

Optimizing Bacterial Transformation in the pDNA-Cationic Lipid - Surfactant Complex Systems by Adjusting CaCl₂ Concentration and Incubation Temperature

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Optimizing Bacterial Transformation in the pDNA-Cationic Lipid - Surfactant Complex Systems by Adjusting CaCl₂ Concentration and Incubation Temperature

Summary : In this study effects of CaCl₂ concentration and incubation temperature on the *E. coli* transformation by plasmid DNA complexes were investigated. Three pDNA formulations (pDNA, pDNA + Zwittergent® 3-08, pDNA: Cationic Lipid (1:1w/w) + Zwittergent® 3-08) were examined. The formulation containing plasmid DNA : cationic lipid complex (1:1 w/w) and Zwittergent® 3-08 showed better transformation efficiency. 50 mM CaCl₂ concentration and 37-40°C incubation temperatures showed positive effects on the number of transformed *E. coli* cells.

Key Words: plasmid DNA; cationic lipids; zwitterionic surfactants; bacterial transformation; *E. coli*

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pDNA - Katyonik Lipid - Surfaktan Kompleks Sistemlerinde CaCl₂ Konsantrasyonunun ve İnkübasyon Sıcaklığının Ayarlanmasıyla Bakteri Transformasyonunun Optimizasyonu

Özet : Bu çalışmada, CaCl₂ konsantrasyonu ve inkübasyon sıcaklığının plazmid DNA kompleksleri ile *E. coli* transformasyonu üzerindeki etkileri araştırılmıştır. Üç plazmid DNA formülasyonu (pDNA, pDNA + Zwittergent® 3-08, pDNA: Katyonik Lipid (1:1a/a) + Zwittergent® 3-08) araştırılmıştır. Plazmid DNA : katyonik lipid kompleksi (1:1 a/a) ve Zwittergent® 3-08 içeren formülasyon, en iyi transformasyon etkinliğini göstermiştir. 50 mM CaCl₂ konsantrasyonu ve 37-40°C inkübasyon sıcaklıklarının transforme olmuş *E. coli* hücre sayısı üzerinde olumlu etkiye sahip olduğu belirlenmiştir.

Anahtar kelimeler : plazmid DNA; katyonik lipidler; zwitteriyonik surfaktanlar; bakteri transformasyonu; *E. coli*

INTRODUCTION

Transformation is an important step in DNA cloning, which is one of the most important components of recombinant DNA technology and many other areas of molecular biology¹. Since transformation of *E. coli* is an essential step in many cloning experiments, it is desirable that it should be as efficient as possible². Most types of bacteria cannot take up DNA efficiently unless they have been exposed to special chemical or physical treatments to make them permeable³. There are two main techniques for transforming bacterial cells. The first and perhaps the most commonly used method is to make bacteria 'competent' by treat-

ing with a chemical such as CaCl₂ to make the bacterial cell wall permeable to DNA transfer^{4,5}. Several groups of workers have examined the factors affecting the efficiency of transformation. It has been found that *E. coli* cells and plasmid DNA can interact effectively in an environment of calcium ions and low temperature (0-5°C), and that a subsequent heat shock (37-45°C) is useful, but not strictly necessary⁶⁻⁹. Several other methods, especially the inclusion of metal ions in addition to calcium, have been shown to improve the process¹⁰.

The other method of transforming bacteria is electroporation, which requires special expensive equip-

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ment and has poor reproducibility¹¹. For years, electroporation has been an effective method for introducing DNA into bacteria, yeast, mammalian cells and plant protoplasts. Electroporation involves the application of an electric field creating pores or holes in the membranes of cells through which DNA is driven as a result of its negative charge¹²⁻¹⁶.

The aim of our study was optimization of a transformation method for recombinant DNA technology by adjusting CaCl₂ concentration and incubation temperature of bacterial growth in the systems of plasmid DNA complexed with cationic lipids and zwitterionic surfactants.

MATERIALS AND METHODS

MATERIALS

ESCORTTM, a lipid suspension, consisting of 2 mg/mL DOTAP/DOPE [(N (1- (2, 3-dioleoyloxy) propyl) - N, N, N, trimethyl ammoniumbromide)/(Dioleoylphosphatidyl ethanolamine)] (1:1 w/w) was obtained from Sigma Chemical Company. Zwitterionic surfactant Zwittergent[®] 3-08 was obtained from Calbiochem. 3 kb, Escherichia coli pBluescript plasmid was selected as the model vector which carries an ampicillin resistance gene and β -galactosidase gene alpha-subunit (Stratagene).

METHODS

Parameters Evaluated for Transformation Efficiency of E. coli

For optimization of transformation three different formulations, four different bacterial growth temperatures and three different CaCl₂ concentrations

were selected. Parameters evaluated for transformation efficiency of E. coli are seen in Table 1.

Plasmid DNA Isolation

Plasmid DNA pBluescript, was selected as a model vector which carries an ampicillin resistance gene and β -galactosidase gene α -subunit. The plasmids were amplified in XL-1 Blue strain of Escherichia coli. Bacteria containing plasmids were grown overnight, plasmids were isolated by alkaline lysis method⁴ and purified using multiple phenol/chloroform extractions. DNA was then precipitated by ethanol and centrifuged. After the removal of the supernatant, the pellet was dried and dissolved in 50 μ L Tris-EDTA (TE) buffer which contains 10 mM Tris pH 7.4, 1mM ethylenediaminetetraacetic acid (EDTA) and RNase at a concentration of 20 μ g.mL⁻¹. The plasmids were stored at -20 °C until use. DNA concentrations were measured by UV absorption at 260 nm, and purity was confirmed by 1% agarose gel electrophoresis⁴.

Preparation of Transformation Systems

Complexes used for transformation of bacteria were prepared by mixing buffer solutions of pBluescript plasmid with the lipid reagent that contains cationic lipid DOTAP and zwitterionic neutral helper lipid DOPE at a ratio of 1:1 (w/w) and zwitterionic surfactant at minimum inhibitor concentration of 3.75 % (w/v). ESCORTTM lipid suspension was the lipid mixture, and Zwittergent[®] 3-08 (Z-8) solution was used as the surfactant. Naked plasmid DNA and plasmid DNA mixed with only zwitterionic surfactant Z-8 without lipid mixture were also used in transformation studies.

Transformation Studies

Escherichia coli XL-1 Blue cells and pBluescript plas-

Table 1. Parameters Evaluated for Transformation Efficiency of E. Coli

Formulations		Bacterial Growth Temperature	CaCl ₂ Concentrations (mM)
Formulation	Code		
(A) Naked pDNA	pDNA	25°C	10 50 100
		30°C	10 50 100
		37°C	10 50 100
		40°C	10 50 100
(B) pDNA + Zwittergent [®] 3-08	pDNA + Z-8	37°C	10 50 100
(C) pDNA:Cationic lipid (1:1 w/w) + Zwittergent [®] 3-08	pDNA:L (1:1w/w) + Z-8	40°C	10 50 100

mid which contains a gene for Ampicillin resistance were used. A single colony of XL-1 Blue was used to inoculate 15 mL of Luria-Bertani (LB) broth growth medium and grown overnight at 25°C, 30°C, 37°C and 40 °C in a shaking incubator. After 5 minutes centrifugation of the overnight culture at 4000 rpm, pellets were resuspended in 2 mL of 10 mM, 50 mM and 100 mM CaCl₂ solutions. 50 µL of bacterial suspension was added to 4.8 µg / 50 µL pDNA samples and after incubation for an hour at 37°C, samples were washed twice with 5% (w/v) BSA solution by centrifugation. Pellets were suspended in 100 µL of LB medium and inoculated on LB agar containing 75µg/mL ampicillin after incubating at 37°C for one hour. The transformed colonies were counted either after 24 hours or 48 hours incubation periods.

RESULTS AND DISCUSSION

Formulation Effects

Three plasmid DNA formulations (pDNA, pDNA + Zwittergent® 3-08, pDNA: Cationic Lipid (1:1w/w) + Zwittergent® 3-08) were investigated regarding their transformation efficiencies. Bacterial transformation could not be achieved by using naked pDNA and the least efficient transformation was observed by using plasmid DNA:Zwittergent® 3-08 mixture. Among these three formulations, plasmid DNA : cationic lipid complex (1:1 w/w) and Zwittergent® 3-08 complex formulation was determined as the most efficient mixture. Complex formulations were prepared by using different amounts of cationic lipids. Optimal lipid - DNA ratio in complex formation was determined according to the results of gel retardation assay, DNase I protection assay and bacterial transformation. By increasing the amounts of lipids in the complex formulations, protection of DNA against DNase I enzyme was achieved. Although all complex formulations could be transformed, due to the cell toxicity of the lipids, transformation efficiency is decreased over 1:1 ratio. After selecting the best pDNA : cationic lipid ratio which was 1:1 w/w (Figure 1), zwittergent was incorporated into the system by considering their MIC values.

Optimum CaCl₂ Concentration

Effect of the 10 mM, 50 mM and 100mM CaCl₂ con-

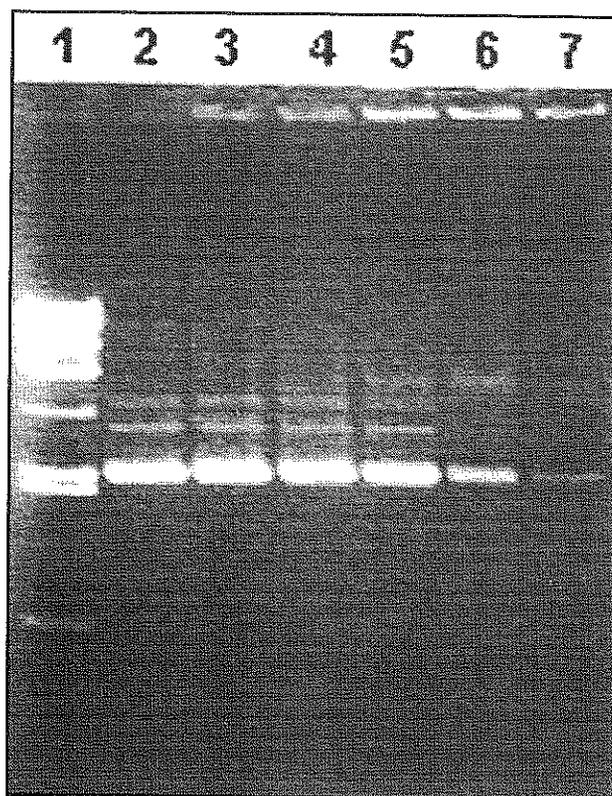


Figure 1. Gel Retardation Assay

- Lane 1: λ DNA Hind III Digest MW Marker
- Lane 2: Plasmid DNA
- Lane 3: Plasmid DNA:Lipid → 1: 0,125 w/w
- Lane 4: Plasmid DNA:Lipid → 1: 0,25 w/w
- Lane 5: Plasmid DNA:Lipid → 1: 0,5 w/w
- Lane 6: Plasmid DNA:Lipid → 1: 1 w/w
- Lane 7: Plasmid DNA:Lipid → 1: 2 w/w

centrations on the bacterial transformation was studied. As seen from Figure 2-A, only a few bacteria can be transformed with pDNA at 10 mM CaCl₂ concentration whereas at 100 mM CaCl₂ concentration which is very high, transformant bacterial cells are less than at 50 mM CaCl₂ levels (Figure 2-C). The optimum CaCl₂ concentration for an efficient transformation was found to be 50 mM (Figure 2-B). Factors such as plasmid characteristics, amount, purity, cell density and marker all influence the transformation efficiency. Transformation efficiency can be affected by the Ca²⁺ induced phosphatidylethanolamine mechanism⁸. Here CaCl₂ concentration is a critical parameter in the phosphoethanolamine dependent membrane enzyme activity.

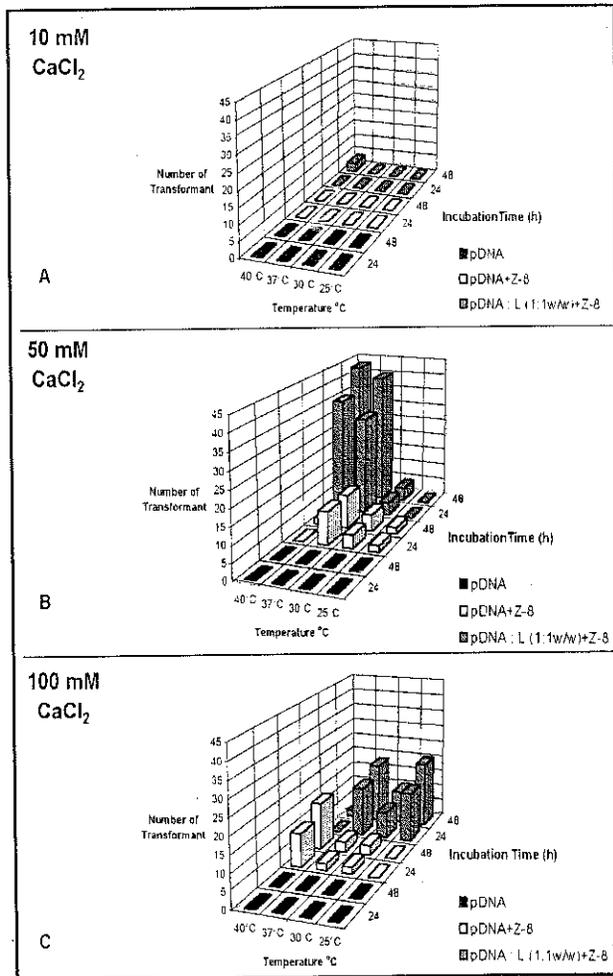


Figure 2. Effect of CaCl₂ concentrations on transformation efficiency of formulations which were transformed to bacteria grown at different temperatures.
 (A) 10 mM CaCl₂ concentration
 (B) 50 mM CaCl₂ concentration
 (C) 100 mM CaCl₂ concentration

Optimum Incubation Temperature

Incubation temperature is also an important factor in the transformation efficiency of *E. coli* cells. Among the temperatures investigated, 25°C was found least efficient as transformation temperature for 10 mM and 50 mM concentrations (Figure 3). 30°C, 37°C and 40°C temperatures were found suitable for transforming bacteria for 50 mM and 100 mM concentrations. In Figure 3 the transformation profile of bacteria grown at different temperatures is seen. CaCl₂ concentration and composition of the pDNA formulation have additional effects on the incubation temperatures.

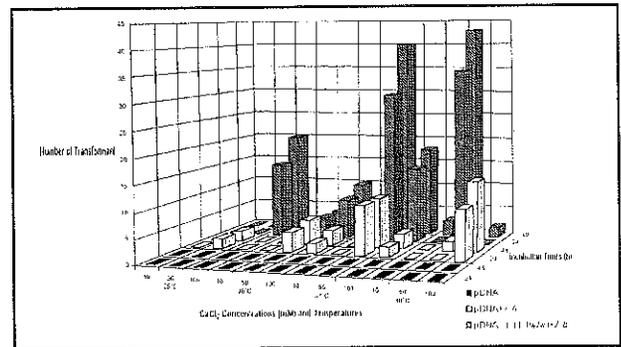


Figure 3. Effect of different CaCl₂ concentrations and bacteria incubation temperatures on transformation efficiency.

By considering all the factors, 50 mM CaCl₂ concentration, plasmid DNA : cationic lipid (1:1 w/w) + Zwittergent® 3-08 complex formulation and 37°C - 40°C temperatures seem to be productive conditions for optimal transformation studies.

CONCLUSION

It can be concluded from the results that cationic lipid and surfactant mixtures and optimized CaCl₂ concentration and temperature conditions may provide an efficient way for transforming bacteria with pDNA. The positively charged lipids can attach to pDNA, and also bind to cells to simplify DNA transfer into the cells. Cationic lipids have been used for transfection of eucaryotic cells for years, but in this study it has been established that these systems are also useful for transformation of procaryotic cells. CaCl₂ concentration and incubation temperatures are important parameters in optimising bacterial transformation.

REFERENCES

1. Brock TD, Smith DW, Madigan MT, Biology of Microorganisms, Prentice-Hall, Inc., New Jersey, 1984.
2. Old RW, Primrose SB, Introduction to Gene Manipulation, Carr NG, Ingraham JL, Rittenberg SC (eds.), Principles of Gene Manipulation: An Introduction to Genetic Engineering, Blackwell Scientific Publications, Oxford, 1-14, 1989.
3. Synder L, Champness W, Molecular Genetics of Bacteria, ASM Press, Washington, 1997.
4. Sambrook J, Fritsch EF, Maniatis T, Molecular Cloning, Cold Spring Harbor Laboratory Press, New York, 1989.
5. Nakata Y, Tang X, Yokoyama KK. Preparation of competent cells for high-efficiency plasmid transformation of *Escherichia coli*, *Methods. Mol. Biol.*, 69, 129-137, 1997.

6. Huff JP, Grant BJ, Penning CA, Sullivan KF. Optimization of routine transformation of *Escherichia coli* with plasmid DNA, *Biotechniques*, 9 (5), 570-577, 1990.
7. Tomley FM. Transformation of *E. coli*, *Methods. Mol. Biol.*, 58, 241-247, 1996.
8. Kanipes MI, Lin S, Cotter RJ, Raetz CR. Ca²⁺-induced phosphoethanolamine transfer to the outer 3-deoxy-D-manno-octulosonic acid moiety of *Escherichia coli* lipopolysaccharide. A novel membrane enzyme dependent upon phosphatidylethanolamine, *J. Biol. Chem.*, 276 (2), 1156-1163, 2001.
9. Lin SY, Kondo F. A simple classification method for residual antibiotics using *E. coli* cells transformed by the calcium chloride method and drug resistance plasmid DNA, *Microbios*, 104 (409), 149-158, 2001.
10. Viljanen P. The effect of polymyxin B nonapeptide (PMBN) on transformation, *Biochem. Biophys. Res. Commun.*, 143 (3), 923-927, 1987.
11. Brooks G., Gene Therapy, *The Pharm. J.*, 252, 256-260, 1994.
12. Swartz M, Eberhart J, Mastick GS, Krull CE. Sparking New Frontiers: Using In Vivo Electroporation for Genetic Manipulations, *Developmental Biology*, 233, 13-21, 2001.
13. Lurquin PF. Gene transfer by electroporation, *Mol. Biotechnol.* 7 (1), 5-35, 1997.
14. Somiari S, Glasspool-Malone J, Drabick JJ, Gilbert RA, Heller RA, Jaroszeski MJ, Malone RW. Theory and in vivo application of electroporative gene delivery, *Mol. Ther.*, 2 (3), 178-87, 2000.
15. Dezawa M, Takano M, Negishi H, Mo X, Oshitari T, Sawada H. Gene transfer into retinal ganglion cells by in vivo electroporation: a new approach, *Micron* 33, 1-6, 2002.
16. Taketo A. Properties of electroporation-mediated DNA transfer in *Escherichia coli*, *J Biochem.*, 105 (5), 813- 817, 1989.