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2

3 **Title:** Visualization of the synergistic effect of lithium acetate and single-stranded  
4 carrier DNA on *Saccharomyces cerevisiae* transformation

5

6 **Short title:** Visualization of the synergistic effect

7

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17

## 1     **Abstract**

2           Transformation is an indispensable method for the genetic manipulation of cells.  
3     *Saccharomyces cerevisiae* can be transformed by incubating intact cells and plasmid  
4     DNA in the presence of polyethylene glycol alone. Lithium acetate and single-stranded  
5     carrier DNA enhance the transformation efficiency, but the mechanism underlying this  
6     enhancement has remained elusive. In this study, we first confirmed that lithium acetate  
7     and single-stranded carrier DNA synergistically improve the transformation efficiency  
8     of *S. cerevisiae* intact cells. We then used transmission electron microscopy to observe  
9     the cell walls of yeast incubated with both lithium acetate and single-stranded carrier  
10    DNA in the presence of negatively charged Nanogold (in this context, a mimic of DNA).  
11    Under these conditions, the cell walls exhibited protruded, loose, and porous structures.  
12    The Nanogold was observed within the cell wall, rather than on the surface. We also  
13    made observations using YOYO-1, a fluorescent DNA probe. Based on the transmission  
14    electron microscopy and fluorescence data, we speculated that single-stranded carrier  
15    DNA covers the whole cell and enters, at least partially, into the cell wall structure,  
16    causing the cell wall to become protruded, loose, and porous; meanwhile, LiAc gives  
17    effect on the cell wall. Together, the two compounds synergistically enhance  
18    transformation efficiency and frequency.

19  
20    Key words: *Saccharomyces cerevisiae*, transformation, lithium acetate, single-stranded  
21    carrier DNA, transmission electron microscopy, Nanogold.

# 1 **Introduction**

2

3 Transformation, in which exogenous plasmid DNA is introduced into cells, is an  
4 indispensable technique for genetic modification. In *Saccharomyces cerevisiae*, the  
5 spheroplast transformation method was established in 1978 (Hinnen, et al. 1978). Later,  
6 the lithium method for transforming intact *S. cerevisiae* cells was developed (Ito, et al.  
7 1983). In the lithium method, monovalent cations (generally lithium, but also rubidium),  
8 enhance the transformation efficiency (the number of transformants per  $\mu\text{g}$  plasmid  
9 DNA) of intact *S. cerevisiae* cells. Ito et al. described the effect of transformation  
10 reagents in detail, demonstrating (i) incubation of intact cells with polyethylene glycol  
11 (PEG) and plasmid DNA is essential for transformation; (ii) short-term incubation of  
12 intact cells with PEG and plasmid DNA at 42°C (heat shock) enhances the  
13 transformation efficiency; (iii) transformation of the cells is most effective at the  
14 mid-log phase; and (iv) lithium acetate (LiAc) is the most effective monovalent cation  
15 (Ito, et al. 1983). Subsequently, Gietz et al. modified the original lithium method and  
16 succeeded in improving its efficiency by immediately mixing washed intact cells with  
17 PEG, LiAc, plasmid DNA, and single-stranded carrier DNA (ssDNA), and incubating  
18 them at 42°C for 40–60 min (Gietz, et al. 1992, Gietz, et al. 1995, Gietz and Woods  
19 2002, Schiestl and Gietz 1989). The modified lithium method has been referred to as the  
20 LiAc/ssDNA/PEG method (Gietz and Woods 2002). Intact cells can be transformed  
21 without the addition of LiAc by incubating the cells with PEG and plasmid DNA at  
22 30°C and then heat-shocking at 42°C (heat shock) (Hayama, et al. 2002, Yamakawa, et  
23 al. 1985). From these results, we can conclude that PEG is required for transformation,  
24 while LiAc and ssDNA both serve to enhance transformation efficiency. The

1 mechanism underlying this enhancement has remained elusive.

2 LiAc increases the permeability of intact cells to nucleic acid (Brzobohaty and  
3 Kovac 1986) and YOYO-1 (Zheng, et al. 2005) but has no effect on the transformation  
4 frequency (the transformation efficiency per viable cell) of spheroplasts (Chen, et al.  
5 2008). Furthermore, Chen et al. observed by scanning electron microscopy that the  
6 surfaces of intact cells incubated with LiAc at 30°C for 30 min became much rougher  
7 and more wrinkled, whereas the surfaces of intact cells incubated without LiAc as well  
8 as those of spheroplasts remained homogenous and smooth (Chen, et al. 2008). Thus, it  
9 has been proposed that LiAc helps plasmid DNA to pass through the cell wall, but not  
10 the plasma membrane (Chen, et al. 2008, Kawai, et al. 2010). However, it should be  
11 stressed that the aforementioned studies examined the role of LiAc alone, in the absence  
12 of ssDNA. Previous studies reported that ssDNA had no effect on the transformation  
13 efficiency of spheroplasts (Schiestl and Gietz 1989), leaving open the possibility that  
14 ssDNA could have an effect on the cell wall in the presence of LiAc.

15 In this study, we focused initially on the effect of ssDNA alone on transformation  
16 efficiency; our results revealed that ssDNA enhanced transformation, albeit less  
17 efficiently than LiAc alone. Subsequently, we shifted focus to the synergistic effect of  
18 LiAc and ssDNA together. Furthermore, we visualized the synergistic effects of LiAc  
19 and ssDNA on the cell wall using transmission electron microscopy (TEM).

20

1 **Materials and methods**

2

3 **Strains**

4

5 *S. cerevisiae* BY4742 (*MAT $\alpha$  leu2 $\Delta$ 0 lys2 $\Delta$ 0 ura3 $\Delta$ 0 his3 $\Delta$ 1*) was purchased from  
6 EUROSCARF and cultivated under standard yeast growth conditions (Sherman 2002).

7

8 **Transformation**

9

10 Transformation of intact *S. cerevisiae* cells was performed as described (Gietz and  
11 Woods 2002), with slight modifications. In brief, for each transformation reaction, *S.*  
12 *cerevisiae* cells were collected from 5.5 ml yeast–peptone–dextrose (YPD) culture  
13 (Sherman 2002) during log phase (approximate  $A_{600}$  of 0.5). The cells were washed  
14 once with sterilized distilled water (SDW) and incubated at 42°C for 20 min in 42  $\mu$ l  
15 suspension consisting of 2.0  $\mu$ l plasmid DNA [pRS415 (Sikorski and Hieter 1989); 0.10  
16  $\mu$ g/ $\mu$ l], 30  $\mu$ l 50% (w/v) PEG (P3640; Sigma, St. Louis, MO), 4.5  $\mu$ l 1.0 M LiAc, and  
17 6.0  $\mu$ l ssDNA (2.0 mg/ml; boiled and cooled salmon testis DNA, D1626, Sigma). Final  
18 concentrations of reagents in the suspension were 4.76  $\mu$ g/ml pRS415, 36% PEG, 10.7  
19 mM LiAc, and 0.29 mg/ml ssDNA. To examine the individual contributions from LiAc  
20 and ssDNA, LiAc (1.0 M, 4.5  $\mu$ l) was replaced with rubidium acetate (RbAc) (1.0 M,  
21 4.5  $\mu$ l) or SDW (4.5  $\mu$ l), and ssDNA (6.0  $\mu$ l) was replaced with SDW (6.0  $\mu$ l). After  
22 incubation, cells were resuspended in 1.0 ml SDW. After 10  $\mu$ l of the suspension was  
23 removed and diluted in SDW, the cells in the remaining suspension were collected and  
24 spread onto selective solid synthetic complete (SC) medium without leucine (Sherman

1 2002). Viable cells were counted by spreading the diluted suspension on YPD solid  
2 medium.

3 In this article, transformation efficiency is defined as the number of transformants  
4 per  $\mu\text{g}$  plasmid DNA (pRS415), while transformation frequency is defined as the  
5 transformation efficiency divided by the number of viable cells.

6

### 7 Transmission electron microscopy (TEM)

8

9 Cells were subjected to the transformation procedure (Gietz and Woods 2002) as above,  
10 but were incubated in 44  $\mu\text{l}$  suspension containing 4.0  $\mu\text{l}$  of negatively charged  
11 Nanogold (100 pmol/ $\mu\text{l}$  in ultra pure water, 1.4 nm in diameter; Nanoprobes, Yaphank,  
12 NY) instead of 2.0  $\mu\text{l}$  pRS415. Cells were collected by centrifugation, and the  
13 supernatant was removed completely. Next, cells were resuspended in 2%  
14 paraformaldehyde and 2% glutaraldehyde, fixed overnight, frozen in liquid propane  
15 using copper disks (Maxtaform Grid II HF51, Nisshin EM), substituted in 2% osmium  
16 tetroxide in acetone plus 3% water, dehydrated, and embedded in Quetol-651 (Nisshin  
17 EM) (Baba 2008). Ultra-thin sections (approximately 80 nm thick) were cut with a  
18 diamond knife on an LKB2088 ultramicrotome V (LKB-Produkter AB, Stockholm,  
19 Sweden), treated with GoldEnhance (Nanoprobes) for 1 min at room temperature, and  
20 stained with 2% aqueous uranyl acetate followed by lead staining (Sigma). The sections  
21 were examined using a JEM-1200EX microscope (JEOL, Tokyo, Japan) at 80 kV.

22

### 23 Fluorescence microscopic observation

24

1 Fluorescence microscopy was performed using a BX51 system (Olympus, Tokyo,  
2 Japan) equipped with a CoolSNAP camera (Quantitative Imaging Corporation, Burnaby,  
3 Canada) and filter (U-MNIBA2). YEp13 (Broach, et al. 1979) was labelled with  
4 YOYO-1 as described (Pham, et al. 2011). Transformation of *S. cerevisiae* cells was  
5 conducted by incubating 42  $\mu$ l of cell suspension at 42°C for 20 min as above, but  
6 substituting 2.0  $\mu$ l 1.0  $\mu$ M YOYO-1 alone or 2.0  $\mu$ l YOYO-1-labelled YEp13 (90  $\mu$ g  
7 [12.7 pmol] in 1.0  $\mu$ M YOYO-1) for 2.0  $\mu$ l pRS415.

8

## 9 **Results and discussion**

10

### 11 Synergistic effect of LiAc and ssDNA

12

13 To understand the individual contributions of LiAc and ssDNA, and of both compounds  
14 in combination, we compared the transformation efficiency and frequency obtained by  
15 incubating cells at 42°C for 20 min in the presence of PEG alone, PEG plus ssDNA,  
16 PEG plus LiAc, PEG plus ssDNA and LiAc, or PEG plus ssDNA and RbAc (Table 1).  
17 As shown in Table 1, we observed that ssDNA alone could enhance transformation  
18 efficiency and frequency by 7- and 6-fold, respectively; addition of LiAc increased  
19 these values by 42- and 60-fold. When used together, ssDNA and LiAc improved  
20 efficiency and frequency by 501- and 560-fold, i.e., they had a synergistic effect on both  
21 efficiency and frequency. ssDNA and RbAc in combination also displayed a synergistic  
22 effect (enhancement by 276- and 321-fold; Table 1).

23

24 Visualization of the synergistic effect of ssDNA and LiAc in combination

1

2 To clarify the mechanism underlying the synergistic effect of ssDNA and LiAc, cells  
3 were subjected to same transformation procedure used to generate Table 1, in the  
4 presence of 400 pmol of negatively charged Nanogold instead of plasmid DNA  
5 (pRS415). Negatively charged Nanogold consists of gold particles that carry a negative  
6 charge; it has been previously used as a mimic for plasmid DNA in studies of  
7 transformation (Pham, et al. 2011). Immediately after incubation at 42°C for 20 min,  
8 cells were fixed and observed using TEM. The signal from the Nanogold was amplified  
9 using GoldEnhance.

10 Cells incubated with PEG alone exhibited cell walls with a smooth surface (Fig.  
11 1A), as did cells that did not receive any treatment (data not shown). In contrast, cells  
12 incubated with PEG and either ssDNA or LiAc had cell walls with slight protrusions  
13 (Fig. 1B,C). Notably, incubation of the cells with PEG, ssDNA and either LiAc or RbAc  
14 (conditions that gave the highest efficiency; Table 1), caused the cell wall structures to  
15 be the most extremely protruded, loose, and porous (Fig. 2A,B). In addition, the porous  
16 cell wall appeared to be partially removed from plasma membrane (Fig. 2A,B). Thus,  
17 taking these data together with the absence of an effect of ssDNA and LiAc on the  
18 transformation efficiency of spheroplasts (Schiestl and Gietz 1989), we concluded that  
19 the target of both ssDNA and LiAc/RbAc is the cell wall, and that the function of these  
20 compounds is to cause the structure of the cell wall to become protruded, loose, and  
21 porous (Fig. 2A,B).

22 The modification of the cell wall by LiAc alone (Fig. 1C) was consistent with  
23 previous reports demonstrating that LiAc alone can modify the structure of the cell wall  
24 and increase permeability to nucleic acid or YOYO-1 (Brzobohaty and Kovac 1986,



1 Chen, et al. 2008, Zheng, et al. 2005). Previously, we used TEM to observe intact *S.*  
2 *cerevisiae* cells incubated with Nanogold and PEG in the absence of LiAc and ssDNA  
3 (Pham, et al. 2011). In this study, we used TEM to visualize for the first time the  
4 synergistic effect caused by both ssDNA and LiAc/RbAc.

5 Negatively charged Nanogold was observed as a dot-like signal within the cell wall;  
6 Nanogold in the intracellular space was always associated with a membrane structure,  
7 i.e. Nanogold was found at the periphery of the structures within the yeast cell (Figs. 1,  
8 2), as observed previously (Pham, et al. 2011). These observations support the idea that  
9 Nanogold enters cells via membrane invagination (Kawai, et al. 2004). It should be  
10 noted that the Nanogold appeared to be partially trapped in the pores formed in the cell  
11 wall in response to ssDNA and LiAc/RbAc (Fig. 2A,B, arrows). This led us speculate  
12 that such a binding mode, in which plasmid DNA is similarly trapped in cell wall pores,  
13 is at least one of the factors explaining the synergistic effect of ssDNA and LiAc/RbAc  
14 on transformation efficiency and frequency.

15

16 Behavior of plasmid DNA and ssDNA

17

18 To determine how plasmid DNA and ssDNA behave during transformation, we used  
19 YOYO-1. YOYO-1 is a widely used cell-impermeable fluorescent DNA probe (Gurrieri,  
20 et al. 1997); intercalation of YOYO-1 into DNA increases the probe's fluorescence  
21 intensity by more than 1,000-fold (Rye, et al. 1992). YOYO-1 has been successfully  
22 used to observe the behavior of plasmid DNA during transformation (Chen, et al. 2008,  
23 Pham, et al. 2011, Zheng, et al. 2005). However, no attempt has been made to observe  
24 the behavior of ssDNA using YOYO-1.

1 We incubated cells with PEG, ssDNA, LiAc, and YOYO-1 in the absence of plasmid  
2 DNA at 42°C for 20 min. The whole cell surface was stained with YOYO-1 (Fig. 3A).  
3 When the cells were incubated with PEG, LiAc, and YOYO-1 in the absence of any  
4 DNA, no fluorescent signal was observed (data not shown). Collectively, the fluorescent  
5 signal on the whole cell surface (Fig. 3A) indicated that ssDNA bound to the whole cell  
6 wall (Fig. 3A). Such binding is compatible with a previous report that detected  
7 cell-associated DNA using the diphenylamine assay (Gietz, et al. 1995). Moreover,  
8 when cells were incubated with PEG, ssDNA, and YOYO-1-labelled plasmid DNA, the  
9 whole cell surface was still stained by YOYO-1 irrespective of the presence or absence  
10 of LiAc (Fig. 3B,C) as in Fig. 3A. Thus, fluorescent microscopic observation using  
11 YOYO-1 could neither discriminate plasmid DNA from ssDNA nor visualize the  
12 synergistic effect of ssDNA and LiAc. This emphasizes the advantages of TEM analysis  
13 in elucidating the mechanism of transformation.

14 Negatively charged Nanogold was observed in the cell wall irrespective of the  
15 presence or absence of ssDNA (Figs. 1, 2), suggesting that even when it is bound to the  
16 whole cell wall (Fig. 3A), ssDNA would not inhibit plasmid DNA from binding to the  
17 cell wall. The binding of plasmid DNA to the cell wall is important for transformation,  
18 because it is the bound plasmid DNA, but not plasmid DNA in solution, that is proposed  
19 to enter the cells during transformation (Pham, et al. 2011). Under conditions that  
20 caused higher amounts of the plasmid DNA to be absorbed on the cell wall,  
21 transformation efficiency and frequency were both higher (Pham, et al. 2011). However,  
22 it had been previously proposed that plasmid DNA in solution enters the cell during  
23 transformation (Gietz, et al. 1995, Gietz and Woods 2001). Based on this proposal, it  
24 has also been supposed that the role of ssDNA or double-stranded carrier DNA is to

1 saturate the DNA-binding sites on the cell wall and to increase the probability of uptake  
2 of plasmid DNA from solution (Gietz, et al. 1995, Gietz and Woods 2001). Contrary to  
3 these hypotheses, our previous results (Pham, et al. 2011), taken together with our  
4 results obtained from TEM analysis (Figs. 1, 2) and microscopic observation (Fig. 3),  
5 clearly indicate that although ssDNA binds the cell surface, ssDNA and LiAc serve  
6 primarily to modify the structure of the cell wall itself. This is in contrast to a  
7 mechanism in which ssDNA and LiAc serve to saturate the DNA-binding sites of cell  
8 wall in order to increase the probability of uptake of plasmid DNA from solution. Thus,  
9 this study clearly demonstrates a new role for ssDNA, based on our previous reports  
10 (Pham, et al. 2011).

11 Nanogold was found in cell wall structures, but not on the surface of the cell wall  
12 (Figs. 1, 2), suggesting that both plasmid DNA and ssDNA can at least partially enter  
13 into the cell wall, and that ssDNA causes the cell wall to become protruded, loose, and  
14 porous. Moreover, TEM analysis conducted after removal of PEG detected Nanogold in  
15 the cells, but not on the surface of the cell wall, even in the absence of LiAc and ssDNA  
16 (Fig. 1A). This would be in good agreement with previous reports by Bruschi et al.  
17 (Bruschi, et al. 1987). In that study, the authors transformed intact *S. cerevisiae* cells by  
18 incubating the cells with plasmid DNA in the presence of PEG alone, and demonstrated  
19 that plasmid DNA was DNase-resistant after PEG was removed, but DNase-sensitive in  
20 the presence of PEG (Bruschi, et al. 1987). We ascribe the DNase-resistance of plasmid  
21 DNA to the probable location of the DNA within the structures of the cell wall, as  
22 observed for Nanogold (Fig. 1).

23

24

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2

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11

1 **Figure legends**

2

3 **Fig. 1.** Visualizing the effects of ssDNA and LiAc. Cells were incubated at 42°C for 20  
4 min in the presence of negatively charged Nanogold, with PEG (A), PEG plus ssDNA  
5 (B), or PEG plus LiAc (C), treated, and observed by TEM. Left panel indicates the  
6 image at 12,500-fold magnification; scale bar is 2  $\mu\text{m}$ . A portion of the left panel image  
7 is further magnified and shown in the right panel (47,800-fold magnification; scale bar  
8 is 0.50  $\mu\text{m}$ ). Signals from Nanogold were enhanced using GoldEnhance, and are  
9 observed as dots.

10

11 **Fig. 2.** Visualization of the synergistic effect of ssDNA and either LiAc or RbAc. Cells  
12 were incubated at 42°C for 20 min in the presence of negatively charged Nanogold,  
13 with PEG plus ssDNA and LiAc (A) or PEG plus ssDNA and RbAc (B), treated, and  
14 observed by TEM. Images are displayed as in Fig. 1. The negatively charged Nanogold  
15 appeared to be partially trapped in spaces formed by porous structures (arrows) in the  
16 cell wall that formed in response to ssDNA and LiAc/RbAc.

17

18 **Fig. 3.** Behavior of ssDNA visualized using YOYO-1. Cells were incubated at 42°C for  
19 20 min in the presence of PEG and ssDNA with (A) LiAc and YOYO-1 alone, (B) LiAc  
20 and YOYO-1-labelled plasmid DNA, or (C) YOYO-1-labelled plasmid DNA alone.  
21 After incubation, cells were gently washed in sterilized water and observed by  
22 fluorescence microscopy.

23

24

1

Table 1. ssDNA and LiAc in combination synergistically enhance transformation efficiency and frequency

Composition <sup>a</sup>	Transformation efficiency		Viable cells		Transformation efficiency (A/B)
	cfu/ $\mu$ g pRS415 (A)	fold	cfu ( $\times 10^4$ ) (B)	fold	fold
None <sup>b</sup>	2,008	1	1,592	1.00	1
	$\pm 1,606$		$\pm 8$		
ssDNA <sup>c</sup>	13,613	7	1,821	1.14	6
	$\pm 8,226$		$\pm 141$		
LiAc <sup>d</sup>	84,888	42	1,128	0.71	60
	$\pm 37,692$		$\pm 400$		
LiAc + ssDNA <sup>e</sup>	1,007,500	501	1,428	0.90	560
	$\pm 657,319$		$\pm 76$		
RbAc + ssDNA <sup>f</sup>	555,625	276	1,372	0.86	321
	$\pm 239,194$		$\pm 220$		

<sup>a</sup> The cells were incubated together with 0.2  $\mu$ g pRS415 at 42°C for 20 min in 42  $\mu$ l suspension containing 36% PEG alone <sup>b</sup>, 36% PEG plus 0.29 mg/ml ssDNA <sup>c</sup>, 36% PEG plus 10.7 mM LiAc <sup>d</sup>, 36% PEG plus 10.7 mM LiAc, and 0.29 mg/ml ssDNA <sup>e</sup>, 36% PEG plus 10.7 mM RbAc, and 0.29 mg/ml ssDNA <sup>f</sup>

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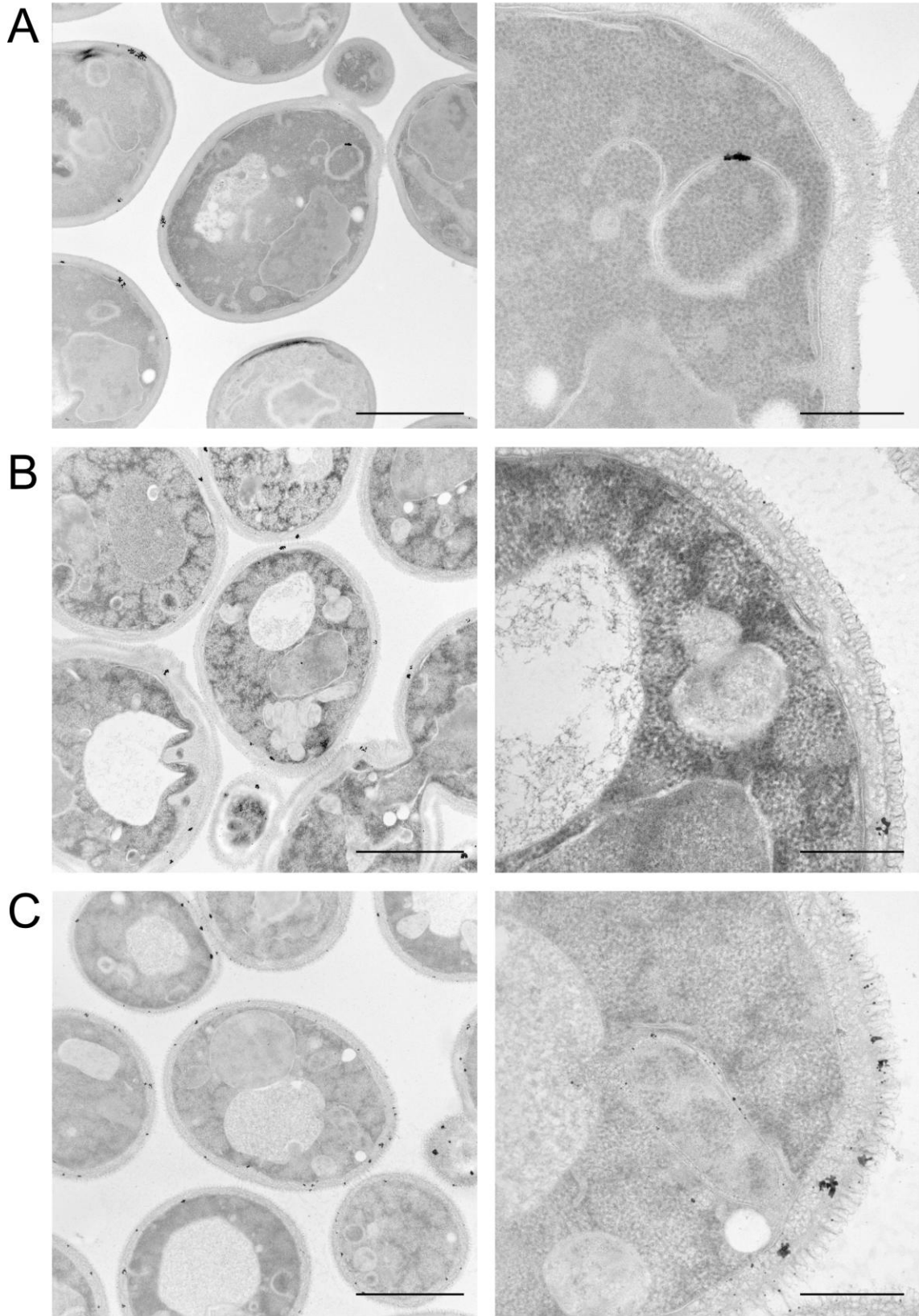
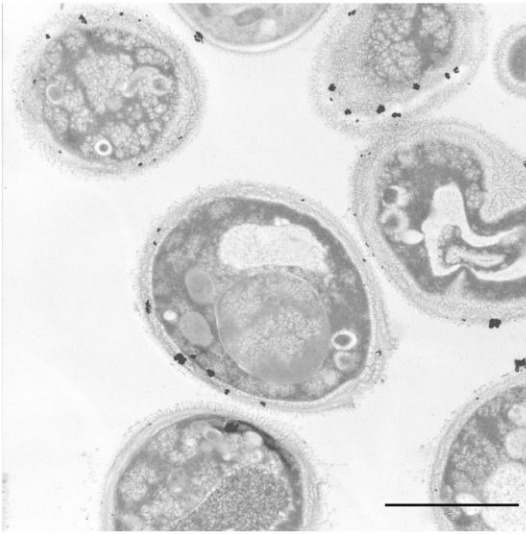


Fig. 1 Tuan et al.

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A



B

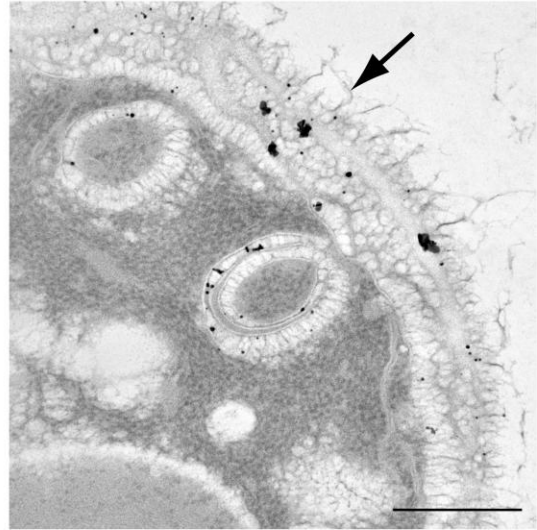
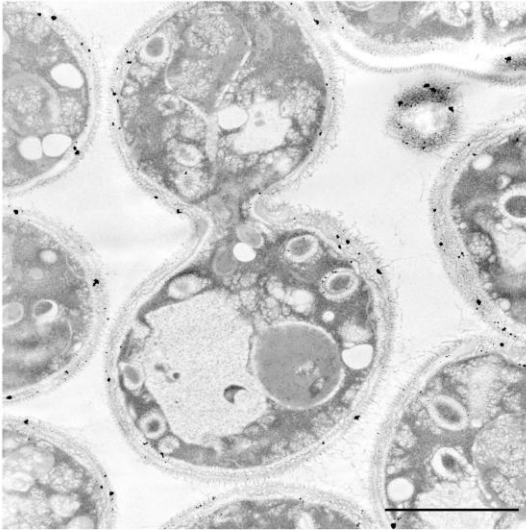


Fig. 2 Tuan et al.

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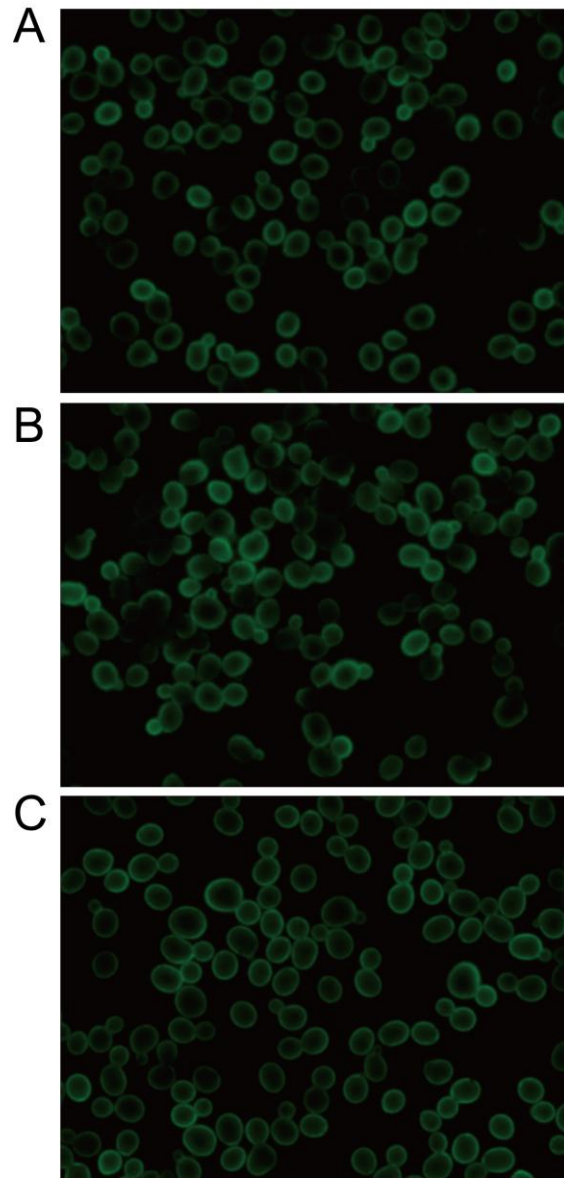


Fig. 3 Tuan et al.

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