



## Cloning and Expression of Lip-A Gene From *Streptomyces Coelicolor* to *Escherichia Coli*

### KEYWORDS

Lip A gene, restriction digestion, transformation, lipases

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### ABSTRACT

The exponential increase in the application of lipases in various fields in the last few decades demands extension in both qualitative improvement and quantitative enhancement. Strain improvement is an essential part of process development for fermentation products. Developed strains can reduce the costs with increased productivity and can possess some specialized desirable characteristics and such improved strains can be obtained by mutation. Thus in the present investigation the *E.coli* DH5 $\alpha$  were cloned with LipA gene isolated from *Streptomyces* species using a PUC 18 plasmid. The transformation was confirmed by culturing the cloned cells on lipase medium and restriction digestion of the recombinant cells and un cloned cells by HIND III restriction enzyme where in the *E.coli* DH5 $\alpha$  from the wild strains were not cloned. The cloned *E.coli* DH5 $\alpha$  could be one of the sources for lipolytic enzymes for the industries

### INTRODUCTION

Lipases catalyze both the hydrolysis and the synthesis of esters formed from glycerol and long-chain fatty acids. These enzymes usually exhibit broad substrate specificity and degrade acyl *p*-nitrophenyl esters, Tweens, and phospholipids, often with positional selectivity, stereo selectivity, and chain length selectivity. Lipase's are the enzyme that have been proved to play a very important role in oil degrading activity.

The versatility of lipases have resulted in their emergence as one of the leading biocatalysts with definite potential to contribute in exploitation of the less utilized lipid technology and multibillion dollar versatile industrial applications (Joseph *et al.*, 2008). Lipases occur widely in nature, but only microbial lipases are commercially significant.

Commercially useful lipases are usually obtained from microorganisms that produce a wide variety of extracellular lipases. Many lipases are active in organic solvents where they catalyze a number of useful reactions including esterification (Chowdary *et al.*, 2001; Hamsaveni *et al.*, 2001; Kiyota *et al.*, 2001; Krishna and Karanth, 2001; Rao and Divakar, 2001), transesterification, regioselective acylation of glycols and menthols, and synthesis of peptides (Ducret *et al.*, 1998; Zhang *et al.*, 2001) and other chemicals (Therisod and Klibanov, 1987; Weber *et al.*, 1999; Berglund and Hutt, 2000; Liese *et al.*, 2000; Azim *et al.*, 2001). The expectation is that lipases will be as important industrially in the future as the proteases and carbohydrases are currently.

*Streptomyces* species are known to be lipolytic (Sztajer *et al.*, 1988) and two different types of lipase genes have been cloned from this genus. lipase genes of *Streptomyces exfoliatus* M11 (Pérez *et al.*, 1993) and *Streptomyces albus* G (Cruz *et al.*, 1994), and shown that these highly similar to a psychrophilic lipase from *Moraxella* sp. TA144 (Feller *et al.*, 1990). A second type of lipase gene encoding lipase has been related to group II *Pseudomonas* lipase that has been recently cloned from *Streptomyces cinamomeus* (Sommer *et al.*, 1997).

### MATERIALS AND METHODOLOGY

#### Isolation of Genomic DNA

*Streptomyces* culture was procured from NCIM Pune. The mycelia mat of the pure culture was homogenized in a mortar and pestles along with the addition of CTAB extraction buffer. The homogenate was incubated for 30 minutes at 65°C using water bath. After incubation equal volume of chloroform isoamylalcohol was added and centrifuged for 10 minutes. The process was repeated twice. The aqueous phase was separated to which isopropanol was added and incubated overnight.

The contents were then transferred to eppendorf tubes and centrifuged at 4°C at 10000 rpm for 10 minutes. The supernatant was discarded and the pellet was precipitated with 70% ethanol and kept for air drying. The pellet was then dissolved with TE buffer and stored in refrigerator until further use. The DNA was qualitatively determined by running on 1% agarose gel electrophoresis.

#### Isolation of plasmid from *E.coli* by alkaline lysis method

1.5 ml of overnight culture of *E.coli* (TBI) carrying plasmid (pUC18) was centrifuged for 10 minutes at 5000 rpm. The pellet was suspended in 1 ml of solution A and 2ml of solution B. they were then incubated on ice for 30 minutes till the solution becomes viscous. 0.5 ml of solution c was added, kept on ice for 60 minutes. The suspension was the centrifuge for 10 minutes at 4°C. Phenol: chloroform (1:1) was added to the supernatant then it was centrifuge for 10 minutes at 10,000 rpm. The supernatant was discarded and 10 $\mu$ l of 70% ethanol was added to the pellet for removing other contaminants. The ethanol was then air dried and pellet was re-suspended in 30  $\mu$ l of TE buffer. The obtained plasmid (pUC18) was analysed by electrophoresing in 1% agarose gel electrophoresis

#### Restriction digestion of the Genomic DNA and plasmid

The restriction of the genomic DNA and the plasmid was carried out by using Hind III restriction enzyme. The fractions obtained were determined on agarose gel and the ligation was carried out.

#### Ligation of Genomic DNA and plasmid

The reaction mixture required for ligation was mixed in ice cold condition. The ligase buffer, the digested plasmid (pUC 18) and digested genomic DNA was thawed and kept in ice. To the fresh sterile 1.5 ml eppendorf tube 8.1  $\mu$ l of sterile distilled water was taken. To this 1.5 $\mu$ l of 10x ligase buffer was added. 3  $\mu$ l of digested genomic and 1.5  $\mu$ l digested plasmid pUC 18 was mixed gently by tapping. Finally 3 $\mu$ l (1 unit) of T<sub>4</sub> DNA ligase was added and mixed gently. The ligation mixture was incubated at 20°C for overnight and later used for *E.coli* (DH 5 $\alpha$ ) transformation.

#### Transformation

24 hours overnight culture of *E.coli* (DH5 $\alpha$ ) was taken. From that 0.1 ml was taken and inoculated in 25 ml of fresh LB medium. The culture was allowed to grow till its OD reaches 0.5-0.6 at 540 nm. The culture was chilled in ice for 30 minutes. Cells were harvested by centrifugation at 5000 rpm for 10 minutes. Supernatant was discarded and the pellet was suspended in 5 ml of 100mM MgCl<sub>2</sub>. Once again the pellet was kept in ice for 60 minutes. Culture was spinned, discarded the supernatant and the pellet was suspended in 1 ml of 100mM MgCl<sub>2</sub>. From this 0.2 ml of culture was taken and 50 $\mu$ l of plasmid pUC18 was added. Suspension was kept in ice for 30 min. Heat shock is given at 42°C for 2 minutes. Then it is again transferred to ice and kept for 5 minutes. Then 2 ml of sterile LB broth was added to the above culture and incubated at 37°C for 1 ½ hour. 0.5 ml of these solutions was taken and plated it on agar plates containing ampicillin (50 $\mu$ g/ml). The plates were then incubated at 37°C overnight. For control 0.2 ml of the culture with only buffer and not with plasmid and sterile LB was added. After incubation 0.5 ml of the control culture was plated on the selective plate. The plates were then incubated at 37°C along with the test plates. After incubation colonies that were formed on the ampicillin plates were observed. In the control plate, no colonies were observed.

Plasmid DNA isolation from cloned cells: 1ml of overnight culture was taken into an eppendorf tube and was centrifuged at 12000rpm for 1minute. The pellet was dissolved in 100 $\mu$ l of double distilled water by gently tapping at the bottom of the tube. 100 $\mu$ l of lysis buffer was added. The eppendorf tubes top was wrapped with parafilm and punctured with a fine needle to avoid popping of the solution. The mixture was kept immediately in boiling water bath for 2 minutes. 50 $\mu$ l of 1 M MgCl<sub>2</sub> was added, mixed by tapping and kept on ice for two minutes. The mixture is centrifuged at 12000 rpm for two minutes. 50 $\mu$ l of 5M potassium acetate was added to the same tube and mixed without disturbing the pellet, by tapping gently and is kept on ice for two minutes. It was centrifuged for two minutes at 12000rpm. The supernatant was taken (leaving the bottom portion that contains the pellet) and transferred to a fresh tube. To this 600 $\mu$ l isopropanol was added. Mixed by inverting slowly and the tubes were kept on ice for two minutes. Centrifuged at 12000 rpm for two minutes and the supernatant was discarded. The pellet was washed with 1ml 70% alcohol and centrifuged for 10 minutes 12000rpm. The pellet were dried completely and re suspended the pellet in 50 $\mu$ l water containing 20 $\mu$ g/ml RNase A.

Restriction Digestion of the plasmid isolated from cloned cells: The following mixture was prepared for restriction of the plasmid in a sterile 1.5 ml eppendorf and tube kept in ice cold condition:

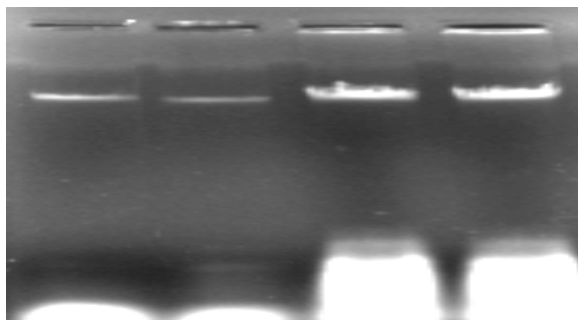
Recombinant Plasmid (pUC18) (0.2-1.0 $\mu$ g)	2 $\mu$ l
10 x Restriction buffer	2 $\mu$ l
Restriction enzyme Hind III (3 units / $\mu$ g plasmid)	1 $\mu$ l
Distilled water	20 $\mu$ l

**Table-1 Reaction Mixture**

The eppendorf tubes were incubated the tube at 37°C for 3 hr. The reaction was stopped by adding 0.5 M EDTA (pH 8.0) to give a final EDTA to 20  $\mu$ l reaction volume. The restricted plasmid was analysed on agarose gel.

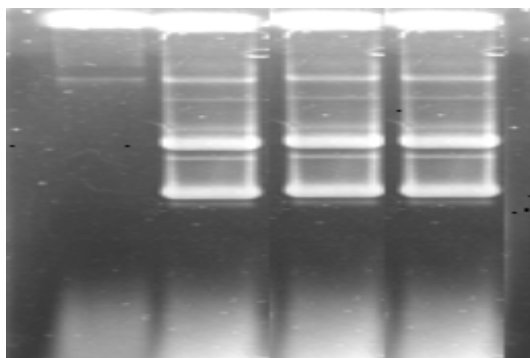
#### RESULTS

Isolation of Genomic DNA: The Genomic DNA was isolated from *Streptomyces coelicolor* by following CTAB method. The isolated DNA was preserved in TE buffer at 4°C and it was electrophorised for qualitative estimation (Fig-1)



**Figure-1: Qualitative estimation of Genomic DNA.**

Isolation of plasmid from *E.coli*: Plasmid Puc18 was isolated from *E.coli*(TB1) by alkaline lysis method. The isolated plasmid was stored at 4°C in TE buffer and electrophorised in 0.8% agarose gel to determine its purity (Fig-2).



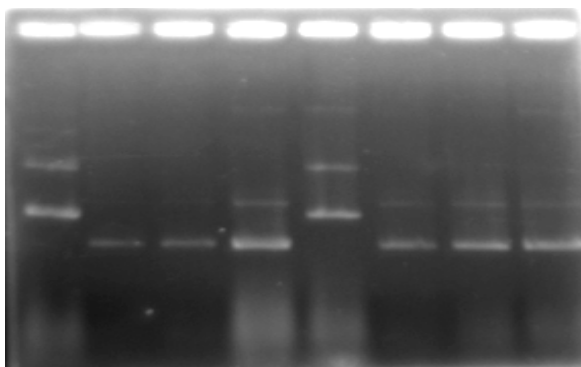
**Figure-2: PUC18 Plasmid isolated from *E.coli***

Restriction Digestion and ligation of the Genomic DNA and Plasmid: The fragments were obtained by restriction digestion of the genomic DNA by Hind III enzyme and were ligated into the plasmid DNA using T4 DNA ligase such that the plasmid carries the gene of interest, that was transformed into *E.coli*

Transformation: The transformation was carried out by competence cell preparation of *E.coli*, these competent cells were then inserted with the cloned plasmid. The cells were cultured on specific media plate to determine the cells with transformed plasmid.

Lipolytic activity of the recombinant cells: The recombinant cells were plated in lipolytic medium and lipolysis was occurred on the plates indicates the successful transformation of target gene, where native wild type *E.coli* (DH5 $\alpha$ ) does not show lipolytic action.

Restriction digestion of the cloned plasmid from the recombinant *E.coli*: The isolation of the plasmid was carried out from the recombinant cells and restriction digestion was carried out with Hind III enzyme and was determined on agarose gel electrophoresis to indicate the presence of the gene in the recombinant cells (Fig-3).



**Figure-3: Restriction Digestion of the recombinant cells. (Lane 1- Recombinant plasmid (pUC18-1), Lane 2 and 3-Non recombinant plasmids, Lane 4 - Control plasmid (pUC18), Lane 5- Recombinant plasmid (pUC18-2), Lane 6,7&8 - Non recombinant plasmids.)**

## DISCUSSION

Henne, A et al., 1999 constructed DNA libraries for *E.coli* DH5 from soil samples as a host and pBluescript SK (1) as a vector. The gene studies was carried out and was found that the product of *lip* (281 amino acids) is 30% identical (35.9% similar) to the lipase (Lip) of *S. albus*. Similar amino

acid identities were obtained for the lipases of *Streptomyces coelicolor* (Valdez, F et al., 1999) and a *Moraxella* sp. (Feller, G et al., 1990).

There are several examples where the presence of a restriction system in the recipient is found to have a marked effect on the conjugal transfer of plasmids from *E. coli* to other bacteria. The 12- fold reduction in *S. coelicolor* ex-conjugant frequency observed with pDJ50 would be consistent with that reported for other intergeneric conjugation systems (that is, - 15-fold reduction per restriction site (Elhai, J et al., 1997).

Chung et al, 1991, isolated lipase gene from *P. fluorescens* SIK W1 which consisted an open reading frame 1347 base pairs long commencing with an ATG start codon encoding a polypeptide of 499 amino acid residues and a TGA stop codon.

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