

Optimization using RSM and partial purification of lipase from oil contaminated soil samples

Dr.P. Joyruth¹ and Dr. Lali Growther²

1. Assistant Professor, Department of Microbiology, The American College, Madurai.
2. Professor and Head, PG and Research Department of Microbiology, Hindusthan College of Arts and Science, Coimbatore.

Mail ID: pjoyruth10@gmail.com, lalips@gmail.com

Phone Number: 9047472797, 9843225410

Abstract:

Enzymes are protein with highly catalytic functions and produced by all living organisms. Commonly occurring natural enzymes are lipase, lactase, amylase and cellulase. Lipases catalyze the hydrolysis and acyl glycerol synthesis. Forty-eight strains were identified from oil contaminated soil samples. Out of forty – eight isolates fifteen organisms form clear halo zone around the colonies. Maximum activity of pH 8 was noted for *Acinetobacter indicus*. *Acinetobacter indicus* obtained maximum activity at 24hrs with 17.96U/ml. Glucose and peptone confirmed the maximum activity of enzyme. CaCl₂ obtained maximum activity of lipase for the isolates. DEAE Cellulose column chromatography produced the activity of 720U/g with purification fold of 7.81 and a yield of 20.5% for *Acinetobacter indicus*. The molecular weight of lipase produced by *Acinetobacter indicus* was 60KDa.

Keywords: Enzyme, Lipase, Optimization, Chromatography, Molecular weight

1. Introduction

Enzymes are chemical catalysts and used for biotechnological applications. They are eco – friendly and gaining interest in industries in recent period. Based on report, market of enzyme revenue reach \$ 20.31 billion by 2028 (<https://www.grandviewresearch.com/> accessed on Nov 21, 2022). Enzymes are produced by fermentation of bio – based sources. Enzymes are applicable in various industries such as textiles, pharmaceutical, dairy, cosmetic, detergents and in synthesis of agro – chemical. Commonly occurring natural enzymes are lactase, amylase, cellulase and lipase (Corleone, 2018). Microbial lipases are important because of their unique properties and produce bulk extracellular lipase (Joyruth and Lali Growther, 2018). Example of bacteria that produce lipase includes *Bacillus*, *Serratia*, *Pseudomonas*, *Halomonas*, *Staphylococcus* etc., and fungi include *Rhizopus*, *Penicillium*, *Aspergillus* etc., (Najjar et al., 2011). Lipase (EC 3.1.1.3), catalyze the hydrolysis and acylglycerol synthesis (Lajis, 2018). Availability of lipase are increased due to the achievement in cloning and expression of enzymes from microorganism and also having specific properties such as specificity, pH, temperature, stability (Menoncin *et al.*, 2008). Organisms grown in tributyrin agar lacking Victoria Blue B produce clear zone around colonies that indicate production of lipase. In presence of Victoria blue B, lipolytic colonies produce dark zone against light blue background (Dogan and Boor, 2003).

Lipases are produced by both solid state and submerged fermentation. Submerged fermentation is favorable than solid state fermentation for enzyme production (Iso *et al.*, 2001) and microbial cells are distributed evenly throughout the medium in submerged fermentation. (Hashemi *et al.*, 2013). Growth of organism in nutrient enriched liquid medium is easy for sterilization (Dutra *et al.*, 2008). Production has been reported as batch, repeated-batch, fed batch and continuous fermentation (Treichel *et al.*, 2010). Media contain carbon, nitrogen, phosphorous sources and mineral salts are necessary for production of lipase. Optimization of fermentation medium and process is important to increase the profit (Schmidt 2005).

RSM uses statistical experiment design such as Box – Behnken Design (BBD), Central Composite Design (CCD) (Dahiya and Purkayastha, 2011). Central composite design obtained by addition of points to estimate interaction effects and curvature. RSM can be combined with CCD, so designed the experiments by CCD and optimized by RSM (Dahiya *et al.*, 2010). Activities of lipases were determined spectrometrically at 30 °C using p-nitrophenol palmitate as substrate. Lipase activities were performed by titrimetrically using olive oil hydrolysis (Borkar *et al.*, 2009). Several techniques are reported for purification of lipases such as hydrophobic interaction chromatography, precipitation, gel filtration, ion exchange chromatography and affinity chromatography. Purified Protein samples are important before determining protein properties and activities. These methods help to understand the functions of enzyme (Reis *et al.*, 2009). In extracellular lipases, first step in purification is removal of cells by filtration or centrifugation. Cell lysis is needed additionally, if it is intracellular lipases. After precipitation ion exchange chromatography and gel filtration are preferred (Bisht *et al.*, 2014). For effective purification ion exchange chromatography or hydrophobic interaction chromatography (Aravindhan *et al.*, 2007) are used. Extracellular lipases produced by *Rhizopus oryzae* are purified by ammonium sulfate precipitation, Sephadex G-75 gel filtration, sulfopropyl-sepharose chromatography and second sulfo-propyl-sepharose chromatography. Strategies of purification involve rapid, inexpensive, high yielding and amendable to large scale of operation (Ji *et al.*, 2010).

2. Materials and Methods

2.1. Sample collection and molecular characterization

Soil samples were aseptically collected from oil factories in Salem, Vellore, Peelamedu and Namakkal. The diluted samples were cultured on tributyrin agar plates by spread plate methods for isolation of bacteria after 24 hours of incubation (Mahima *et al.*, 2016). Clear zone forming colonies are lipase producers. Molecular characterization was determined by 16S rRNA gene sequencing, BLAST search analysis and construction of phylogenetic tree was proceed by following standard methods. (Mellmann *et al.*, 2006). 16S r RNA sequencing was carried at MACROGEN, Korea. Genomic DNA was extracted by Sambrook and Russell, 2001. Similar sequences in BLAST program were retrieved and aligned with multiple alignment programs CLUSTAL W. The multiple alignment file was then used to create Boot – strapping tree using MEGA version 6 software. (Tamura *et al.*, 2007).

2.2. Identification of lipase gene from bacterial strains

Genomic DNA from *Acinetobacter indicus* was extracted and amplified the lipase gene by PCR.

2.3. Production of enzyme under submerged fermentation (SMF)

Among forty – eight isolates, sixteen positive bacterial strains were cultivated in production media for enzyme production. Media consists of 3% yeast extract, 3% Sucrose, 0. 1 g (g/l) CaSO₄, 0. 5 g/l-KH₂PO₄, 0. 1g/l-MgSo₄.7H₂O, 1% olive oil and 100 ml of distilled water in 250ml conical flask. Inoculated flask was incubated at 37°C for 24-48 hours (Mobarak-Qamsari *et al.*, 2011).

2.4. Lipase assay

Lipase activity was demonstrated spectrophotometrically at 30 °C by using p-nitrophenol palmitate (PNPP) as substrate. (Ittikhar *et al.*, 2008). The absorbance was measured at 410 nm for first 2 minutes of reaction. One unit was defined as that amount of enzyme liberated 1µmol of PNP per minute (Karadzic *et al.*, 2006).

2.5. Effect of physico-chemical parameters

The culture sample was optimized at different physico – chemical parameters such as temperature (20°C - 50°C), pH (4 -8), Incubation period (12 – 60hrs), Carbon sources (Glucose, sucrose, mannitol, lactose and maltose), Nitrogen sources (Casein, yeast extract, albumin, peptone and urea), metal ions (MgSO₄, CaCl₂, FeSO₄, NaCl₂ and ZnCl₂), natural oils (Coconut oil, Groundnut oil, Castor oil and Neem oil), Surfactants (Tween 20, 40, 60, 80, 100, Triton X – 100 and SDS).

The lipase enzyme production was done by culturing the organisms in medium with optimized conditions. After culture, the supernatant was collected and tested the lipase activity to confirm the highest production of lipase.

2.6. Statistical analysis of optimization factors

2.6.1. Modeling by Response surface methodology (RSM)

RSM is the statistical method for modeling the optimization to identify the interaction between one or more response parameters. RSM is use to study the combined effects of medium components and optimize the parameters in fermentation process. (Gunawan *et al.*, 2005)

2.6.2. Selection of parameters for central composite design

Using submerged fermentation, optimization of lipase depends on various factors such as temperature, pH, incubation time, moisture and chemical factors. The present study was based on central composite design (CCD) of response surface methodology (RSM), to perform response surface model. (Mehta *et al.*, 2019)

2.6.3. Statistical analysis

The interaction terms on response and effect of parameters were studied by Analysis of Variance (ANOVA) and significance test to check the model adequacy. The analysis was made through surface plot using MINITAB 14 software. (Chauhan *et al.*, 2013)

2.7. Enzyme purification

Lipase enzyme was purified by different steps like ammonium sulfate precipitation, dialysis, and DEAE-Cellulose column chromatography. After each step, observed the lipase activity, protein estimation by Lowry's method, specific activity and fold purification. The purification steps were carried out at 4°C (Aravindan *et al.*, 2007).

2.8. Ammonium sulfate precipitation

The crude enzyme was precipitated by ammonium sulfate at 30 - 70% saturation and then kept undisturbed for overnight at 4°C (Joyruth and Lali Growther, 2020). The precipitate was collected by centrifugation at 15,000rpm for 15 minutes. The pellet obtained was dissolved in minimal volume of 10mM Tris HCL buffer at pH 7.5. (Venil *et al.*, 2009).

2.9. Dialysis against buffer

The pellet obtained after ammonium sulfate precipitation was dissolved in 10mM Tris HCl buffer at pH 7.5 and introduced into pre-treated dialysis bag and placed in 10mM Tris-HCl buffer and dialyzed overnight at 4°C. Then the lipase assay and estimation of protein by Lowry's method were done to this sample.

2.10. DEAE - Cellulose column chromatography

Solution of dialysis contains both soluble and insoluble materials. This was centrifuged to remove insoluble material. The supernatant was applied to DEAE - Cellulose column (2.4x45 cm), equilibrated with 10 mM Tris HCl buffer, pH 7.5. Then it was eluted with 10mM Tris HCl buffer. The flow rate was adjusted to 0.5ml/min. About fifteen fractions were collected at a rate of 10ml/hr. The protein estimated by Lowry's method and enzyme assay was performed to each fraction. The lipase which showed highest enzyme activity was pooled and stored at 4°C and used as the purified lipase (Gupta *et al.*, 2004).

2.11. Estimation of protein

Quantitative estimation of the protein content obtained after different steps were done by Lowry *et al.*, (1951) method.

2.12. Determination of molecular weight by SDS-PAGE

The molecular weight of the partially purified lipase from ammonium sulfate precipitation and dialysis was checked by SDS. (Rakesh kumar *et al.*, 2012).

3. Result and discussion

3.1. Isolation of bacterial species from soil sample

Totally 10 soil samples were collected from oil factory in Salem, Vellore, Peelamedu and Namakkal. Totally forty – eight bacterial species were isolated using nutrient agar and listed in table no.1.

Table 1. Isolation of Bacterial species from oil factory

Salem	Peelamedu	Vellore	Namakkal
<i>Bacillus sp.</i> ,	<i>Staphylococcus sp.</i> ,	<i>Micrococcus sp.</i> ,	<i>Staphylococcus sp.</i> ,
<i>Pseudomonas sp.</i> ,	<i>Pseudomonas sp.</i> ,	<i>Staphylococcus sp.</i> ,	<i>Bacillus sp.</i> ,
<i>Staphylococcus sp.</i> ,	<i>Klebsiella sp.</i> ,	<i>Aeromonas sp.</i> ,	<i>Klebsiella sp.</i> ,
<i>Escherichia coli</i>	<i>Bacillus sp.</i> ,	<i>Bacillus sp.</i> ,	<i>Arthobacter sp.</i> ,
<i>Proteus sp.</i> ,	<i>Aeromonas sp.</i> ,	<i>Escherichia coli</i>	<i>Pseudomonas sp.</i> ,
<i>Klebsiella sp.</i> ,	<i>Lactobacillus sp.</i> ,	<i>Halomonas sp.</i> ,	<i>Aeromonas sp.</i> ,
<i>Micrococcus sp.</i> ,	<i>Serratia sp.</i> ,	<i>Pseudomonas sp.</i> ,	<i>Pseudomonas sp.</i> ,
<i>Streptococcus sp.</i> ,	<i>Micrococcus sp.</i> ,	<i>Streptococcus sp.</i> ,	<i>Bacillus sp.</i> ,
<i>Acinetobacter sp.</i> ,	<i>Bacillus sp.</i> ,	<i>Serratia sp.</i> ,	<i>Lactobacillus sp.</i> ,
<i>Staphylococcus sp.</i> ,	<i>Halomonas sp.</i> ,	<i>Klebsiella sp.</i> ,	<i>Serratia sp.</i> ,
<i>Proteus sp.</i> ,	<i>Enterobacter sp.</i> ,	<i>Stenotrophomonas sp.</i> ,	<i>Halomonas sp.</i> ,
<i>Halomonas sp.</i> ,	<i>Serratia sp.</i> ,	<i>Enterobacter sp.</i> ,	<i>Proteus sp.</i> ,

3.2. Screening of microorganisms

All the isolated microorganisms were screened for lipase production using tributyrin agar plates, only sixteen colonies showed clear zone around the colony and were selected for further studies. Clear zone producing strain was shown in figure 1.

Figure 1. Plate showing clear zone around the colony



3.3. Identification of bacterial strains

Based on colony morphology, cell morphology and studied biochemical test, the isolates were identified (table:2).

Table 2. Biochemical results for isolated organisms

S.No	Isolates	Gram staining	Motility test	Indole test	MR test	VP test	Citrate test	Catalase test	Oxidase test	Urease test	Carbohydrate Fermentation test			
											Sucrose	Maltose	Lactose	Fructose
1	<i>Bacillus sp.</i> ,	+	+	-	-	+	+	+	-	-	+	+	-	+
2	<i>Pseudomonas sp.</i> ,	-	+	-	-	-	+	+	+	-	-	-	-	-
3	<i>Staphylococcus sp.</i> ,	+	-	-	+	+	+	+	-	+	+	+	+	+
4	<i>Klebsiella sp.</i> ,	-	-	-	-	+	+	+	-	+	+	+	+	-
5	<i>Proteus sp.</i> ,	-	+	-	+	-	+	+	-	+	-	-	-	-
6	<i>Serratia sp.</i> ,	-	+	-	-	+	+	+	-	+	+	+	-	+
7	<i>Lactobacillus sp.</i> ,	+	-	-	+	-	-	-	+	-	+	+	+	+
8	<i>Streptococcus sp.</i> ,	+	-	-	-	-	-	-	-	-	+	+	+	+
9	<i>Aeromonas sp.</i> ,	-	+	+	-	+	+	+	+	-	+	+	-	+
10	<i>Micrococcus sp.</i> ,	+	-	-	-	-	-	+	+	+	-	-	-	-
11	<i>Arthrobacter sp.</i> ,	+	-	-	-	-	-	+	-	-	+	-	+	-
12	<i>Escherichia coli</i>	-	+	+	+	-	-	+	-	-	+	-	+	-
13	<i>Acinetobacter sp.</i> ,	-	-	-	+	-	+	+	-	-			-	
14	<i>Enterobacter sp.</i> ,	-	+	-	-	+	+	+	-	-	+	+	-	-

15	<i>Stenotrophomonas sp.</i> ,	-	+	-	-	-	+	+	-	-	-	-	+	+
----	-------------------------------	---	---	---	---	---	---	---	---	---	---	---	---	---

3.4. Production of enzyme using submerged fermentation

From forty- eight organisms, selected fifteen organisms were subjected for production of lipase by submerged fermentation and the activity of enzymes were shown in table 3. Out of fifteen strains, five isolates (*Bacillus sp.*, *Acinetobacter sp.*, *Enterobacter sp.*, *Serratia sp.*, *Stenotrophomonas sp.*) were able to produce lipase above 11U/ml. *Acinetobacter sp.*, were selected for further studies.

Table 3. Lipase enzyme production in submerged fermentation of selected bacterial strains

S.No	Strains	Lipase Activity(U/ml)
1.	<i>Bacillus sp.</i> ,	12
2.	<i>Staphylococcus sp.</i> ,	09
3.	<i>Escherichia coli</i>	05
4.	<i>Micrococcus sp.</i> ,	08
5.	<i>Klebsiella sp.</i> ,	04
6.	<i>Serratia sp.</i> ,	13
7.	<i>Proteus sp.</i> ,	10
8.	<i>Pseudomonas sp.</i> ,	10
9.	<i>Enterobacter sp.</i> ,	12
10.	<i>Aeromonas sp.</i> ,	07
11.	<i>Pseudomonas sp.</i> ,	11
12.	<i>Micrococcus sp.</i> ,	05
13.	<i>Acinetobacter sp.</i> ,	15
14.	<i>Lactobacillus sp.</i> ,	09
15.	<i>Stenotrophomonas sp.</i> ,	14

3.5. Strain Identification with 16S r RNA

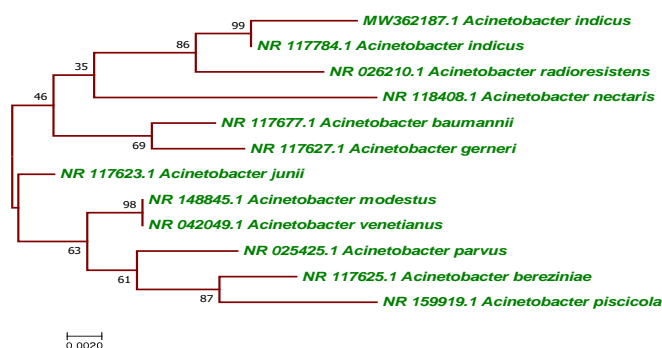
3.5.1. Characteristics of lipase producing bacteria

The enzyme assay revealed that among forty-eight isolates, five isolates (*Bacillus sp.*, *Acinetobacter sp.*, *Enterobacter sp.*, *Serratia sp.*, *Stenotrophomonas sp.*) were found to be more potent for the production of lipase enzyme. The sequences were deposited in Genbank with accession number for *Acinetobacter indicus* was MW362187 (Mellmann *et al.*, 2006).

3.5.2. Phylogenetic analysis of the isolates

The phylogenetic tree of isolates belonging to genera was constructed by Maximum Likelihood method. Scale bar of *Acinetobacter indicus* (0.0020 substitutions/site with identity 99%).

Fig.No:2 - Phylogenetic Tree – *Acinetobacter Indicus*



3.6. Effect of various physico – chemical parameters

Maximum lipase activity was observed at 40°C by *Acinetobacter indicus*. Some microorganisms such as *Bacillus cereus* shows, 30°C as the optimum temperature for growth and production of lipase. Effect of temperature

on lipase activity was optimized by incubated in different temperature (30°C - 45°C) and obtained 13.07U/g ds at 40°C in 48hrs. (Nema *et al.*, 2019).

Maximum activity of pH 8 was noted for *Acinetobacter indicus*. Bacteria prefer a pH around 7 for best growth and lipase production. (Barbaro *et al.*, 2001). Akila Rajan and Jayakumaran Nair, 2011, reported that maximum lipase enzyme activity was observed at pH 8.5, for both submerged and solid-state fermentation. The changes in pH of medium depend on products of metabolism, medium composition and end – product of their metabolism. (Ire Sopuschukwe Francis *et al.*, 2018).

Acinetobacter indicus obtained maximum activity at 24hrs with 17.96U/ml. Chaudhary and Sharma 2020 reported that *Aspergillus costaricaensis* obtain maximum lipase activity at 48hrs of the incubation period. Activity of lipase decreases in longer incubation period due to accumulation of toxic end products, nutritional depletion and pH changes in medium. Production of enzyme begins from the second day of fermentation and maximum lipase activity was observed in log phase of growth (Falony *et al.*, 2006).

Table 4. Effect of incubation period on enzymatic activity

Incubation periods	<i>Acinetobacter indicus</i> (U/ml)
12hrs	13.06±0.3
18hrs	14.93±0.7
24hrs	17.96±0.7
48hrs	14.86±0.8
60hrs	10.2±0.5

Glucose as sole carbon sources for *Acinetobacter indicus*. Glucose is beneficial carbon sources that enhance the maximum production of cold active lipase. Carbon sources provide maintenance of cellular functions and growth such as enzyme production. Carbon source from saccharides influenced the enzyme production. (Soleymani *et al.*, 2017).

Peptone confirmed the maximum activity of nitrogen sources. Peptone as nitrogen sources increased the production of lipases in *Pseudomonas* LSK25. (Leelatulasi Salwoom *et al.*, 2019). CaCl₂ obtained maximum activity of lipase for *Acinetobacter indicus* with 27U/ml, Metal ions form complexes ionized fatty acids and alter the solubility and behavior at oil – water interfaces, so inhibit enzyme activity. The addition of Ca²⁺ resulted in maximum production of lipase in 2% (Salwoom *et al.*, 2019).

Table 5. Effect of metal ions on lipase activity

Metal ions	<i>Acinetobacter indicus</i> (U/ml)
MgSO ₄	24.9±0.36
CaCl ₂	27±0.5
FeSO ₄	17.96±0.35
NaCl ₂	16.06±0.6
ZnCl ₂	16±0.5

Maximum activity in presence of olive oil. Olive oil acts as an inducer for lipase production in *Bacillus mycoides* and *Bacillus coagulans* (Thomas *et al.*, 2003). The presence of olive oil with OD of 1.00 produce maximum lipase activity of 9.50 (EU/ml) (Pallavi *et al.*, 2014).

Tween 80 showed increased activity of lipase in *Acinetobacter indicus* (30U/ml). The presence of surfactants in medium enhances the secretion of protein (Pallavi *et al.*, 2014), by altering the permeability of cell membranes. Dalmau *et al.*, 2000 reported that the presence of Tween 80 in the medium was obtained best inducer for production of lipase in *Candida rugosa*.

Table 6. Effect of surfactants on lipase activity

Surfactants	<i>Acinetobacter indicus</i> (U/ml)
Tween 20	14.03±0.45
Tween 40	10.2±0.52
Tween 60	19.16±0.56
Tween 80	30±0.2
Tween 100	21.13±0.3

Triton X-100	16±0.5
SDS	12.06±0.3

3.7. Optimization of medium for lipase production by the statistical approach

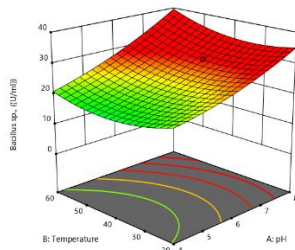
Optimum concentration of individual factors was obtained by subjecting the significant factors to central composite design (CCD). Using RSM the final medium optimization and interaction among the significant factors were studied. Similar report studied by Kiran *et al.*, 2010. *Pseudomonas aeruginosa* AT10 produce rhamnolipids, by optimizing the culture media using RSM. (Abalos *et al.*, 2002).

Table 7 ANOVA for response surface model of *Acinetobacter indicus*

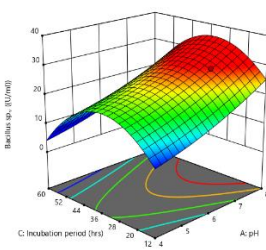
Term	Lipase activity
F value	33.32
Prob> F	<0.0001
Std.Dev.	1.22
Mean	19.35
R ₂	0.9898
Adj.R ₂	0.9601
Ade. precision	18.1589

Fig. No:3 Response surface graph – *Acinetobacter indicus*

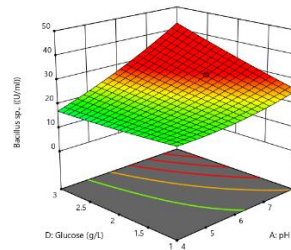
Interaction effect of pH and Temperature



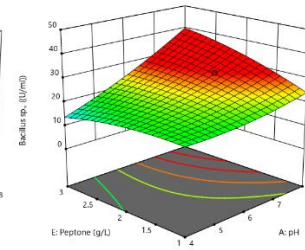
Interaction effect of pH and Incubation period



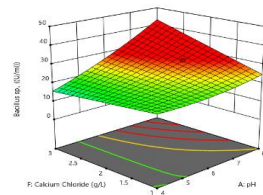
Interaction effect of pH and Glucose (P = 0.0359)



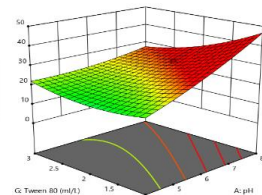
Interaction effect of pH and Peptone



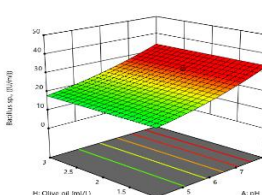
Interaction effect of pH and Calcium chloride



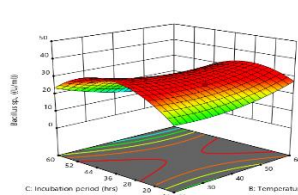
Interaction effect of pH and Tween 80



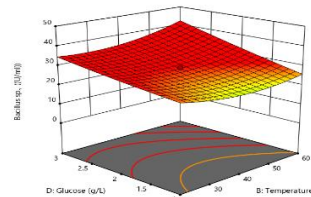
Interaction effect of Olive oil and pH



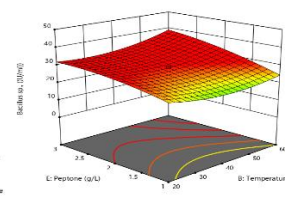
Interaction effect of Temperature and Incubation period



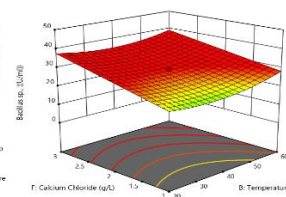
Interaction effect of Temperature and Glucose



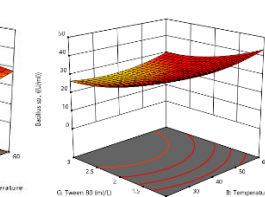
Interaction effect of Peptone and Temperature



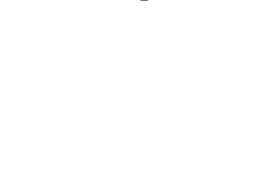
Interaction effect of CaCl₂ and Temperature



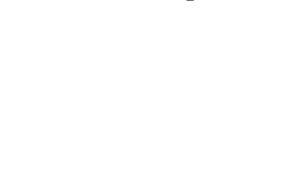
Interaction effect of Tween 80 and Temperature



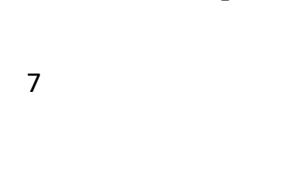
Interaction effect of Olive oil and Temperature



Interaction effect of Glucose and Incubation period

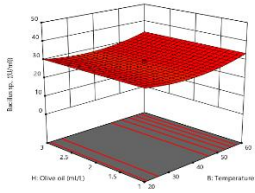


Interaction effect of Peptone and Incubation period

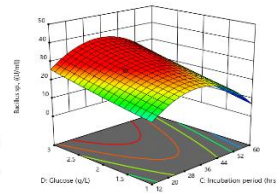


Interaction effect of CaCl₂ and Incubation period

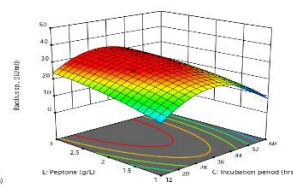




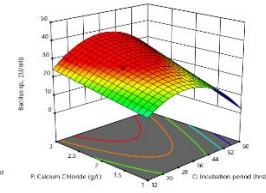
Interaction effect of Tween 80 and Incubation period



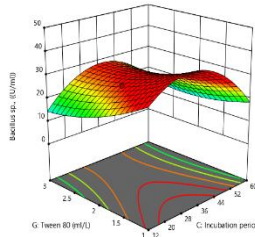
Interaction effect of Olive oil and Incubation period



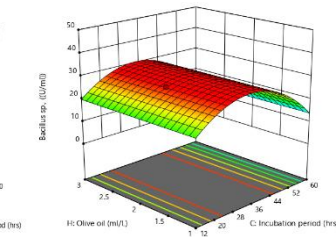
Interaction effect of Peptone and Glucose



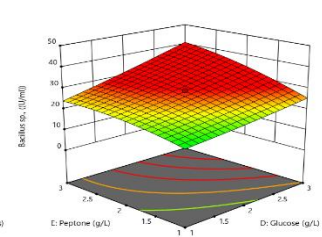
Interaction effect of Glucose and CaCl₂



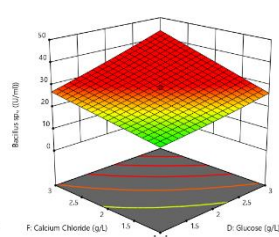
Interaction effect of Tween 80 and Glucose



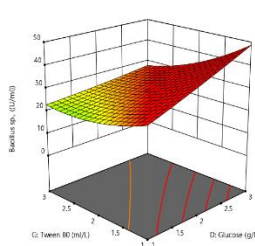
Interaction effect of Olive oil and Glucose



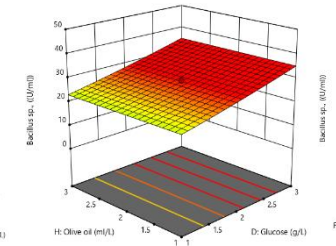
Interaction effect of CaCl₂ and Peptone



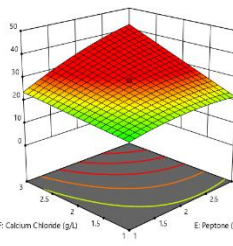
Interaction effect of Tween 80 and Peptone



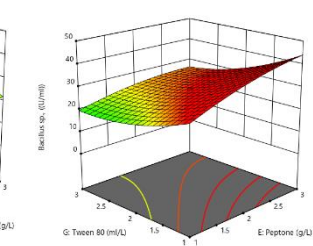
Interaction effect of Olive oil and Peptone



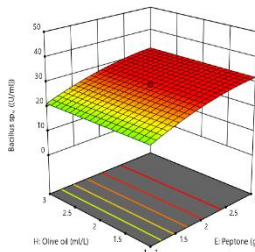
Interaction effect of Tween 80 and CaCl₂



Interaction effect of CaCl₂ and Olive oil



Interaction effect of Olive oil and Tween 80



3.8. Purification of lipase

The enzyme was purified by Ammonium sulphate precipitation, dialysis, DEAE – cellulose column chromatography and SDS – PAGE.

3.8.1. Ammonium sulfate precipitation

Maximum production of lipase was 47U/ml obtained by *Acinetobacter indicus* with 40% concentration of Ammonium sulfate precipitation. Similar results obtained by Massedeh *et al.*, 2012. *Bacillus stearotherophilus* shows 40% of Ammonium sulfate precipitation produce high amount of activity, with 1.03 purification fold.

Table 8. Lipase activity in Ammonium Sulfate precipitation

	<i>Acinetobacter indicus</i>
--	------------------------------

Concentration of Ammonium sulfate	Lipase Activity (U/ml)	Protein (mg/ml)
30%	45	0.21
40%	47.58	0.40
50%	41	0.30
60%	40.5	0.24
70%	39	0.22

3.8.2. Dialysis

Dialysis has increased the activity to 47.58U/ml in *Acinetobacter indicus*

3.8.3. DEAE – Cellulose column chromatography

DEAE – Cellulose column chromatography has increased the activity to 55U/ml for *Acinetobacter indicus*.

In *Acinetobacter indicus*, the crude enzyme showed the specific activity of 72.31U/ml, Ammonium sulphate precipitation has increased the activity to 75.80U/ml with purification fold of 1.048 and yield of 67.14. Dialysis has increased the activity to 92.18U/ml with a purification fold of 1.21 and yield of 42.14%. DEAE – Cellulose column chromatography has increased the activity to 720U/ml with purification fold of 7.81 and yield of 20.5%. The result was tabulated in table no 9. Joseph *et al.*, 2011, reported that DEAE – Cellulose column chromatography obtained 35.64-fold purification with specific activity of 93.38U/mg.

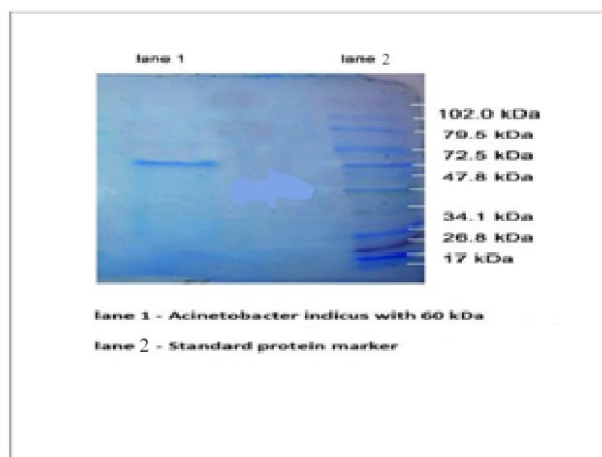
Table 9. Purification of lipase enzyme from *Acinetobacter indicus*

Fractions	Total volume (ml)	Activity (U/ml)	Protein (mg/ml)	Total protein (mg)	Total activity (U)	Specific Activity (U/ml)	Purification (Fold)	Recovery (%)
Culture filtrate	100	35	0.484	48.4	3500	72.31	1	100
Ammonium sulphate	50	47	0.310	31.0	2350	75.80	1.048	67.14
Dialysis	25	59	0.200	20.0	1475	92.18	1.21	42.14
DEAE cellulose chromatography	10	72	0.100	1.0	720	720	7.81	20.5

3.9. SDS – PAGE

The molecular weight of lipase produced by *Acinetobacter indicus* was 60 KDa. Lipase isolated mostly from *Aspergillus* sp., have molecular weight of 25 to 70KDa reported by Contesini *et al.*, 2010. Due to glycosylation of purified protein separated by SDS – PAGE. Lipase from *E. nidulans* NFCCI3643 have a molecular weight around 54KDa (Lanka and Latha, 2015). Hu *et al.*, 2018 observed the molecular weight of protein in SDS – PAGE was 51KDa.

Figure 4. Determination of molecular weight (SDS-PAGE)



4. Conclusion

Oil contaminated soil has a potential application in detergent industries. The enzyme was optimized by various physical parameters. Maximum lipase activity was observed at 40°C at pH 8. Glucose and peptone as sole sources for maximum production of lipase. CaCl₂ and Tween 80 showed increased activity of lipase in *Acinetobacter indicus* (30U/ml), Partial purification of enzyme with ammonium sulfate precipitation, dialysis and DEAE Cellulose column chromatography produced the activity of 720U/g with purification fold of 7.81 and a yield of 20.5%.

5. Conflicts of interest

The authors of the present work report that there is no conflict of interest in this work.

6. References

1. Abalos, A., Maximi, F., Manresa, M.A and J. Bastida, (2002). Utilization of response surface methodology to optimize the culture media for the production of rhamnolipids by *Pseudomonas aeruginosa* AT10. *Journal of chemical Technology and Biotechnology: International Research in process, Environmental and clean Technology*, **77**(7): 777 – 784.
2. Akila Rajan, and Jayakumar Nair, (2011). Study on alkaline lipase production by a newly isolated *Aspergillus fumigates* MTCC 9657 in submerged and solid – state fermentation using economically and industrially feasible substrate. *Turkish Journal of Biology*, **35**(5): 569 – 574.
3. Aravindan, R., Anbumathi, P. and T. Viruthagiri, (2007). Lipase applications in food industry. *Ind. J. Biotechnol*, **6**: 141 – 158.
4. Bisht, D., Yadav, S.K., Darmnal, N.S., *et al.*, (2013). Optimization of immobilization conditions by conventional and statistical strategies for alkaline lipase production by *Pseudomonas aeruginosa* mutant cells: scale up at bench – scale bioreactor level. *Turkish J Biol*, **37**: 392 – 404.
5. Borkar, P.S., Bodade, R.G., Rao., S.R. and C.N. Khobragade, (2009). Purification and characterization of extracellular lipase from a new strain *Pseudomonas aeruginosa* SRT 9. *Braz J Microbiol*, **40**(2): 358 – 366.
6. Chaudhary, P and P. Sharma, (2020). Optimization of cultural parameters of *Aspergillus costaricaensis* CBS115574 for lipase production by solid state fermentation. *Asian Journal of Microbiology and Biotechnology*, **4**(3): 108 – 118.
7. Chauhan, M., Chuhan, R.S and V.K. Garlapati, (2013). Evolution of a new lipase from *Staphylococcus* sp., for detergent additive capability. *Biomed. Res. Int.* 374967.
8. Corleone, J. (2018). What are the benefits of lactose – free milk? Available in: [https:// healthy eating. Sfgate.com / benefits – lactosefree – milk- 7284.html](https://healthy.eating.sfgate.com/benefits-lactosefree-milk-7284.html).
9. Dahiya, P. and S. Purkayastha, (2011). Isolation, screening and production of extracellular alkaline lipase from a newly isolated *Bacillus* sp., PD – 12. *J Biol Sci*, **11**: 381 – 387.
10. Dahiya, P., Arora, P., Chaudhury, A., Chand, S and N. Dilbaghi, (2010). Characterization of an extracellular alkaline lipase from *Pseudomonas mendocina* M – 37. *Journal of Basic Microbiology*.**50**(5): 420 – 426.
11. Dogan, B. and K.J. Boor, (2003). Genetic diversity and spoilage potential among *Pseudomonas* sp., isolated from fluid milk products and dairy processing plants. *Appl Environ Microbiol*, **69**:130 – 138.
12. Falony, G., Armas, J.C., Mendoza, J.C.D. and J.L. Martinez – Hernandez, (2006). Production of extracellular lipase from *Aspergillus niger* by solid – state fermentation. *Food Technology and Biotechnology*,**44**(20): 235 – 240.
13. Gunawan, E.R., Basri, M., Mohd, B.A.R., Abu, B.S., Raja, N.Z.A. R, (2005). Study on surface methodology (RSM) of lipase – catalyzed synthesis of palm – based wax ester. *Enzyme and Microbial technology*, **37**: 739 – 744.
14. Gupta, R., Gupta, N. and P. Rathi, (2004). Bacterial lipases: An overview of production, purification and biochemical properties. *Appl Microbiol Biotechnol*, **64**: 763 – 781.
15. Hashemi, M. *et al.*, (2013). Comparison of submerged and solid-state fermentation systems effects on the catalytic activity of *Bacillus* sp., KR – 8104 amylases at different pH and temperature. *Ind crop prod*, **43**: 661 – 667.
16. Iftikhar, T., Niaz, M., Afzal, M., Haq, I and M.I. Dan Rajoka, (2008). Maximization of Intracellular lipase production in a lipase – over producing mutant derivative of *Rhizopus oligosporus* DGM 31: A Kinetic study. *Food Technol. Biotechnol.* **46**(4): 402 – 412.
17. Iso, T., Kikuta, H., Nagamori, E., Honda, H., Ogini, H., Ishikawa, H. and T. Kobayashi, (2001). Lipase production in two – step fed – batch culture of organic solvent – tolerant *Pseudomonas aeruginosa* LST – 03. *J BiosciBioeng*, **91**: 245 – 250.

18. Ji, Q., Xiao, S., He, B. and X. Liu X, (2010). Purification and characterization of an organic solvent-tolerant lipase from *Pseudomonas aeruginosa* LX1 and its application for biodiesel production. *J Mol Catal B Enzym*, **66**: 264-269.
19. Joyruth, P and Lali Growther, (2018). A review on microbial lipase production, purification, characterization and its applications in various fields. *Asian journal of microb. Biotech.Env. Sc.* 20(1): 294 – 300.
20. Joyruth. P and Lali Growther, (2020). Molecular identification, production and optimization of lipase from oil contaminated soil using submerged fermentation. *J.Pure . Appl. Microbiol.*, 14(1): 341 – 349.
21. Lajis, A.F.B. (2018). Realm of thermoalkaline lipases, in bioprocess commodities Lajis. *Hindawi J Lipids*, 1 – 22.
22. Lanka, S and J.N.L. Latha, (2015). A short review on various screening methods to isolate potential lipase producers: Lipases – the present and future enzymes of Biotechnological industry. *International Journal of Biological chemistry*, **9**(5): 207 – 219.
23. Lowry, O.H., Rosebrough, N.J., Farr, A.L. and R.J. Randall, (1951). Protein measurement with the folin phenol reagent. *J. Biol. Chem.* **193**: 265 – 275.
24. Mahima, G., Hajela, M and G.P. Pandey, (2016). Screening, Identification, characterization and production of bacterial lipase from oil spilled soil. *Int. J. Curr. Microbiol. App. Sci.* **5**(3): 745 – 763.
25. Mehta, A., Sharma, R. and R. Gupta, (2019). Statistical optimization by Response surface methodology to enhance Lipase production by *Aspergillus fumigatus*. The open Microbiology Journals.**13**: 86 – 93.
26. Mellmann, A., Backer, K., Fiff, C., Keckevoet, U., Schumann, P and D. Harmsen, (2006). Sequencing and *Staphylococci* identification. *J. Emerg. Infect. Dis.* **12**(2): 333 -336.
27. Menoncin, S., Domingues, N.M., Freire, D.M.G, Toniazzo, G., Cansian, R.L. and J.V. Oliveira, (2008). Study of the extraction, concentration and partial characterization of lipases obtained from *Penicillium verrucosum* using solid – state fermentation of soybean bran. *Food and Bioprocess Technology*, Doi: 10 – 1007/S 11947 – 008 – 0104 – 8.
28. Mobarak – Qamsari, E., Kasra – Kermanshahi, R. and Z. Moosavi – Nejad, (2011). Isolation and Identification of a novel, lipase – producing bacterium, *Pseudomonas aeruginosa* KM110. *Iran J Microbiol*, **3**: 92 – 98.
29. Najjar, A., Robert, S., Guerin, C., Violet – Asther, M. and F. Carriere, (2011). Quantitative study of lipase secretion, extracellular lipolysis and lipid storage in the yeast *Yarrowialipolytica* grown in the presence of olive oil. Analogies with lipolysis in humans. *Applied Microbiology and Biotechnology*, **89**: 1947 – 62.
30. Nema, A., Patnala, S.H. and S.K. Devaraj, (2019). Production and optimization of lipase using *Aspergillus niger* MTCC 872 by solid – state fermentation. *Bulletin of the national Research centre*, **43**(8). <https://doi.org/10.1186/s42269> - 019 - 0125 – 7.
31. Pallavi, R., Uma, T. and D. Nitin, (2014). Post – harvest fungal diseases of fruits and vegetables in Nagpur. *Int J of Life Sciences*, A2: 56 – 58.
32. Rakesh Kumar, Arpit Sharma, Arun Kumar. and Deepak Singh, (2012). Lipase from *Bacillus pumilus* RK31: production, purification and some properties. *World Appl Sci J*, **16**(7): 940 – 948.
33. Reis, P., Holmberg, K., Watzke, H., Leser, M.E. and R. Miller, (2009). Lipases at interfaces: A review. *Adv Colloid Interface Sci*, **147 – 148**: 237 – 250.
34. Salwoom, L., Abd Rahman, R.N.Z.R., Salleh, A.B., Shariff, F.M., Convey P., Pearce, D. and M.S. Mohamad Ali, (2019). Isolation, characterization and lipase production of a cold – adapted bacterial strain *Pseudomonas* sp., LSK 25 isolated from Signy Island, Antarctica. *Molecules*, **24**(4):715.
35. Soleymani, S., Alizadeh, H. and M. Rabbani, (2017). Efficient media for high lipase production: one variable at a time approach. *Avicenna J of Medical Biotechnology*, **9**(2): 82 – 86.
36. Tamura, K., Stecher, G., Peterson, D *et al.*, (2013). MEGA 6: Molecular evolutionary genetics analysis version 6.0. *Mol. Bio. Evol.* **30**: 2725 – 2729.
37. Thomas, A., Mathew, M., Valsa, A.K., Mohan, S. and Manjula, (2003). Optimization of growth conditions for the production of extracellular lipase by *Bacillus mycoides*. *Indian Journal of Microbiology*, **43**(1): 67 – 69.
38. Treichel, H., Oliveira, D.D., Mazuti, M.A., Luccio, M.D and J.V. Oliveira, (2010). A review on microbial lipases production. *Food Bioproc Technol*, **3**: 182 – 196.
39. Venil, C.K., Nnthalakumar, K., Karthikeyan, K and P. Lakshmanaperumalsamy, (2009). Production of L – asparaginase by *Serratia marcescens* SBOS: Optimization by response surface methodology. *Iranian J. Biotechnol.* **7**: 10 – 18.