Figure 2B). However, the specific signal detected by 9E10 was much weaker than that of the anti-Gts1p antibody because of the low titration performance with this MAb or the labile

nature of Sfl1p. Functional dissection of Gts1p (i.e., Lsr1p) has shown that the C-terminal half plays an essential role for flocculation (data not shown). Though the Myc-tagged Lsr1p showed fully functional flocculation property compared to that of intact Lsr1p, and the Myc-tagged Sfl1p showed the same results (data not shown). In this case, these two proteins were functional whether or not the Myc-tag is connected in their C terminus.

In this report, we describe the construction of a novel small epitope (Myc)-tagging *E. coli*-*S. cerevisiae* shuttle-vector system that is useful in the following ways: (i) relatively small-sized (4.3–6.2 kb) as a shuttle vector for yeast; (ii) five to six unique restriction enzyme sites in its MCS; (iii) four different selective markers (*TRP1*, *HIS3*, *LEU2* and *URA3*); and (iv) two types of promoter (*ADH1* and *GAL1*–10). It is hoped that this system will be a convenient tool for yeast researchers by eliminating the time-consuming antiserum generation.



**Figure 2. Re-evaluation of the primary structure of Gts1p and detection of Myc-tagged Sfl1p.** (A) Western-blot analysis with polyclonal anti-Gts1p antibody. Arrowhead indicates polypeptides recognized with anti-Gts1p antibody (4). (B) Western-blot analysis with 9E10 MAb antibody against Myc-tag. Immunoprecipitates from cell lysates of GDU2168, carrying pGML20-GTS1 (Gts1p-Myc) and pGMT20-LSR1 (Lsr1p-Myc), and SDU2168, carrying YEp51-SFL1 (Sfl1p) and pGMT20-SFL1 (Sfl1p-Myc), were prepared with rabbit polyclonal anti-Myc-tag antibody (MBL, Tokyo, Japan) (see Table 2).

## REFERENCES

1. **Bossier, P., P. Goethals and C. Rodrigues-Pousada.** 1997. Constitutive flocculation in *Saccharomyces cerevisiae* through overexpression of the *GTS1* gene, coding for a 'Glo'-type Zn-finger-containing protein. *Yeast* 13:717-725.
2. **Evan, G.I., G.K. Lewis, G. Ramsay and J.M. Bishop.** 1985. Isolation of monoclonal antibodies specific for human c-myc proto-oncogene product. *Mol. Cell. Biol.* 5:3610-3616.
3. **Fujita, A., Y. Kikuchi, S. Kuhara, Y. Misumi, S. Matsumoto and H. Kobayashi.** 1989. Domains of the SFL1 protein of yeast are homologous to Myc oncoproteins or yeast heat-shock transcription factor. *Gene* 85:321-328.
4. **Mitsui, K., S. Yaguchi and K. Tsurugi.** 1994. The *GTS1* gene, which contains a Gly-Thr repeat, affects the timing of budding and cell size of the yeast *Saccharomyces cerevisiae*. *Mol. Cell. Biol.* 14:5569-5578.
5. **Oliver, S.** 1996. A new approach to the systematic analysis of yeast gene function. *Trends Genet.* 12:241-242.
6. **Sikorski, R.S. and P. Hieter.** 1989. A system of shuttle vectors and yeast host strains designed for efficient manipulation of DNA in *Saccharomyces cerevisiae*. *Genetics* 122:19-27.
7. **Tanaka, K., K. Matsumoto and A. Toh-e.** 1988. Dual regulation of the expression of the polyubiquitin gene by cyclic AMP and heat shock in yeast. *EMBO J.* 7:495-502.
8. **Yaguchi, S., K. Mitsui and K. Tsurugi.** 1997. Reexamination of the nucleotide sequence of *GTS1*, a candidate clock-related gene of *Saccharomyces cerevisiae*. *Biochem. Arch.* 13:97-105.
9. **Zubenko, G.S., A.P. Mitchell and E.W. Jones.** 1980. Mapping of the proteinase b structural gene *PRB1*, in *Saccharomyces cerevisiae* and identification of nonsense alleles within the locus. *Genetics* 96:137-146.

We thank Akio Toh-e for providing plasmid pKT10, Tsutomu Ohta for providing yeast strain BJ2168 and So-ichi Yaguchi for technical support for the immunological experiment. Address correspondence to Dr. Hidekatsu Iha, Department of Biochemistry, Yamanashi Medical University, 1110 Shimokato, Tamaho, Nakakoma, Yamanashi 409-3898, Japan. Internet: hiha@swallow.res.yamanashi-med.ac.jp

Received 15 June 1998; accepted 24 August 1998.

**Hidekatsu Iha and Kunio Tsurugi**  
Yamanashi Medical University  
Yamanashi, Japan