

**Development of Eco-friendly Technique for  
the Bioclarification  
of  
Highly Turbid Drinking Water  
In  
Mining Region of Goa  
Using  
Freshwater Yeast Cultures**

A thesis submitted to Goa University for the award of the degree of

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**In BOTANY**

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Goa University, Goa

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**September, 2022**

## **CERTIFICATE**

This is to certify that **SHEELA PAL** has satisfactorily completed the thesis entitled “**Development of Eco-friendly Technique for the Bioclarification of Highly Turbid Drinking Water In Mining Region of Goa Using Fresh Water Yeast Cultures**” submitted to Goa University for the award of the degree of **DOCTOR OF PHILOSOPHY IN BOTANY** is a record of original and independent work carried out during the period of February 2016 – 2022 in the **DEPARTMENT OF BOTANY, SCHOOL OF BIOLOGICAL SCIENCES**

**GOA UNIVERSITY** under my supervision and that it has not previously formed the basis for the award of any Degree, Diploma, Associateship or Fellowship or any other similar title to any candidate of this or any other university. I affirm that the thesis submitted by **SHEELA PAL** is completely independent research work carried by her under my supervision.

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It is hereby declared that the thesis entitled “**Development of Eco-friendly Technique for the Bioclarification of Highly Turbid Drinking Water In Mining Region of Goa Using Fresh water Yeast Cultures**” submitted to Goa University for the award of the degree of DOCTOR OF PHILOSOPHY IN BOTANY is a record of original and independent work carried out during the period of February 2016 – 2022 in the DEPARTMENT OF BOTANY, GOA UNIVERSITY under the supervision of Dr. NANDKUMAR MUKUND KAMAT, and that it has not previously formed the basis for the award of any Degree, Diploma, Associateship or Fellowship or any other similar title to any candidate of this or any other university.

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**Figure 3.60** Phylogeny analysis of *Candida tropicalis*.

## ABBREVIATIONS

NTU	Nephelometric Turbidity unit
gm	Gram
L	Liter
mm	Millimeter
Tem	Temperature
rpm	Revolution per minute
Mm	Milimole
F.d.	Fractal dimension
MEA	Malt extract agar
ABR	Aclian blue retention
YMB	Yeast mediated biosedimentation
qt	Adsorption efficiency
Ce	NTU at equilibrium
Co	Initial concentration
Cf	NTU final
SMTTW	Simulated mine tailing turbid water

This thesis is organized into four chapters. Chapter I consists of a brief introduction to the work presented, a review of literature, and objectives. Chapter II includes materials and methods involved. Chapter III consists of results obtained, and finally Chapter IV presents a discussion with reference to important results, concluding with prospects / scope for future work. At the end are placed associated sections such as a summary, a bibliography of literature cited, appendices regarding media, reagents, stains used, papers published / communicated in peer-reviewed journals, papers presented at national / international conferences, courses, workshops / other conference participation.

**CHAPTER I**  
**Introduction**  
**Review Of Literature**  
**and**  
**Objective**

# INTRODUCTION

The present work attempts to address the problem of pollution of drinking water resources, in the iron ore mining region of Goa.

## **1.1. Pollution of drinking water resources**

Water on the earth is one of the important needs of living being. While 1.1 billion people live without clean drinking water, 2.6 billion people lack adequate sanitation in the world (Source: World water Council, 2022,).

In India approximately 77 million people lack access to safe water, 769 million lack adequate sanitation (Source: water.org, 2022). Around the world, the water quality in mining area is degrading in response to mining activity.

Organic solvents, petroleum products, and heavy metals from disposal sites or storage facilities can migrate into aquifers. Pesticides and fertilizers. Runoffs of mine dump can be carried into lakes and streams by rainfall runoffs or snowmelt, or can percolate into aquifers (US Environmental Protection Agency., 2021).

30% people of Goa get polluted water encouraging health problems like diarrhoea, dysentery mainly among children and the aged. The presence of excess raw iron is responsible for neurodegenerative diseases (George., 2010).

### **1.1.1. Open cast mining causes pollution in natural water bodies**

The water sources on which we all rely are being threatened more and more by human activities like mining. Water pollution from mine wastewater dumped into the environment, seepage from tailings and waste rock impoundments, and heavy water use in ore processing are all effects of mining that have an impact on fresh water.

Open cast mines affect both surface, water body and agriculture field (Okolo., et al 2018).The environmental impact of mining operations that were conducted with little regard for the environment is becoming more widely known. We have occasionally paid a very high price for using minerals in our daily lives. By its very nature, mining uses up, diverts, and potentially dangerously pollutes water resources. For the purification of surface water there is a need to develop cost effective, easier and eco-friendly process.

**Table. 1.1 Nomenclature related to ecofriendly sedimentation technique**

Nomenclature	Definition	Reference
Bioclarification	<p>Clarification is an essential step in a water or wastewater treatment process to remove suspended solids through gravity settling, providing a clarified liquid effluent.</p> <p>The process to remove suspended solids by biological means followed by gravity settling, is known as bioclarification.</p>	<p><a href="https://www.evopedia.com/en/markets/applications/clarification/">https://www.evopedia.com/en/markets/applications/clarification/</a>.</p>
Biosorption	<p>Both absorption and adsorption are described by the term "sorption". The assimilation of a substance from one state into another is called absorption (i.e., liquids being absorbed by a solid or gases being absorbed by water). Ions and molecules physically bind to the surface of the solid material through adsorption. In this instance, the solid surface is the adsorbent and the material that has gathered at the interface is the adsorbate. In biosorption, a biological matrix is used as the sorbent.</p>	<p>Izabela et al., 2013</p>
Bioadhesion	<p>The phenomena of natural and artificial materials adhering to biological surfaces is referred to as bioadhesion.</p>	<p>Manuel et al., 2012</p>
Bioflocculation	<p>The term "bioflocculation" refers to a method where flocculants are mediated by microbes or biodegradable macromolecular flocculants produced by microorganisms.</p>	<p>Zayad et al., 2018</p>
Biosedimentation	<p>Sedimentation is the process in which particles separate from a liquid because of gravity.</p> <p>Biosedimentation is the process in which particles separate from a liquid mediated by biological agents followed by gravity.</p>	<p>Haan (2020)</p>

## 1.2. Problems caused by colloidal Turbidity

Turbidity is the cloudiness or haziness of a fluid. Turbidity in water results from the presence of colloidal particles that scatter light. As a result, objects in water become indistinct. The turbidity of water measured with Turbidity meter, is calculated based on the amount of light scattered by particles in the water column.

Turbidity of water is a prevalent problem these days near industrial regions, such as mining and textile industries. Chemical and physical coagulants are now widely used to treat turbidity. Chemical procedures are unfriendly to the environment, whereas physical approaches cover a vast region. The use of microorganisms to treat turbidity hasn't been much applied. Previously, bacterial and fungal strains were utilized to remove turbidity, but they were found to be low range tolerant and produce a large amount of biomass, respectively (Keenan., 2000, Nazareth et al. 2001).

Soil colloids with size range 0.001 $\mu$ m to 1  $\mu$ m are complex mixtures of organic and inorganic entities (Everett., 1972) and causes turbidity in solution (Sakhawoth et al., 2017).

**Table.1.2 Classification of textural classes**

S.N.	Textural Class	Particle size diameter (mm)
1	Sand	0.02 - 2.0 mm
2	Silt	0.002 - 0.02 mm
3	Clay	0.002 mm

### 1.2.1. Suspended particulate matter (SPM)

More the particles present in water, more will be the light scattered. Hence, turbidity and total suspended solids are related.

## LITERATURE REVIEW

### 1.3. Goa

Goa is an important Indian state having natural magnificent beaches, a plateau, estuaries, and mangrove environment, as well as a wide range of plant and animal species. It is rich in minerals such as iron, manganese, bauxite, limestone, dolomite, refractory clays, limonite sand, steatite, silica sand, feldspar, graphite, talc, quartz, soapstone, etc. Due to the abundance of minerals, the mining sector in Goa is vital to the state's economy (Nayak., 2002).

Mining industry in Goa is mainly centered in four talukas: Bicholim, Sattari, Quepem and Sanguem, accounting for around 16 percent of the overall geographical area (AEQM.,1997). Aside from development, mining has irreversibly harmed forests, agriculture and water resources. Ore was retrieved from turbid pits below the water table in some mines using water pumped from the pits. Pumped water has the potential to be detrimental to both biotic and human life. Metals such as Fe, Mn, Cr, Pb, Ni, and Zn are adsorbed on clay and transported from mines (Nayak et al., 1995). During the rainy season, mines discharge runoff into local water bodies, causing turbidity. The Selaulim and Bicholim rivers were clogged with silt, while the Zuari and Mandovi rivers were poisoned with arsenic (levels as high as 50 micrograms/gram due to mining activity) (Nayak et al., 1995). Approximately 90% of Goa's iron and ferromanganese is carried to the Mormugao seaport via these two rivers. The Mollem National Park, the Bhagwan Mahaveer animal sanctuary, and protected regions in Bondla, Neturlim, and Cotigao wildlife sanctuaries are among the sensitive zones where mining is steadily making inroads (AEQM., 1997).



# ELECTORAL MAP OF GOA STATE

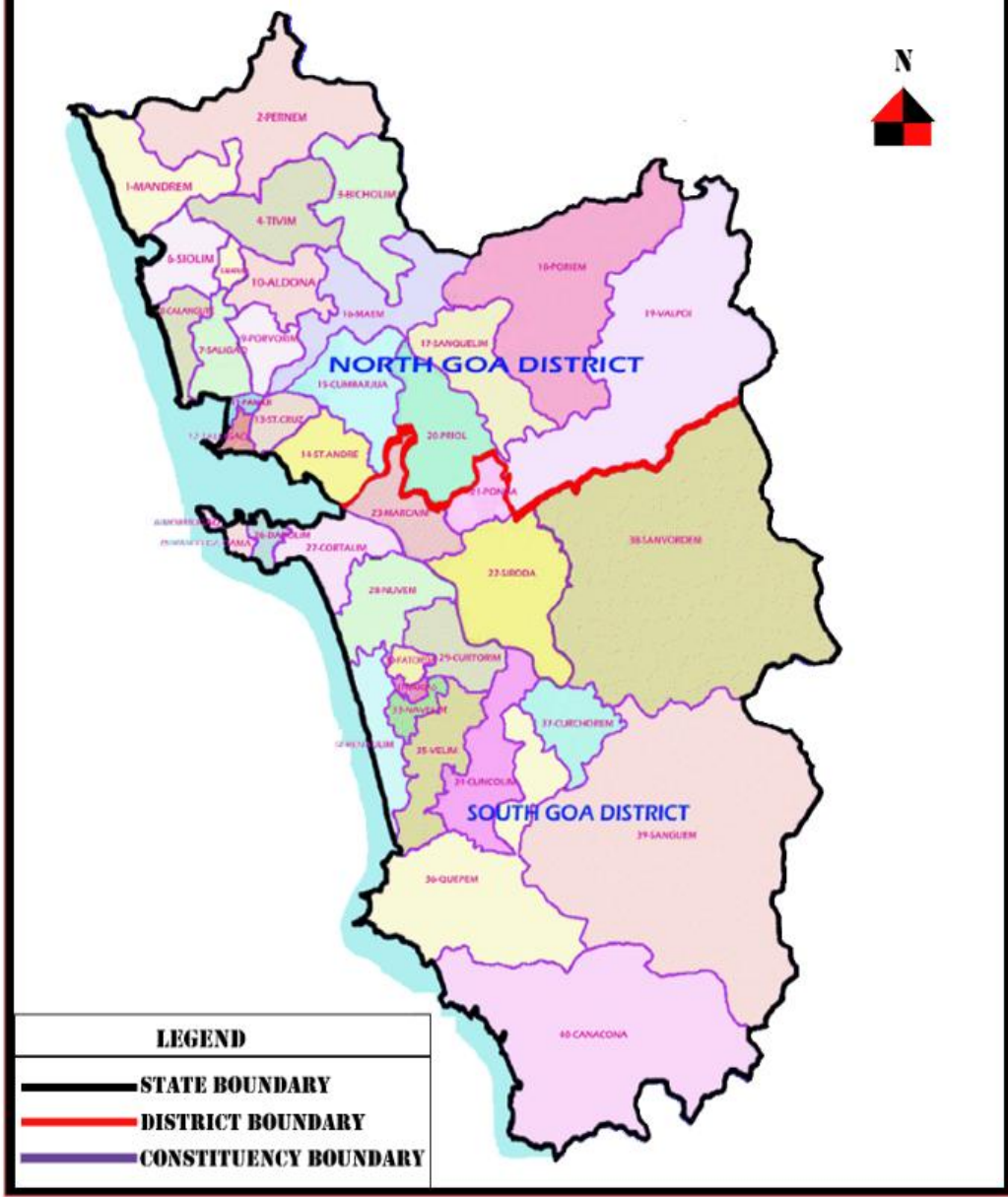


Figure.1.1: Map of Goa

(Image courtesy: <https://ceogoa.nic.in/appln/ui/GoaMap.aspx>)

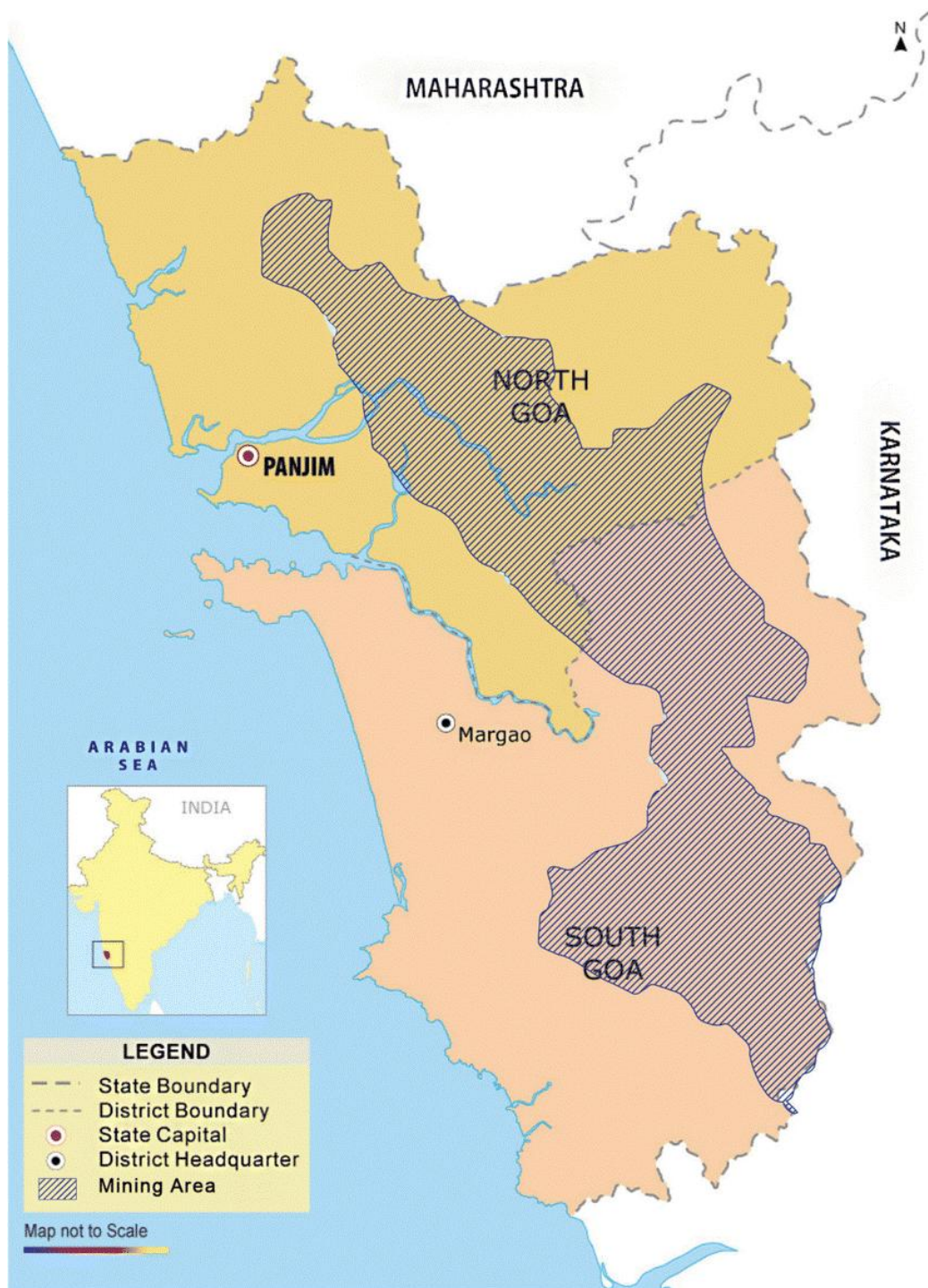


Figure.1.2: Mining area of Goa (Image courtesy: Sarupria et al., 2018)

### **1.3.1. Freshwater bodies in Goa**

Goa is blessed with a plethora of freshwater bodies including lentic and lotic e.g. natural springs, lakes, ponds, waterfalls and streams. The water reservoirs having low salinity concentration are known as freshwater bodies.

### **1.3.2. Present status of Fresh water bodies in Goa**

Freshwater bodies have a lot of biodiversity. It depicts seasonal changes in the environment (Udayashankara.,2015). Water quality monitoring using physiochemical indicators reflect contemporary water quality. However biotic parameters created in recent years have proven to be a great tool for predicting water quality, trophic level evolution and pollution status. Biological monitoring could be a good and affordable way to assess the effects of environmental influencers (Robert 1974). The abundance of phytoplanktons and other bacteria has an impact on trophic levels in the environment (Krishnamurthy et al., 2000). Many bacteria can be extracted from fresh water bodies for potential application in human wellbeing. Nearly all of the freshwater bodies in the area are affected by mining operations.

The Western Ghats's Freshwater Biodiversity Assessment was conducted by the IUCN Global Species Programme's Freshwater Biodiversity Unit in collaboration with the Zoo Outreach Organization (ZOO) to review the global conservation status and distribution of 1,146 freshwater species belonging to four taxonomic groups: fishes (290 taxa), mollusks (77 taxa), odonates (171 taxa) and aquatic plants (608 taxa) in response to this need for information and raised awareness (Molur., 2011).

During summer, the temperature of the water in Goa's temple tanks was higher than usual (Pariolkar, 2014). The presence of yeasts in seasonal freshwater bodies in Goa has received little attention. The ideal time for this type of fungi to recuperate is during the monsoon season (Naik KS., 2016). There is a scarcity of information about freshwater biodiversity.

**Table 1.3 Water parameters values in freshwater bodies of Goa**

<b>Mg<sup>+2</sup></b>	<b>Fe<sup>+3</sup></b>	<b>Cl<sup>-1</sup></b>	<b>Ca<sup>+2</sup></b>	<b>Hardness</b>	<b>Alkalinity</b>	<b>pH</b>	<b>Temp (°C)</b>	<b>Parameter</b>	<b>Water body</b>
120.9	0.32	218.7	169.3	317.4	166.1		28.5	Vikrant et al., 2015	Vaddem lake
13.9	0.46	30.7	20.1	34.1	59	6.56	28.6	-do-	Mayem lake
49	0.40	123.3	51.5	103	95.6	6.66	28.7	-do-	Santacruz lake
52.9	0.4	65.4	67.3	123.8	168.9	7.38	25.5	-do-	Someshwa tank
-	-	-	-	-	-	6.20	28	Sawaiker et al., 2016	Syngenta lake
-	-	--	-	-	-	6	25	-do-	Khandola Pond
-	-	-	-	-	-	6.3	25.5	-do-	Lotus lake
-	-	-	-	-	-	6.72	26.2	-do-	Curtorim lake

NO <sub>3</sub> <sup>-1</sup>	DO	TDS	TSS	SO <sub>4</sub> <sup>-2</sup>	PO <sub>4</sub> <sup>-3</sup>
	-	-	-	225	0.478
-	-	-	-	84.8	0.27
-	-	-	-	103.2	0.43
-	-	--	-	170.1	0.37
0.20-0.7	8-12	600	-	-	0.1
0.4	9	45	-	-	0.07
2.5	6.60	616	-	-	0.01-0.7
1.43	9.19	1210	-	-	0.49

\*Temp=Temperature ,°C All values are in mg/L except temperature, electrical conductivity,pH

### 1.3.3. Natural factors affecting water quality in Goa

Natural factors, affecting water quality are related to the hydrography and the topography of the region. Lakes, ponds, ditches, water reservoirs, tanks etc receive pollutants from catchment area and are less diverse in pollutants as compared to rivers, stream, creeks, brooks, etc.

Stream frequency are often subjected to a greater degree of pollution. More the amount of runoff making its way into the water, more will be the increase in pollutant diversity.

The intensity of rainfall varies in Goa from place to place. Rainfall is the main cause of aquatic pollution of water bodies as it causes runoff from sand, laterite and bauxite in mining areas.

The topography of Goa is highly undulating with vast areas of uneven terrain. Mining activity is considered as the primary cause of altering the natural topography in some mining areas of Goa. Due to resulting heavy siltation, the health of water bodies is degrading (Nayak., 2002).

Physico-chemical analysis of seven riverine water samples from Bicholim taluka carried out by Nayak (1995), show that Velguem and Harvalem rivers exhibit a marked seasonal variation in pH and TSM as they run through active mine areas. Locals in some rural areas of Goa use tank water (natural spring water) for drinking, and they suffer greatly during the wet season due to lack of clean drinking water, as it gets contaminated with run offs of silt or from mine dumps.

Water treatment is a critical issue in the mining industry because it is used in the mineral ore treatment process. Mine dump runoff pollutes water by contaminating it with heavy metals, producing leaching, erosion, and sedimentation. Clay, colloids, and minerals are commonly found in mine effluent (Ghose et al., 2000), which increase water turbidity and hence have an influence on public health.

Studies are being conducted to remove colloidal turbidity caused by mining operations. Heavy metal contamination and its effects on the environment and living creatures are being examined by research institutions in various parts of Goa. Chemical, physical, and biological approaches are commonly employed to eliminate colloidal turbidity.

Some chemical flocculants used in water purification such as polyacrylamide are non-biodegradable. Chemical flocculants also produce colours in treated water effluent and build fragile flocs that are difficult to settle (Bhatti et al., 2009).

Physical wastewater treatment technologies have been criticised for having problems such as sludge settleability, a big footprint, a long retention time and high-volume inefficiency (Odegard et al., 1990). Heavy metals are being removed from aqueous solutions using polymer-based nano composites (Guixia et al., 2018). Wang (2018) examined the research on removing heavy metals from wastewater using nanoscale zero valent iron. Metal organic framework compounds can be thought of as an alternative

method to get heavy metals out of water (Paulina et al., 2018). Yihan (2018) examined the work on heavy metal removal from wastewater utilizing novel materials such as metal organic framework.

Because of their economic viability and ecological superiority, biological techniques have got a lot of attention. Microbial species are being employed as sustainable ways in the biotreatment of very turbid water resources, utilizing the ability of diverse species of bacteria, fungi, algae, and other microorganisms to aid the flocculation process and remove turbidity. When it comes to reducing colloidal turbidity, coagulation and flocculation have an advantage. Microbial treatment plants take up less room and are environmentally favourable. Microbes work by contacting positively charged particles with negatively charged microbial cell surfaces, resulting in the formation of clumps, that settle down quickly (Hattori et al., 1970; Verspagen., 2006).

#### **1.4. Composition of muddy water and soil in mining area and mine tailing soil**

Goa is rich in natural mineral wealth. Iron ore, Bauxite, China clay and silica sand are important deposits. Iron ore and manganese ore have resulted from the residual concentration of banded manganiferrous quartzites and phyllites respectively. Mining reserves are prominent in Bicholim, Sanguem and Sattari. Important mineral formations in Goa is significantly observed in the Vageri formation, Bicholim formation, Sanvordem formation and Barcem formation (Nayak et al., 1995).

**Table 1.4 Types of clays and their composition in Iron ore mining (Nayak et al., 1995).**

Name of clay	Colour	Percentage of Metal Elements (W/W )
Intrusive	Pale Pink with yellow spots	Fe: 18-25; Al <sub>2</sub> O <sub>3</sub> : 30-35; Mno: 0.02- 0.5 SiO <sub>2</sub> : 10-15
Lateritic	Brown pink	Fe: 40- 45; Al <sub>2</sub> O <sub>3</sub> : 20- 25; Mno: 0.25- 0.5 SiO <sub>2</sub> :10-20
Limonitic	Yellowish orange	Fe: 45- 56; Al <sub>2</sub> O <sub>3</sub> : 14- 17 Mno: 0.25- 0.5; SiO <sub>2</sub> : 7-9
Manganiferous	Black, yellow, brown sticky with oily appearance	Fe: 35- 43; Al <sub>2</sub> O <sub>3</sub> : 5- 10 Mno: 5- 12; SiO <sub>2</sub> : 5- 9
Phyllitic	Pink	Fe: 11- 12; Al <sub>2</sub> O <sub>3</sub> : 17 Mno : traces; SiO <sub>2</sub> : 25- 30

**Table 1.5 Physical Properties of natural soils dump waste and mine tailings dump (Nayak et al., 1995)**

Soil Sample	Particle Size Analysis			Bulk Density g/cm <sup>3</sup>	Particle Bulk Density g/cm <sup>3</sup>	Porosity%	Water holding capacity %
	Sand%	Silt%	Clay %				
Natural Soil	51.2	24.7	20.3	0.96	1.9	49.48	44.1
Dump Waste (Sandy Loams)	56.9	21.9	18.1	1.3	2.8	53.5	40.6
Tailings (Sandy Clay Loam)	48.6	25.0	26.1	1.7	3.0	44.4	31.7

**Table 1.6 Metals found in agricultural soil**

**from mining and non mining areas (Ratha et., al 1994)**



Metal found in agricultural soil affected by mining				Metal found in agricultural soil non affected by mining			
Element	Concentration	Element	Concentration	Element	Concentration	Element	Concentration
Na %	0.35	Ca %	0.208	Na %	0.275	Ca %	0.274
Mg %	0.77	Ti %	1.581	Mg %	1.035	Ti %	1.295
Al %	17.92	Mn %	0.360	Al %	17.405	Mn %	0.247
Si %	45.688	Fe %	31.924	Si %	43.00	Fe %	34.824
P %	0.147	Ni ppm	1.422	P %	0.162	Ni ppm	1.345
K %	1.045	Pb ppm	0.641	K %	1.168	Pb ppm	0.642
Cr %	0.253	Zn ppm	0.210	Cr %	0.204	Zn ppm	0.188
Co%	0.087	-	-	Co%	0.066	-	-

### 1.5. Rainfall as a factor for transportation of mine tailing dump in water bodies

Goa received annual rainfall of 3663.9 mm in 2016, 3443.4 mm, in 2017, 2671.3 mm in 2018 (IMD, <https://hydro.imd.gov.in/hydrometweb>) so average annual rainfall of Goa is approximately 3500 mm (Kamal et al., 2015). During rainy season, mine dump or ore runoff enters into the freshwater bodies of these areas. The high concentration of ore tailing cause turbidity in the water bodies. According to analytical reports of mining companies and IBM Nagpur, suspended solids were in the range of 102 to 4212 mg/L in the river stream nallahs of Goa (AEQM, 1997).

### 1.6. Conventional and natural methods for removal of colloidal turbidity

Coagulation or flocculation, sedimentation, and sand filtration, followed by disinfection, are some of the traditional methods for removing turbidity from water (Raus et al., 2016). Chitosan and its derivatives are utilised as flocculants in a variety of applications (Yang et al., 2016). Among all chemical-based flocculants, such as ferric chloride, aluminium sulphate, copper sulphate, and ferrous sulphate; Al-based flocculants are frequently used because they can change the surface's charge property, causing colloidal particles to coagulate and form flocs (Wue et al., 2007). However, aluminum residues have been linked to Alzheimer's disease in studies (Xu et al, 2014).

**Table.1.7 Turbidity removal efficiency of different natural plant coagulants**

Name of coagulants from plant	Turbidity removal efficiency (%)	References
<i>Okra mucilage</i>	98.7%	Ani et al., 2012
Bean seed	80%	Antov et al., 2012
<i>Moringa oleifera</i> seeds	97%	Azni et al., 2006
Date palm rachis	95%	Mhenni., 2010
<i>Hibiscus rosa sinensis</i>	99%	Jinisha et al., 2017
Chestnut and acorn	90%	Skirbic et al., 2009
Cactus and hyacinth beans peels	89.03 %, 77.10 %	Girish., 2012
<i>Plantago ovata</i>	95.6%	Ramavandi et al., 2014
Water hyacinth ( <i>Eichhornia crassipes</i> )	99.5% Cr (IV) removal	Shaha et al., 2017 Shaha et al., 2017

Natural coagulants, based on plants researched for removing turbidity from wastewater, produce huge biomass, making it difficult to remove from drinking water. Seaweeds have been examined for their biosorption qualities due to their adsorbent properties and the fact that they are readily available in large quantities (Davis et al., 2000). Cellulosic algal cell walls contain carboxylate, amine, imidazole, phosphate, sulphhydryl sulphate and hydroxyl moieties (Christ et al., 1981). The decrease of NTU of colloidal turbidity by *Plantago ovate* seed extract has been researched, and displayed a substantial reduction rate of NTU after doping with NaCl (Ramesh et al., 2017). Table 1.7 contains a list of natural coagulants as reported by various studies.

### 1.7. Algae as bioremediation agents

Because algae undergo photosynthesis, they transform solar energy into useful biomass and absorb nutrients like nitrogen and phosphorus, creating eutrophication. Microalgae are an important part of the aquatic food chain because they produce C and N. (Lange et al., 1993). In wastewater treatment, the bio-treatment of sewage using algae is particularly appealing. The use of algae to clean wastewater has been studied for over forty years, with Oswald et al. (1957) reporting one of the first descriptions of this application. In the presence of *Lemanea*, *Stigeoclonium* and other algae, wastewater treatment is mediated by a combination of nutrient intake, higher pH and high dissolved oxygen concentration. *Micrasterias*, *Staurastrum*, *Pinnularia*, *Meridionand*, *Surirella*,

and other *Micrasterias* species have been extensively used in this area (Al-Homaidan et al., 2014).

Algae have the ability to remove large amounts of nitrogen, sulphur, and phosphorus from the environment (Chevalier et al., 2000; Mulbry et al., 2008; Su, Mennerich, & Urban, 2012; Woertz et al., 2009). The presence of *Botryococcusbraunii sp.* was reported in wastewaters contaminated with nitrogen, phosphorous, sulphur, ammonium phosphate and nitrate (An et al., 2003; Salloum et al., 2002), as it blooms when these elements are present in high concentrations in wastewater (An et al., 2003; Salloum et al.,) *Chlorella sp.* removed phosphate from wastewater and decreased nitrates in amines (Wang et al., 2010). Walker et al. (1975). Semple et al. (1996) investigated *Prothecazopfi* for the degradation of aromatic compounds in oil and crude oil reduction. The bioremediation process of *Chlorella spp.* has been researched in both live and dried (frozen) cell stages (Wang et al., 2011).

Ribson (1998) studied the techniques for immobilizing algae cells for improved wastewater treatment performance and conducted an experiment using *Chlorella emersonii* for removal of phosphorous from wastewater. Chen (2003) investigated the marine microalgae *Isochrysis galbana* and successfully immobilised the algae to manage the water quality using clam cultures cultured in alginate beads for long-term storage. *Chlorella vulgaris* and the macrophyte *Lemnaminuscule* were used to clean wastewater containing high quantities of organic materials, such as ethanol and citric acid by-products (Valderrama et al., 2002). Photosynthetic bacteria and green microalgae have also been found to have the ability to remove nutrients. Under aerobic dark heterotrophic conditions, bacterial species, such as *Rhodobacter sphaeroides* and *Chlorella sorokiniana* were found to efficiently extract nutrients (Ogbonna et al., 2000). Andriana et al., (2012) cultivated freshwater and marine water algae to assess their potential for wastewater treatment and biofuel production and discovered that their growth rate is unrelated to the strain's nitrogen removal capacity. In 55 days, microalgae were able to remove 66.98 percent of chloride from chlorinated wastewater (Yang et al., 2016). The essential elements for the cultivation of microalgae for wastewater treatment, abiotic and biotic, operative conditions, and bioreactor design were examined by Jeon de la et al., (1992). The primary problems in successful waste remediation by algae are

how to preserve the mixed consortium's long-term effectiveness and homeostasis. (Goncalves et al., 2016).

**Table.1.8 Algae reported for pollutant**

Algal Species	Bioremediation targets	References
<i>Prototheca zopfii</i>	Crude oil	Walker et al., 1975
Diatoms	Naphtalene	Cerniglia et al., 1980
<i>Phormidium</i> , <i>Oscillatoria</i> , <i>Nostoc</i> and <i>Synechococcus</i>	Hydrocarbons	Abed et al., 2002
Algal mixed consortium <i>Chlorella</i> and <i>Scenedesmus</i>	N, P.	Shene et al., 2016
<i>Spirulina</i>	Domestic wastewater treatment	Laliberte et al., 1997
<i>Phormidium sp.</i>	N, Orthophosphate	De la Nou et al., 1988
<i>Coelastrum proboscideum</i>	Pb (II)	Mac hardy et al., 1980
<i>Cladophora glomerata</i>	Zn	Vymazal et al., 1984
<i>Dunaliella sp.</i>	pyrethroid insecticide Deltamethrin	Baeza-Squiban et al., 1990
<i>Chlamydomonas reinhardtii</i>	Cu, Hg, Cd, Pb	Bayramo et al., 2006
<i>Monoraphidium braunii</i>	bisphenol	Gattullo et al., 2012
<i>Euglena</i> , <i>Oscillatoria</i> , <i>Chlamydomonas</i> , <i>Scenedesmus</i> , <i>Chlorella</i> , <i>Nitzschia</i> and <i>Navicula</i>	Water pollution Indicator, Cr (IV) and Cl removal	Al-Homaidan et al., 2012; Alfonso., 2018

### 1.8. Bacterial Coagulants for Removal of Colloidal Turbidity of Wastewater

The bioremediation properties of several chemo heterotrophic bacteria were investigated. Bacterial colonies grow quickly and are easily genetically modified. Furthermore, extremophiles are important in decomposing dangerous substances or in killing other species in the environment. They are easily able to thrive in locations that are heavily polluted (Ahmad et al., 2012; Brim et al., 2000). *Pseudomonas*, *Rhodococcus*, and *Acinetobacter* are some of the bacterial genera used most often. (Shourian et al., 2009). Exopolymer binding has been shown to immobilize lead in a variety of bacteria, including *Staphylococcus aureus*, *Micrococcus luteus* and *Azotobacter species* (Maier et al., 2009). Many bacteria are lithotrophic in addition to adsorption of chemicals. They manufacture a variety of hazardous substances (Nakajima et al., 2008). Some bacteria enable for easy maintenance and collection of organisms due to their biofilm-forming capabilities (Radwan et al., 2002). The degradation or removal of wastewater constituents such as organic and inorganic components, such as ammonium and phosphorus, is a common subject of bacterial bioremediation research. The bacterial communities

operating in wastewater treatment plants, according to Wagner & Loy (2002), were linked with Betaproteobacteria, Alphaproteobacteria, Gammaproteobacteria, Bacteroidetes, and Actinobacteria. Ammonia oxidizers Betaproteo and Gammaproteo bacteria, in particular, are aerobic chemolithoautotrophs. In freshwater habitats, certain Alphaproteobacteria taxa, particularly *Nitrobacter* and *Nitrospira*, are known to oxidise nitrite to nitrate (Schmidt et al., 2003).

In order to eliminate colloidal turbidity from surface water, Wasify (2015) used *Bacillus licheniformis*, *Bacillus insolitus*, and *Bacillus alvei* bacterial species to create exopolysaccharides.

Bacterial exopolysaccharides, which operate as a natural coagulant, exhibited measurable efficacy in the treatment of surface water. *Bacillus sp.* exopolysaccharide was utilised as a flocculating agent (Kanmani., 2017). Heavy metal removal from wastewater has been studied using a *Klebsiella varriicola* bacterial strain isolated from the textile industry (Aslam et al., 2017). Peptidoglycan, carboxyl, amine, and phosphonates are components of bacterial cell walls that are thought to aid contaminant biosorption from wastewater (Vijay et al., 2004). Bacterial strains have been isolated from activated sludge from local hoggery displayed chromium removal and turbidity removal efficiency at 28% and 90% respectively (Zhang et al., 2012). Olanira (2009) studied bacterial strain coagulant property. Marine bacterial isolates assessed for turbidity removal of kaoline clay and reduction of COD of dairy industry flocculation efficiency have been observed as 99.7% and 93.9%, respectively (Onukwuli., 2012). *Rhodopsuedomonas sphaeroides* was used for the flocculation of coal slurry (Zhang et al., 2012). *Arthrobacter sp.* isolated from Tuyme river, South Africa was studied for flocculant production on lactose and urea as a sole source of nitrogen and carbon and showed flocculating activity of 75.4% and 83.4%, respectively (Mabinya et al., 2012). *Pseudomonas aeruginosa* and *Bacillus subtilis* showed great potential for heavy metal adsorption (Alencar., 2017). *Fusarium solani*, *Bacillus* and *Arthrobacter* have been used for the efficient treatment of mine tailing turbidity on a lab scale (Nazareth et al., 2001). The biosorption study conducted by Ansari et al., (2007) using isolates isolated from agriculture soil irrigated with industrial wastewater contaminated with toxic metals

showed resistance to metals. Under anaerobic conditions, *Desulfovibrio* spp., a sulfate reducing bacteria, can cause metal precipitation (Roane and Pepper, 2000).

Challenges were encountered during the large scale production of bacterial strains, for effective biosorption or flocculation. Bacterial strains were found to be pH specific, and could not tolerate pH variation, osmoregulation and physical robustness (Keenan., 2000).

**Table1.9 List of some Metal adsorbing Bacteria**

<b>Bacteria</b>	<b>Heavy metal</b>	<b>References</b>
<i>Aeromonas caviae</i>	Cd, Cr (IV)	Loukido et al., 2004
<i>Alcaligenes eutrophus</i>	Cd	Mahvi et al., 2004
<i>Aphanothece halophytica</i>	Zn	Incharoensakdi et al., 2002
<i>Bacillus firmus</i>	Pb, Zn, Cu	Salehizadeh et al., 2003
<i>Bacillus licheniformis</i> , <i>Escherichia coli</i> , <i>Pseudomonas fluorescens</i>	Cd, Pb, Zn	Basha et al., 2014
<i>Bacillus licheniformis</i>	Cu, Cr, Fe	Samarth et al., 2012
<i>Bacillus licheniformis</i>	Cr (IV)	Zhou et al., 2007
<i>Bacillus subtilis</i> , <i>Micrococcus</i>	Cu	Nakajima et al., 2001
<i>Bacillus thuringiensis</i>	Cr (IV)	Sahin et al., 2005
<i>Bacillus thuringiensis</i>	Ni, Hg	Ozturk et al., 2007
<i>Corynebacterium glutamicum</i>	Pb	Choi et al., 2004
<i>Nostoc muscorum</i>	Cr (VI)	Gupta et al., 2008
<i>Ochrobactrum anthropi</i>	Cd	Ozdemiret al., 2003
<i>Pseudomonas aeruginosa</i>	Cu, Pb	Kazyet al., 2002; Lin et al., 2006
<i>Pseudomonas cepacia</i>	Cu	Savvaidis., et al., 2003
<i>Pseudomonas putida</i>	Pb, Cu	Pardo et al., 2003
<i>Pseudomonas</i>	As	Nitish et al., 2018
<i>Sphaerotilus natans</i>	Cu	Beolchini.,et al., 2006
<i>Sphingomonas paucimobilis</i>	Cd	Tangaromsuk., et al., 2002
<i>Staphylococcus xylosus</i>	Cd, Cr (IV)	Ziagovaet al., 2007
<i>Streptomyces coelicolor</i>	Cu	Ozturk et al., 2005
<i>Streptomyces pimprina</i>	Cd	Puraniket al., 1995
<i>Streptomyces rimosus</i>	Fe (III)	Selatniaet al., 2004

## 1.9. Fungi in Wastewater Treatment

Since the fungi kingdom is very diverse with both unicellular yeast and branching hyphae, it produces several remarkable reproductive structures. Fungal biomasses are reported highly efficient for the sequestration of Zn (II) and Pb (II) from wastewater (Aftab. et al., 2017). Much attention has been paid to the removal of heavy metal from wastewater via fungal strains as they have high percentage of cell wall materials, which increases the possibility of the presence of a number of functional groups involved in binding the metal from wastewater (Gadd. 1998; Dhankhar. 2011). They can be easily applied practically because a large number of fungal strains are reported as non-pathogenic (Dhankhar., 2011). Many fungal species such as *Rhizopus*, *Aspergillus*, *Streptoverticillum*, *Saccharomyces* etc. have been explored for biosorption of heavy metals (Volesky et al., 1990). *Trichoderma* sp. have been studied for the absorption of copper from wastewater (Saxena., 2006). Krauss (2011) reviewed the work done for mineralization of xenobiotic compounds to carbondioxide via fungal species. Fungal strains *Fomitopsis meliae*, *Trichoderma ghanense* and *Rhizopus microsporus* isolated from gold mining sites in South Africa, and studied for their metal tolerance efficiency were found to show remarkable tolerance in heavy metal rich media (Oluwatosi., 2016). Fungi have been studied for enzymatic treatment to remove chemicals from water (Becker. 2017). Various species of terrestrial, wood and soil basidiomycetes show outstanding mineralization of xenobiotics, while aquatic fungi have yet to be studied (Krauss et al., 2011). First report on fungal wastewater treatment had been reported by Curtis 1969. Yoshizawa in the end of 1970 drew attention for the use of yeast in wastewater treatment.

Furthermore, researchers have found yeasts produce lipids (Chung., 2016) glycolipids (Yang., 2013) and enzymes. Therefore, it is widely used in the treatment of high concentration organic wastewater, heavy metal ions wastewater and domestic sewage.

### 1.10. Challenges for implementation of fungal biomass at field level

Despite the fact that fungal biomasses have been investigated for the removal of heavy metals from wastewater, field application remains a barrier. The mechanism of bioremediation and technological constraints could be the major reasons behind this (Dhankhar et al., 2011).

Direct microbiological application is also tricky. As a result, several aspects of the implementation of fungal biotreatment of wastewater must be investigated. There are physiochemical aspects to be considered, such as pH, as well as the major difficulty of biomass immobilization in wastewater treatment plants. Because it creates a large volume of sludge, improvements in reuse methods should be investigated and optimized.

**Table 1.10 List of metal adsorbing Fungi**

<b>Fungi</b>	<b>Heavy Metals Resistant</b>	<b>References</b>
<i>Trichoderma atroviride</i>	Cu, Zn, Cd	Lopez et al., 2013
<i>Trichoderma atroviride</i> , <i>Mortierella exigua</i>	Zn, Ba, Fe	Malina et al., 2005
<i>Aspergillus niger</i> , <i>Trichoderma asperellum</i> , <i>Penicillium simplicissimum</i>	Cu, Pb	Burton et al., 2003
<i>Trichoderma</i> , <i>Aspergillus</i> , <i>Mortierella</i> , <i>Paecilomyces</i> , <i>Penicillium</i> , <i>Pythium</i> and <i>Rhizopus</i>	Co	Ross et al., 1986
<i>T. virens</i> (PDR-28)	As, Cu, Cd, Ni, Pb, Zn	Babu et al., 1998
	<b>Metal adsorbent</b>	
<i>Aspergillus brasiliensis</i> , <i>Penicillium</i>	Cu, Mn, Zn	Pereira et al., 2014
<i>Aspergillus niger</i>	Pb, Cd, Cu, Ni	Kapoor et al., 1999



<b>Fungi</b>	<b>Heavy Metals Resistant</b>	<b>References</b>
<i>Aspergillus niger</i> , <i>Aspergillus flavus</i> ,	Pb, Ni, Cr (V)	Dwivedi et al.,2012
<i>Aspergillus terreus</i>	Ni, Fe, Cr (V)	Dias et al., 2007
<i>Aspergillus terreus</i>	Pb, Cd	Joshi et al., 2011
<i>Gliocladium roseum</i>	Cd	Massaccesi et al., 2002
<i>Mucor rouxii</i>	Pb, Cd, Ni, Zn	Yan., 2008
<i>Rhizopus arrhizus</i>	Zn	Zhou 1999
<i>Rhizopus delemar</i>	Cu, Co, Fe	Tsekova et al, 2002
<i>Rhizopus oligosporus</i>	Cd	Aloysius et al., 1999
<i>Saprolegnia delica</i> , <i>Trichoderma viride</i>	Zn (II), Pb (II),	Ali et al., 2007
<i>Talaromyces helicus</i>	Cd	Massaccesi et al., 2002
<i>Trichoderma longibrachiatum</i>	Cr(V)	Joshi et al., 2011
<i>Trichoderma viride</i>	Cd	Joshi et al., 2011
<i>Trichoderma asperellum</i>	Pb	Zhu et al., 2018
<i>Aspergillus niger</i> , <i>Rhizopus</i> , <i>Penicillium</i>	Petroleum pollutant	Duniya et al., 2018

### 1.11. Yeast

Yeast is a polymorphic, unicellular creature. It keeps its round, ovoid, or cylindrical shape. It belongs to the ascomycetous and basidiomycetous fungal groups.

Yeast is divided into two types: active yeast and non-active yeast.

I. Fermented yeast (Baker's yeast, for example) is a type of yeast that can only ferment six-carbon sugar into alcohol and carbon dioxide (Mohmd., et al 2017). *Candida sp.* is an example of oxidised yeast, which has high oxidation ability. The chemical makeup of the cell wall changes as yeast develops (Lebrun et al., 2018). Mannan, glucan, and mannoproteins are thought to be the main components of *Saccharomyces cerevisiae's* cell wall (Northocote et al., 1952). Calcium is essential for flocculine activation (Hattori et al., 1970). Older cell forms flocculate earlier and is stronger than younger cell flocculate, according to Herrere et al. (1991). Younger cells lack flocculin and have lower cell sizes.

### **1.11.1. Aquatic natural yeast**

The biodiversity of aquatic environmental specifically freshwater yeast community haven't been well studied (Yorkov, et al 2017). A study by Libkind et al. (2017) looked at the diversity and ecology of yeasts in freshwater habitats (temperate and tropical rivers, lakes, and lagoons). In terms of conformity, yeast identification was mostly based on differences in morphological and physiological features. As such the identification, was difficult in several cases with inconclusive results (Kurtzman & Robnett.,1998). Morphological and physiological testing were introduced in 2001. After the use of the more trustworthy molecular data in identification was introduced, most studies focused on identifying yeasts (Libkind et al., 2017).

### **1.11.2. Yeasts in Rain Water**

Microorganisms in the air, clouds, or precipitation have not yet been given a clear definition of their makeup. Despite the fact that there have been several studies on bacterial communities, bacterial strains from the clouds were isolated, and their role in shaping the atmosphere's chemical makeup was investigated (Tina et al, 2015). After expressing bacteria in an active form from an inactive state in a suitable environment, Morris et al. (2008) showed atmospheric dispersal may play a significant influence in bacterial biogeography patterns. Bacterial strains are shielded from UV radiation and given nutrients to ensure their survival in fog and clouds (Tina et al., 2015). Clouds facilitate long-distance microbial movement (Griffin 2006). In 2002, Baurer and colleagues recovered bacterial and fungal spores from cloud water.

### **1.11.3. Application of yeast for removal of colloidal turbidity**

Yeasts are unicellular eukaryotic organisms but at times they form hyphae or pseudo-hyphae according to environmental conditions. Aquatic yeast abundance and diversity depend on the water reservoir whether it is fresh water or marine water source (Hagler et al., 1987). Yeasts are polyphyletic members of the kingdom fungi's phyla ascomycetes and basidiomycetes.

According to Mueller and Schimt (2007), as the kingdom fungus was identified only 40 years ago and its variety is understudied, only 7% of the total species have been

described, making it critical to document the undescribed, particularly aquatic species (Shearer., 2007).

Smirnov (1964) proposed introducing yeast into lakes as fungal propagules through inflowing stream rain water and wind. Freshwater yeast ecology, such as distribution patterns, diversity, and importance in aquatic systems, has yet to be well researched (Carlos et al., 2012).

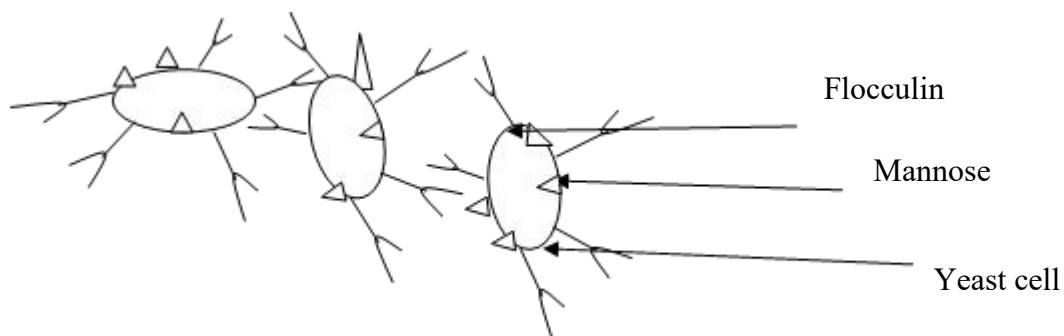
Monika N. discovered black yeast in tap water in 2016. *Slovenian Exophiala sp.* and *C. parapsilosisa* white yeast. Garg discovered *Rhodotorula graminis*, a red coloured yeast, from fresh water flowing through the mining area of Goa in 2011. Chin (2015) isolated 43 species of yeast from the sea surface microlayer and underlying water on Taiwan's northern coast using a polyphasic molecular method. Currently there are approximately 12000 yeast species (Hawkworth 1991), among which few have been studied for the cleaning of wastewater. Many industries use full scale bioclarification wastewater treatment plants based on this process (Yoshizawa et al., 2014). Many studies of yeasts associated with polluted water have been performed eg. *Candida*, *Rhodotorula* screened to find out the pollution level (Slavík et al., 1998).

Niljana et al., (2017) extracted yeasts from pharma industry effluent to breakdown cefdinir. *Candida* spp. had the highest cefdinir degradation efficiency of 84 percent. Zhang Fan (2015) discovered that mixed strains of bacteria and yeast are more efficient than single strains of bacteria and yeast for wastewater treatment. Bread yeast was reported to be an efficient natural coagulant for removing 99.6% colloidal turbidity from synthetic turbid water (Northocote 1952). *Saccharomyces cerevisiae* and *Torula sporadel brueckii*, utilised to treat pharmaceutical effluents, showed the highest percentage reductions of 52.5, 52.5, and 58.7%, for BOD, COD, and nitrate respectively (Abiye et al., 2015). The yeast nanocomposite system was built utilising a mixture of hydrophilic and hydrophobic nano-silicates and yeast cell mixture to grow yeast cells in a hydrocarbon polluted aqueous medium to breakdown the hydrocarbon (Klymenko et al., 2017).

*Candida tropicalis* showed reduction of copper levels in the growth medium by nearly 80%. (Rahman et al., 2007). In studies involving *Fusarium solani* and *Bacillus arthrobacter* for efficient treatment of mine tailing on a lab scale at Goa University (Nazareth et al. 2001; Deshpande, 1990), fungal strains were found to be more efficient. Biosorption may be primarily a function of the binding of metal cations to chemical functional groups on the yeast cell wall via ionic and coordinate bonds (Brady et al., 1994). Particles having opposite charges were attached to the surface of microbial biomass (Len et al., 2006).

Soh (2022) studied dye removal approach using spent brewer yeast and concluded that adsorption was not affected by changing the operating temperature. Chaudhary et al., 2022 found an excellent removal of Zn, Pb, Cd and Ni with the help of fungal consortia i.e. 80%, 70%, 80% and 75% respectively.

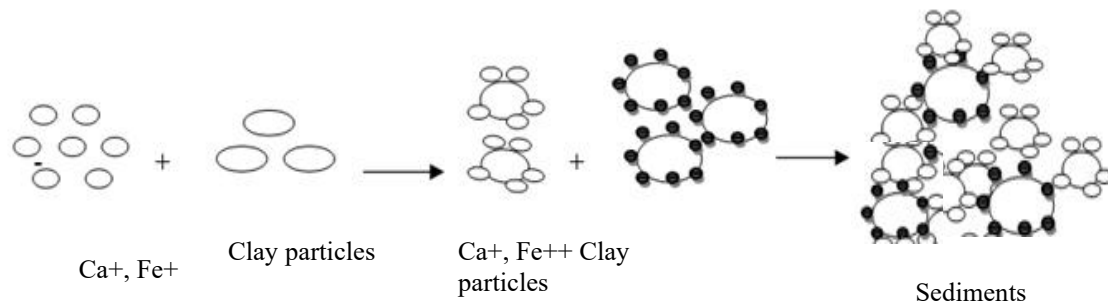
Yeasts have been examined for a variety of purposes. For example, Emila (1979) used yeast to assess the nitrogen budget in estuarine sediments. *Rhodotorula* strains, isolated from soil, were studied for lead reduction efficiency with phosphate, reported a removal of 98% of lead cations (Tian et al., 2022). *Sterigmatomyces halophilus* reported 100% acid orange dye removal by (Ali et al., 2021). Aibeche et al., (2021) reported aquatic yeasts for removal of toxic lead. Artifon et al., (2022) used spent brewer yeast for removal of textile dyes and attained 80-90% dye removal efficiency.



**Figure.1.3: Mechanism of flocculation of cells**

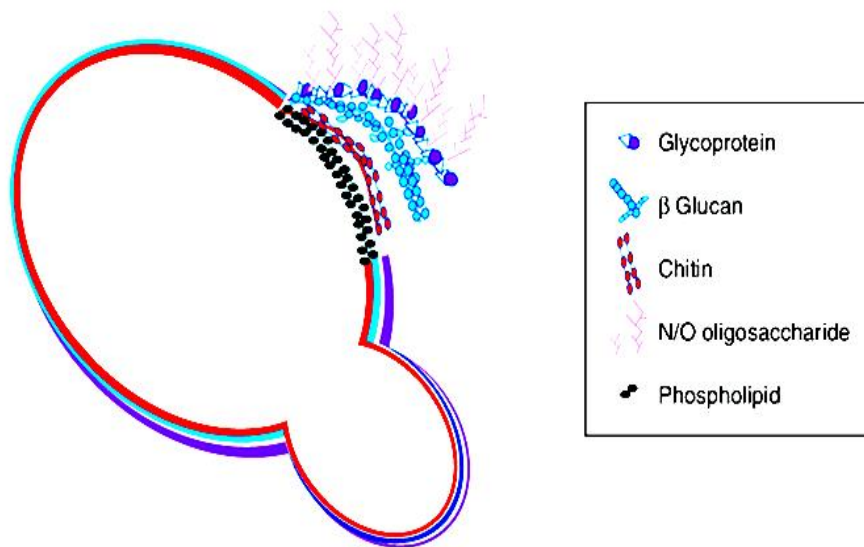
The primary characteristic of water clarifying by yeast is based on the unique structure of the microbial cell surface, morphology, and composition of the cell wall, all of which

must be carefully studied in order to achieve maximum performance (Verstrepen., 2004). Aside from cell-cell adhesion, yeast cell walls interact with their environment and have exceptional abiotic surface attachment abilities. This would reduce sludge generation, the treatment plant's ecological footprint, hydraulic retention time, and capital costs, as well as assist rural consumers. Yeast strains remove a lot of turbidity while reducing sludge generation.



**Figure.1.4: Mechanism of adhesion of clay minerals on microbial cell surface and sedimentation**

According to Oh et al., (2009), the main source of metal biosorption is ionic interaction and complex formation between metal cations and acidic sites on the microorganism's surface. Because the potential of freshwater yeast in bioremediation has been underutilized, particularly in removing colloidal particles from turbid water, yeast's flocculating qualities may play a significant role in bioremediation. Millions of yeast cells are present in 1ml of yeast suspension ( $1 \times 10^7$  number of yeast cell in yeast suspension with absorbance 1), are capable of binding clay colloids to its surface. Predicted mechanism of cell mediated biosedimentation is electrostatic force between anions and cations present on the surface of yeast and clay colloids respectively



**Figure.1. 5: Yeast wall composition demonstration**

The fungal cell wall can compose approximately 40% of the total volume and range in thickness from 0.1 $\mu$ m to 1.0  $\mu$ m. The main component of the cell wall of most fungi is chitin and  $\beta$  (1)-glucan are surrounded by a gel like substrate comprised primarily of  $\alpha$  (1)-glucans and galactomannan proteins (Hasim et al., 2019).

After thorough literature survey, it has been observed that yeasts were studied for removal of metals from water. For successful sedimentation of the colloidal particles, further removal of heavy metals of wastewater utilising freshwater yeast strains and its mechanism for particle adsorbent can be examined.

Turbidity caused by clay particles is a big issue in Goa's mining districts when it comes to providing potable water during the rainy season. The flocculation and sedimentation of colloidal turbidity present in water polluted by mine rejects are highly dependent on yeast adhesion properties.

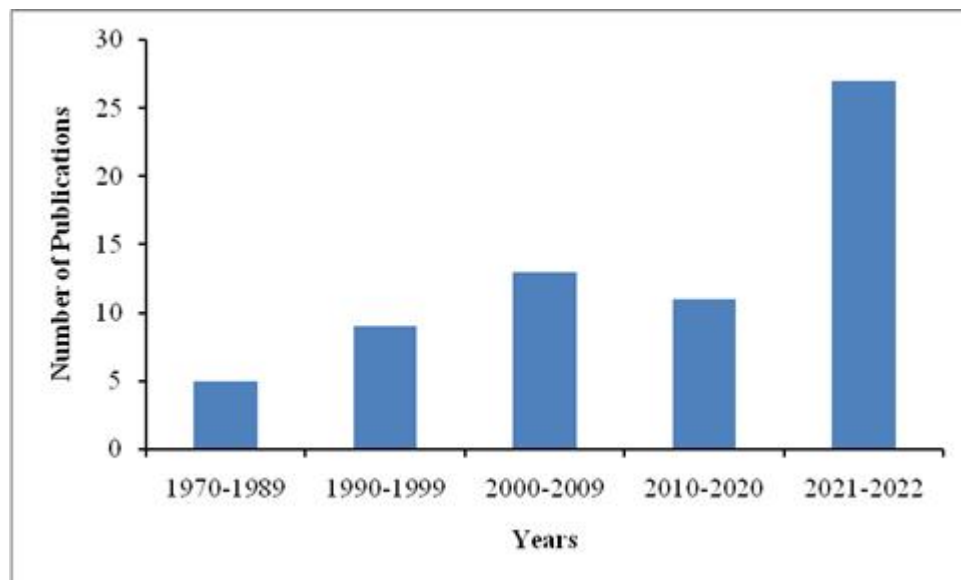
Since chemical coagulants are not ecofriendly; use of plants generates sludge and bacterial strains have difficulty in large scale production for effective biosorption or flocculation, a study can be performed to evaluate yeast strains for effective bioflocculation. Yeasts could prove more advantageous as it can be produced on large scale with less effort and avoiding the production of secondary pollutants.

### 1.11. Present methods reported for the removal of turbidity

Method	Limitations
Chemical coagulant eg. alum salt	Non-biodegradable, Not ecofriendly, Chemical coagulant produces colours in effluent of treated water and forms fragile flocs difficult to settle. (Bhatti, Z. A 2009)
Physical	Long time required, Dissolved molecules not removed by conventional physical treatments.
Biological eg. Moringa	Produces huge sludge (Azni et al., 2006)
Bacteria	Less efficient than fungi (Nazareth et al., 2001)

Biosedimentation is the process to achieve the destabilization of colloidal particle due to contact with biosedimentation to form large sediments that can be easily removed by from water. Particle are having opposite charges are attached to the surface of microbial biomass (Len et al., 2006). Clay minerals are having positive charge while microbial cell surface is having negative charge, clump together get settled down efficiently.

The literature, on the other hand, lacks a comprehensive assessment of the relative performance of yeast and water treatment systems for turbidity removal, as well as the consequences of integrating them in a continuous flow system.



**Figure.1.6: Number of publications for bioremediation of wastewater using yeasts**  
(Accessed from different search engines)

### **1.12. General mechanism of metal tolerance by resistant species**

Several studies suggested that bacteria developed metal tolerance after being exposed to harmful metals shortly after life began, maybe as a result of the abundance of metal on the planet (Girno et al., 2002). Metal resistance in bacteria is assumed to have evolved as a result of recent exposure to metal pollution over the last 50 years. The current environmental degradation caused by anthropogenic metal activities has heightened the need for research into microbial metal resistance and treatment (Silva et al., 2012). Metals in the environment are directly influenced by microorganisms. During adaptation, microorganisms have created innovative metal resistance and detoxifying methods. Plasmids can be used to codify the metal resistance of microbes (Galetti., 2019). It can be metal specific at times, or it can exhibit resistance to a variety of metals at other times. Metal resistance is caused by metals binding to extracellular materials, which immobilizes the metals. Cadmium, lead, zinc, and iron are a few of the cationic metals that can bind to anionic cell surfaces. Metals can bind strongly to carboxylic, amino, thio, hydroxo, and hydroxyl carboxylic phosphoryl groups on algal cell surfaces, while phospholipids in bacterial lipopolysaccharides in the outer membrane also interact strongly with cationic metals (Sathendra et al., 2018). The distribution of metals is influenced by cell surface binding, which is especially relevant in the aquatic environment. In practice, microorganisms are investigated for their ability to sorb metals for the aim of bioremediation, or the removal of metal pollutants from the environment.

#### **1.12.1. Exopolymer binding**

The most prevalent approach of lowering metal bioavailability is through this mechanism. Polysaccharides, carbohydrates, nucleic acids, and fatty acids make up the majority of exopolymeric compounds. Exopolysaccharides protect cells from desiccation, phagocytosis, and parasitism in addition to their role in heavy metal binding. The pH of the environment has a big impact on microbial exopolymers. Metal detoxification via EPSs causes metal immobilization and prevents metal entrance into the cell (Sathendra et al., 2018).

**1.12.2. Siderophore complexation** – Siderophores are low-molecular-weight organic compounds that chelate iron. Their biological role is to collect iron in low-concentration



environments and deliver it into the cell. Siderophores, on the other hand, may interact with metals that are chemically related to iron, such as aluminium, gallium, and chromium. Siderophores diminish metal bioavailability and thus metal toxicity by binding to metals. In cyanobacteria, siderophores, for example, minimise copper toxicity (Roane and Pepper, 2000).

**1.12.3. Biosurfactants complexation** – Several bacteria create this class of chemicals, which are expelled outside the cell. Biosurfactants have recently been studied for their role in the creation of complex metals such as cadmium, lead, and zinc (Miller, 1995). Metals become more soluble when they are complexed with biosurfactants, and the complexed metal is non-toxic to the cell. Metal reduction precipitation – Common metabolic by-products that result in metal reduction can affect metal bioavailability (Gupta et al., 1993). Soluble metals are converted to less soluble metal salts, such as sulfidic and phosphidic metal salts, in this situation. *Citrobacter spp.* can bind lead and copper in aerobic conditions due to phosphate synthesis by enzymes (Macaskie., et al 2000).

**1.12.4. Metal dependent mechanism of metal resistance** – The mechanisms of intracellular metal resistance in microbes are not well known. Metal binding and sequestration by metallothioneins or related proteins is perhaps the best-known process (Prasad et al., 2004).

When metal pollution is present in the environment, these ways are triggered in the bacterial system. Heavy metal metabolism and the management of various forms of stress are both aided by MTs. Plant phytochelations are comparable to metallothioneins in many aspects, including the high amount of cysteine residues in the protein and the fact that both are involved in heavy metal detoxification. Three traits are shared by all metallothioneins - They are proteins with a low molecular weight. They are high in cysteine residue, high in metal content, and exhibit metal ion coordination in metal thiolate clusters. The metallothioneins can be categorised into two groups. All animal metallothioneins are classified as Class I, while those found in plants and other microbes are classified as Class II (Wilfried et al. 2000) The mammalian archetype is represented by the Class-I MTs, which have been fully described (Wilfried et al., 2000).

The thiol for mercaptide bonding in metal-thiolate clusters is provided by the invariant alignment of cys inside the protein. The alpha-domain in the carboxy terminal region of Cd and Zn is a metal cys cluster.

The metal 3-cys-9 cluster is found in the amino terminal beta-domain. 2-4 amino acids connect the two domains. At least 62 metallothioneins have been identified in mammals, fish, crabs, oysters, and mussels, among other animals (Fowler et al. 1987). *Synechococcus spp.*, *E. coli*, and *Pseudomonas putida* have all been found to produce metallothionein-like proteins (Gupta et al., 1993).

The efflux system is another metal-dependent mechanism of metal resistance. To remove metals from the cell, several microbes use plasmid-encoded energy-dependent metal efflux mechanisms. ATPases are used in some efflux systems, while chemiosmotic ion/proton pumps are used in others. These processes use active transport (ATPase pump) or diffusion (chemiosmotic ion/proton pump) to aggressively return harmful ions that have entered the cell out of the cell. The three metals most typically related with efflux resistance are arsenate, chromium, and cadmium (Nies., 2003).

#### **1.12.5. Yeast in phosphate solubilization**

As Phosphate is an important nutrient for plant growth, its solubility is slow in soil. Plant can take up phosphate only when it is in dissolved form. Many researchers have studied the adsorption of P from wastewater (Neal., 2001; Loganathan, 2014).

Phosphorus is an important component of plant growth, but excessive usage of phosphate fertilizers results in runoff and leaching during rainstorms, as well as overloading and eutrophication of aquatic habitats. Than. et al., (1973) studied yeast for the wastewater treatment and concluded, that water samples collected from different treatment sites, showed huge reduction in Phosphorous, ammonia and nitrogen after treatment with yeast. Aquatic hypomyces *Tetracladium setigerum* studied for phosphate solubilization efficiency (Sati et al., 2018) found 3.15 mg/L phosphate soluble capability. Soil fungi were explored for phosphate solubilization by many researchers eg. *Trichoderma* studied for its phosphate solubilization efficiency and impact on growth rate of *Avicennia marina* (Rawat. et al., 2011). Sharma, (2011)

isolated soil fungi and studied it for phosphates solubilization efficiency in both solid and liquid medium and reported rock phosphate solubilization upto 61.6% gm.

**Table.1.12 Reported technique for aquatic yeast isolation**

Process	I	II	III		References
Name of sampler	Niskin	Van Dorn	Kemmerer	-	Jannasch et al., (1973)
Sampling	Deep sea	Surface	Near Shore	-	Reference/Remark (Jannasch et al., (1973)
Isolation	1. Water passing through membrane filter 2. Filter kept on media.	Water sample plating directly, or after dilution	1. Water passing through membrane filter 2. Filter resuspended in sterile water for low density yeast population. 3. Water plated on media.	-	-
Medium	YPD Jannasch et al. (1973)	PDA	2% MEA	Y M	<a href="http://deskuervis.nic.in/pdf/manual_for_yeast_work_sept_2008.pdf">http://deskuervis.nic.in/pdf/manual_for_yeast_work_sept_2008.pdf</a>
Strain maintenance	MEA	-	-	-	<a href="http://deskuervis.nic.in/pdf/manual_for_yeast_work_sept_2008.pdf">http://deskuervis.nic.in/pdf/manual_for_yeast_work_sept_2008.pdf</a>

### 1.13.1. Fractal characters

Yeast are known to create cell wall polysaccharides that are useful in a variety of cell-cell and cell-to-inorganic particle interactions. Cell surface geometry and morphology, in addition to chemical appropriateness, influence the biosedimentation process.

The study of mycelial complex structures is difficult. Cultures were exhibiting fractal growth on the surface of their colony. Fractal geometry has made significant contributions to understanding of microbial growth patterns.

Fractal geometry allows the irregularity and complexity of mycelial surface structures to be measured in relation to growth, metabolic activity, enzyme synthesis, and coloration (Obert et al., 1990). The fractal dimension of mycelium can be related to its form and productivity, such as citric acid fermentation and different antibiotic fermentations (Papagianni., 2004).

Fractal growth patterns can also be used to evaluate the availability of nutrients.

Low nutritional environments result in low hyphal mass, while rich nutrient environments result in high hyphal mass (Ritz et al., 2001). Branches begin to develop in diverse directions once microbial mycelia have begun to proliferate.

Mandelbrot's fractal theory presented a novel technique to defining the geometry of systems with no specified geometry. Puchkov (2016) has provided examples of using image analysis in the study of both the macroscopic and the microscopic microbiological objects obtained by various imaging techniques.

The fractal dimension is the most essential numerical parameter in calculating the fractal of any mass. The number of independent quantities required to specify the location or arrangement of points on an item is referred to as fractal dimension (Shirali 2014). For calculating the fractal dimension of any fractal mass, a variety of concepts have been offered.

Cluster to cluster collision results in the creation of aggregates with a low fractal dimension. Yan L. X (2005) devised a new method for calculating the fractal dimension of bio flocs in activated sludge suspensions. The fractal dimension was used to distinguish between the flocs and granule particles generated in an anaerobic digester used for water treatment (Bellouti. et al., 1997). Using fractal geometry theory, Saiedi (2017) investigated temporal variation in aggregate stabilization, carbon content and microbial respiration. Balaban (2018) investigated the evolution of spatial arrangement in *Enterobacter cloacae* aggregates using multifractal analysis to evaluate its radial growth on semi-solid substrates.

The type of microorganisms and the method employed to aggregate them affect the fractal behaviour of microbial aggregation (Logan., 1991). Because colony and density

heterogeneities, as well as vacuoles within mycelia, are difficult to quantify, averages are assessed using image-based approaches (Boddy, 1999). Mycelia are also fractal in nature, like other irregular structures. Fractal geometry and fractal dimension may be a useful tool for determining the mycelial development pattern or extent in space because it involves their space filling capacity. According to Barry (2009), filamentous bacteria fermentation process is influenced by their size and morphological behavior.

Because traditional methods such as projected area, length, and perimeter are limited, fractal dimensions can be used to assess morphological development productivity, which is connected with morphological adaptability and can be determined by fractal dimensions (Barry 2009). Computer models have been created to replicate fungal colonies, which have been researched for the nature of hyphal branching points, growth, and next-door neighbour characteristics, as well as growth factors in relation to fractal dimension (Obert.,1993). The diameter and division of single yeast cells were measured using fractal dimensions (Tomankova., 2006; Vesela., 2001).

Worix (2009) used the fractal dimension of bacterial colonies to compare biochemical and physiochemical changes in bacterial colonies as well as how they responded to environmental changes.

*Achlya bisexualis* morphology studied, the fractal dimension of the response to heavy metal concentration decreases as concentration rises (Lundy 2001). Gonzalez (2016) used polymerosomes to mimic biofilm formation and the aggregation process was investigated using a fractal framework, as well as antibacterial properties. Fractal dimensions can be a useful tool for analysing aggregates since they allow for quantitative evaluation of aggregate shape (Logan., 1991). However, due to their fragile nature, flocs cannot be investigated. Particle aggregates formed during the wastewater treatment process have fractal characteristics (Ganczarzy., 1989).

The fractal dimension tool was used to characterise *Trichiurus haumela* in a frozen state. During storage, it was discovered that when the temperature dropped, the quality (hardness, springiness, and fractal dimension) of the product diminished (Luan et al., 2018).

The whey protein gel structure and visco elasticity have been described using the fractal dimension idea. The fractal dimension has been used to investigate soil characteristics. Different forms of aggregates form as a result of soil particle association, mineral and organic materials, and soil structure. Using fractal dimensions, the water retention of soil particles was compared. The soil bulk density and clay silt content were found to be connected to the fractal dimensions of the soil water retention curve (Pazira et al., 2018). Lestrai (2018) and Malekanis (1996) evaluated fractal analysis techniques for determining clay material fractal dimensions.

The fractal dimension of flocs changes as the method of floc creation changes. The geometric characteristics of flocs can be understood using the fractal analysis theory (Li., 2006).

The lowest fractal dimensions were detected in biological flocs and the flocculation process due to bridging the aggregates, while larger fractal dimensions were observed in aggregates created by sweep substrate transfer. The fractal dimensions of flocs are also determined by hydraulic conditions (Sun, 2013).

Furthermore, after churning, the fractal dimension initially grows and subsequently declines. There are several methods for calculating fractal dimensions, but all of them are based on the power law connection (Kramer., 2004). The irregularity and complexity of mycelia surface structures are related to growth, metabolic activity, enzyme synthesis, and coloration, according to fractal geometry (Martin et al., 1990).

Hypothesis was made to determine whether the fractal dimensions of yeast colony edges could serve and assist in screening and selecting strains capable of bioclarification of turbidity.

### **1.13.2. Fractality Index**

Fractality index is a four digit number obtained by multiplying the score produced by the fractal analysis software JFRAD by 1000 with the last number being rounded up. (Kamat., 2019).

#### **1.14. Drinking water status in mining areas of Goa**

Goa is blessed with a plethora of freshwater bodies. In Goa, various surface water reservoirs are used as a source of drinking water. The physiochemical characteristics of water are a critical aspect in determining the water quality for various applications. Analyzing the physiochemical properties of water aids in the understanding of water quality and the development of techniques to address problems. Water quality is well understood to be as vital as its quantity. As a result, an examination of the physiochemical properties of water bodies was required.

Manderker et al. (2012) assessed ground and surface water in Goa's mining sites and found that all water parameters were within acceptable limits.

Moreover, they also discovered that the pH of ground water in mining regions was somewhat acidic. Further, Krishan et al. (2016) investigated the quality of Goa's groundwater and created a water quality index tool that combines data from various water quality criteria into a single number. According to them, these indices are temporal and spatial based and may vary depending on local factors. Their research indicated that the water quality was good and safe to drink. The quality of Goa's coastal water was also assessed (Sampath et al., 2018).

(Sawaiker. et al., 2016) investigated the lake's physiochemical characteristics as well as its phytoplankton diversity, concluding that seasonal fluctuations in the lakes' physiochemical characteristics were detected. The physiochemical parameters of tap and filter water were studied by Singh et al., (2015). While tap water had a pH that was acceptable, filter water had a pH that was less than ideal. Both tap and filter water had alkalinity and total hardness within acceptable limits. Sawaiker et al., (2017) assessed water quality using microbes as a bio monitor and found that diatoms are excellent indicators of water contamination. Sreevnivasa. et al., (2013) investigated the chemical properties of water samples obtained from a bore well and an open well in Bicholim, Goa. The majority of the parameters in the water sample were found to be within acceptable limits. Furthermore, bicarbonate levels were found to be over 1000 ppm in several areas, such as Shikheri and Gaonkarwada. The water quality of four beaches was studied by Saxena et al., (2018). Because the urban discharge point was closer to this location, beaches had lower water quality and a higher quantity of coliforms.



**Figure.1.7: Water tank at Morlem**

### **1.15. Bioflocculation test need at laboratory scale**

The purposes of these laboratory-scale biosedimentation tests are to assess the viability of wastewater treatment using SMTTW and to look into the characteristics of the process, including the microscopic mechanism and parameters influencing the process. Cultures are first screened as bioflocculants. Second, the goal is to identify any differences in treatment intensity across the different characteristics of the investigated cultures, such as biomass dose, age, and pH of wastewater. Procedures can be optimised by testing at various scales.

#### **1.15.1 Rapid mixing**

The quick mixing stage is perhaps the most critical point of the coagulation-flocculation/biosedimentation process because it is here that destabilizing reactions take place and where primary floc particles are generated, the features of which have a significant impact on later flocculation kinetics.

### **1.16. Molecular Study of yeast strains**

Several yeast species from various environments have yet to be identified and used for commercial purposes. The majority of therapeutically or medically important yeasts are identified using molecular sequencing. Most species found in the aquatic environment are still unidentified (some species are included in the table 1.13) in the soil, and in the



air. After yeast has been identified, it can be classified and graded in comparison to other species. Some yeast species have antifungal properties, meaning they are resistant to fungus, whereas others have particular colours or can be fatal. As a result, adequate management is essential to keep the yeast culture in line with their nonviolent property. Butler et al. (2009) sequenced the genome of *C. tropicalis*, a diploid yeast with 12 chromosomes per cell, a genomic size of 14.5 Mb, 6,258 genes producing proteins, and a guanine-cytosine concentration of 33.1 percent, according to Doi et al. (1992). The number of chromosomes is not precisely known. *C. tropicalis* was formerly thought to be an asexual yeast. Some research has suggested that tetraploid cells can be produced by mating between diploid cells and (Porman et al., 2011; Xie et al., 2012; Seervai et al., 2013). The cause of cells turning from white to opaque, according to the researchers, could be mating, which is controlled by colony phenotypic flipping. According to Seervai et al. (2013), *C. tropicalis* strains can be made to go through a parasexual cycle without meiotic reduction from a tetraploid state polyploidy, which affects gene expression and protein production in the cells (Morrow, 2013). *C. tropicalis* showed reduction in ploidy and considered a mechanism of adaptation and may be associated with cell stress (Bermanet et al., 2012). *C. tropicalis* has showed huge genetic similarity with *C. albicans* to other *Candida* species. (Butler et al., 2009). This minute different can be noticed with the help of molecular identification.

**Table.1.13 Aquatic yeast reported from different regions**

Name of Aquatic yeast	Source	Place	Reference
<i>Aureobasidium pullulans</i>	Sea water Lake River	Bohai Sea, Nahuel Huapi Lake Argentina Brazil	Zhou et al., 1991 Carlos et al., 2010 Adriana et al., 2012
<i>Candida azyma</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Candida curvata</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Candida famata</i>	Sea water	Bohai Sea	Harrison et al., 2017
<i>Candida guilliermondii</i>	Sea water	Bohai Sea	Guerra et al., 1938
<i>Candida humicola</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Candida insectorum</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Candida krusei</i>	Sea water	Bohai Sea	Zhou et al., 1991

<i>Candida multis-gemmis</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Candida parapsilosis</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Candida pelliculosa Redaelli</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Candida sake</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Candida terebra</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Candida tropicalis</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Candida valdiviana</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Cryptococcus albidus</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Cryptococcus ater C</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Cryptococcus hungaricus</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Cryptococcus infirmominiatus</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Cryptococcus laurentii</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Cryptococcus luteolus</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Cryptococcus magnus</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Cryptococcus uniguttulatus</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Debaryomyces hansenii</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Debaryomyces polymorphus</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Pichia burtonii</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Pichia carsonii</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Pichia etchellsii</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Pichia guilliermondii</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Pichia heimii</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Pichia ohmeri</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Pichia philogaea</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Pichia scolyti</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Rhodotorula aurantiaca</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Rhodotorula glutinis</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Rhodotorula graminis</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Rhodotorula minuta</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Rhodotorula rubra</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Saccharomyces cerevisiae</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Saccharomyces kluyver</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Torulaspora delbrueckii</i>	Sea water	Bohai Sea	Zhou et al., 1991
<i>Candida austromarina</i>	Sea water	-	Yarrow et al., 1978
<i>Candida maritima</i>	Seawater	Australia.	Zhou et al., 1991
<i>Leucosporidium antarcticum</i>	Antarctic sea water	Antarctic	Fell et al., 1970

<i>Metschnikowia bicuspidata</i>	Sea Water	USA	Zhou et al., 1991
<i>Metschnikowia krissii</i> Uden	Seawater,	USA.	Zhou., et al 1991
<i>Metschnikowia zobellii</i>	Seawater,	USA.	Zhou et al., 1991
<i>Rhodospiridium dacryoideum</i>	Seawater	Antarctic	Fell et al.,1991
<i>Metschowinkia australis</i>	Sea Water	Antarctic	Zhou., et al 1991
<i>Rhodospiridium sphaerocarpum</i>	Seawater	Antarctic	Zhou., et al 1991
<i>Candida Mycoderma</i>	Atlantice ocean	-	Zhou., et al 1991
<i>C. membranaefaciens</i>	Arbian Sea	India	Gupta., 1990
<i>Ambrosiozyma platypodis</i>	River	South Santiam, USA	Zhou et al., 1991
<i>Candida methanosorbosa</i>	River mud	Japan.	Zhou et al., 1991
<i>Kurtzmanomyces tardus</i> Gimenez-Jurado & van Uden	Contaminated demineralized water	Portugal	Zhou et al., 1991
<i>Lindnera sargentensis</i>	Fresh Water	USA	Zhou et al., 1991
<i>Metschnikowia bicuspidata</i> var. <i>chathamia</i>	Fresh water of lake,	New Zealand	Zhou et al., 1991
<i>Sporopachydermia lactativora</i>	Seawater	Antarctic	Zhou et al., 1991
<i>Sterigmatomyces halophilus</i>	Seawater.	-	Zhou et al.,1991
<i>Ogataea minuta</i>	Water	USA	Zhou et al., 1991
<i>Ogataea minuta</i> var. <i>nonfermentans</i>	Water	USA	Zhou et al.,1991
<i>Pseudozyma antarctica</i>	Sediment of Lake Vanda	Antarctica	Zhou et al., 1991
<i>Sympodiomyces parvus</i>	Seawater	Antarctic	Zhou et al.,1991
<i>Debaryomyces</i>	Central Pacific Ocean	-	Zhou et al., 1991
<i>Candida torulopsis</i>	Pacific Ocean	-	Zhou et al., 1991
<i>Torula</i>	Atlantice Ocean	-	Zhou et al.,1991

<i>Debaryomyces fabryi</i>	Sea	Arabian, Bay of Bengal	Kuriokose., et al 2012
<i>Debaryomyces nepalensis</i>	Sea	Arabian, Bay of Bengal	Kuriokose ., et al 2012
<i>Debaryomyces subglobosus</i>	Sea	Arabian, Bay of Bengal	Kuriokose., et al 2012
<i>Rhodotorula ferulica</i>	Polluted river water,	Portugal	Zhou et al., 1991
<i>Trichosporon coremiiforme</i>	Water bug,	Thailand	Zhou et al., 1991
<i>Candida fukuyamanensis</i>	Pond	Japan	Zhou et al., 1991
<i>Rhodotorula mucilaginosa</i>	River	-	The yeasts : A Taxonomic Study. (1952)
<i>Candida krusei</i>	River,	Brazil	Adriana et al., 2012
<i>Candida parapsilosis</i>	River, lake	Brazil USA	Adriana et al., 2012 Den et al., 1963
<i>Cryptococcus laurentii</i>	River	Brazil	Silva et al., 2012
<i>Cryptococcus luteolus</i>	River	Brazil	Silva et al., 2012
<i>Cryptococcus hungaricus</i>	River	Brazil	Silva et al., 2012
<i>Cryptococcus albidus</i>	River	Brazil	Silva et al., 2012
<i>Kloeckera japonica</i>	River	Brazil	Adriana et al., 2012
<i>Kodmaeae</i>	River	Brazil	Adriana et al., 2012
<i>Metschnikowia</i>	River	Brazil	Adriana et al., 2012
<i>Pichia</i>	River	Brazil	Adriana et al., 2012
<i>Rhodotorula</i>	River	Brazil	Adriana et al., 2012
<i>Issatchenkia sp.</i>	lake	Brazil	Silva et al., 2012
<i>Candida diversa</i>	lake	Brazil	Silva et al., 2012
<i>Torulaspota pretoriensis</i>	lake	Brazil	Silva et al., 2012
<i>Candida pseudolambica</i>	lake	Brazil	Silva et al., 2012
<i>Cryptococcus podzolicus</i>	lake	Brazil	Silva et al., 2012
<i>Hanseniaspora uvarum</i>	lake	Brazil	Silva et al., 2012
<i>Trichosporon jirovecii</i>	lake	Brazil	Silva et al., 2012
<i>Williopsis saturnus</i>	lake	Brazil	Silva et al., 2012
<i>Hanseniaspora thailandica</i>	lake	Brazil	Silva et al., 2012
<i>Trichosporon laibachii</i>	lake	Brazil	Silva et al., 2012
<i>Bullera sp.</i>	lake	Brazil	Silva et al., 2012
<i>Cryptococcus rajasthanensis sp.</i>	lake	Brazil	Silva et al., 2012

<i>Cryptococcus podzolicus</i>	lake	Colombian, Brazil	Silva et al., 2012
<i>Candida lipolytica</i>	Guanabara Bay	Brazil	Hagler et al., 1979
<i>Sporobolomyces</i>	Lakes	Argentina	Libkind et al., 2004
<i>Cryptococcus agrionensis</i>	River Agario	Argentina	Libkind et al., 2014
<i>Rhodotorula mucilaginosa</i>	River Agario, Lake	Argentina	Libkind et al., 2014 Carlos et al., 2014
<i>Holtermanniella festucosa</i>	River Agario	Argentina	Libkind et al., 2014
<i>Cystofilobasidium capitatum</i>	River Agario	Argentina	Libkind et al., 2014
<i>Cystofilobasidium. macerans</i>	River Agario	Argentina	Libkind et al., 2014
<i>Cryptococcus albidus</i>	River Agario	Argentina	Libkind et al., 2014
<i>Cryptococcus antarcticus</i>	River Agario	Argentina	Libkind et al., 2014
<i>Cryptococcus cylindricus</i>	River Agario	Argentina	Libkind., et al 2014
<i>Cryptococcus laurentii</i>	River Agario, River Suwanee	Argentina, Florida	Libkind et al., 2014 Lazarus et., al 1974
<i>Cryptococcus victoriae</i>	River Agario	Argentina	Libkind et al., 2014
<i>Rhodotorula colostri</i>	River Agario	Argentina	Libkind et al., 2014
<i>Rhodosporidium babjevae</i>	River Agario	Argentina	Libkind et al., 2014
<i>Rhodosporidium toruloides</i>	River Agario	Argentina	Libkind et al., 2014
<i>Holtermanniella festucosa</i>	River Agario	Argentina	Libkind et al., 2014
<i>Cystofilobasidium capitatum</i>	River Agario	Argentina	Libkind et al., 2014
<i>Cystofilobasidium macerans</i>	River Agario	Argentina	Libkind et al., 2014
<i>Candida austromarina</i>	River Agario	Argentina	Libkind et al., 2014
<i>Bauerago sp</i>	River Agario	Argentina	Libkind et al., 2014
<i>Sporobolomyces roseus</i>	River Agario	Argentina	Libkind et al., 2014
<i>Cryptococcus agrionensis</i>	River Agario	Argentina	Russo et al., 2009
<i>Delphinella strobiligena</i>	Lake	Nahuel Huapi lake Argentina	Carlos et al., 2010
<i>Dioszegia hungarica</i>	Lake	Nahuel Huapi lake Argentina	Carlos et al., 2010
<i>Candida.sp.</i>	Lake	Nahuel Huapi lake Argentina	Carlos et al., 2010
<i>Cystofilobasidium capitatum</i>	Lake	Nahuel Huapi lake Argentina	Carlos et al., 2010

<i>Cystofilobasidium. Infirmominiatum</i>	Lake	Nahuel Huapi lake Argentina	Carlos et al., 2010Libkind., et al 2014
<i>R. colostri</i>	Lake	Nahuel Huapi lake Argentina	Carlos et al., 2010
<i>Rhodospiridium diobovatum</i>	Lake	Nahuel Huapi lake Argentina	Carlos., et al 2010
<i>Bullera dendrophila</i>	Lake	Nahuel Huapi lake Argentina	Carlos et al., 2010
<i>Cryptococcus adeliensis</i>	Lake	Nahuel Huapi lake Argentina	Carlos et al., 2010
<i>C. diffluens</i>	Lake	Nahuel Huapi ake Argentina	Carlos et al., 2010
<i>C. festucosus</i>	Lake	Nahuel Huapi ake Argentina	Carlos et al., 2010
<i>C. heveanensis</i>	Lake	Nahuel Huapi lake Argentina	Carlos et al., 2010
<i>Cryptococcu magnus</i>	Lake	Nahuel Huapi lake Argentina	Carlos et al., 2010
<i>C. saitoi</i>	Lake	Nahuel Huapi lake Argentina	Carlos et al., 2010
<i>C. stepposus</i>	Lake	Nahuel Huapi ake Argentina	Carlos et al., 2010
<i>C. wieringae</i>	Lake	Nahuel Huapi lake Argentina	Carlos et al., 2010
<i>Guehomyces pullulans</i>	Lake	Nahuel Huapi lake Argentina	Carlos et al., 2010
<i>Candida parapsilosis</i>	Lake	Nahuel Huapi lake Argentina	Carlos et al., 2010
<i>C. railenensis</i>	Lake	Nahuel Huapi lake Argentina	Carlos et al., 2010
<i>C. sake</i>	Lake	Nahuel Huapi lake Argentina	Carlos et al., 2010
<i>C. carnescens</i>	Lake	Nahuel Huapi lake Argentina	Carlos et al., 2010
<i>Cryptococcus tephrensii</i>	Lake	Nahuel Huapi lake Argentina	Carlos et al., 2010 2010
<i>Debaryomyces hansenii</i>	Lake	Nahuel Huapi lake Argentina	Carlos et al., 2010

<i>Hanseniaspora uvarum</i>	Lake	Nahuel Huapi lake Argentina	Carlos et al., 2010
<i>Pichia fermentans</i>	Lake	Nahuel Huapi lake Argentina	Carlos et al., 2010
<i>Candida famata</i>			Harrison et al., 1961
<i>C. glabrata</i>	Coastal water	Taiwan	Sheng ., 2012
<i>Saccharomyces yakushimaensis</i>	Coastal water	Taiwan	Sheng ., 2012
<i>Kazachstania jainicus</i>	Coastal water	Taiwan	Sheng ., 2012
<i>Kodamaea ohmeri</i>	Coastal water	Taiwan	Sheng ., 2012
<i>Pichia anomala</i>	Coastal water	Taiwan	Sheng., 2012
<i>Issatchenkia orientalis</i>	Coastal water	Taiwan	Sheng., 2012
<i>Hanseniaspora uvarum</i>	Coastal water	Taiwan	Sheng 2012
<i>Rhodotorula pacifica</i>	Deep sea	Pacific ocean	Nagahama et al., 2006
<i>Kluyveromyces nonfermentans</i>	Deep sea	Pacific ocean	Nagahama et al., 1999
<i>Cryptococcus surugaensis</i>	Suruga Bay	Japan	Nagahama et al .,2003
<i>Cryptococcus</i>	Sea	Antarctic	Vishniac., et al 1979
<i>Candida pulcherrima</i>	lake	USA	Den. et al., 1963
<i>Cryptococcus albidus</i>	lake	USA	Den. et al., 1963
<i>Cr. diffluens</i>	lake	USA	Den. et al., 1963
<i>Cr. gastricus</i>	lake	USA	Den. et al.,1963
<i>Cr. laurentii</i>	lake	USA	Den. et al 1963
<i>Rhodotorula glutinis</i>	lake	USA	Den. et al 1963
<i>R. pilimanae</i>	lake	USA	Den. et al 1963
<i>R. rubra</i>	lake	USA	Den. et al 1963
<i>Trichosporon cutaneum</i>	lake	USA	Den. et al 1963
<i>Candida sp.</i>	Marine	Nicobar Islands	Rao et al., 2011

<i>Candida albicans, Candida tropicalis, Debaryomyces hansenii, Geotrichum sp., Pichia capsulata, Pichia fermentans, Pichia salicaria, Rhodotorula minuta, Cryptococcus dimenna and Yarrowia lipolylica</i>	Mangrove sediment	South east coast, India	Saravanakumar et al., 2013
<i>Candida spp, Pichia spp., Cyberlinera spp. , Meyerozyma spp.,Clavispora spp.,Saccharomyces spp.,Kluyveromyces spp.,Yamadazyma spp. ,Trichosporon spp. and Wickerhamomyces spp.</i>	River	South Africa	Monapathi et al., 2020
<i>Torula , Pichia Wickeramomyces, Candida</i>	River	Japan	Naito et al., 2019
<i>Yamadazym, Candida atmosphaerica, C. spencermartinsiae, C. atlantica, C. oceani and C. taylorii</i>	Mid-Atlantic Ridge ocean water	Atlantic	Burgaud et al., 2016
<i>Rhodotorula mucilaginosa, Clavispora lusitaniae, Wickerhamomyces anomalus</i>	Ghellaze lake	Algeria	Aibeche et al., 2021

Considering the gap in knowledge regarding the use of freshwater yeast for the removal of colloidal turbidity from waste water, the present work aimed to fill the gaps by targeting freshwater yeasts as bioflocuants for wastewater treatment. The lead of work was obtained by Kamat et al., (2010) during the dissertation work of Post graduate students.



The main aim of this work was to test this hypothesis **“Whether freshwater yeast cultures could effectively reduce turbidity in freshwater, polluted due to entry of mine tailing water.”**

The specific objectives of the work were formulated:

#### **OBJECTIVES OF THE PRESENT STUDY**

1. Survey of Freshwater habitats from mining and non mining areas of Goa and isolation of natural aquatic yeasts culture.
2. Systematic screening, detection and evaluation of promising yeast strains capable of bioflocculation of colloidal turbidity.
3. Assessment of drinking water quality of mining areas of Goa.
4. Lab scale bioflocculation using natural yeast isolates upto a scale of 10-100 L.
5. Standardization of bioflocculation process for field level application scale 500 L-1000 L
6. Molecular study of only the selected strain.

**CHAPTER II**  
**MATERIALS AND METHODS**

## **2.1. Literature surveyed**

Fresh water yeast and wastewater treatment using yeast, related scientific literature was surveyed using a variety of online querying techniques, with the search terms bioclarification, biosedimentation, and turbidity reduction used in various databases, including general search engines, academic / institutional databases, specific journal search, patent websites, culture collections, biological, and chemical databases. Google. (<https://scholar.google.co.in>) Scholar's growth rate of online sources relevant to particular search phrases till August 2022.

### **2.1.1. Freshwater habitats surveyed**

Water habitats were selected based on their location, availability and water use or source. The main source of freshwater reservoirs was rainwater. Lake, temple tank, water fall's stream were among them. Water tanks were built in rectangular and in square shapes. Some of them were transparent and some of them were greenish in appearance (Fig.2.3, karapur Pond).

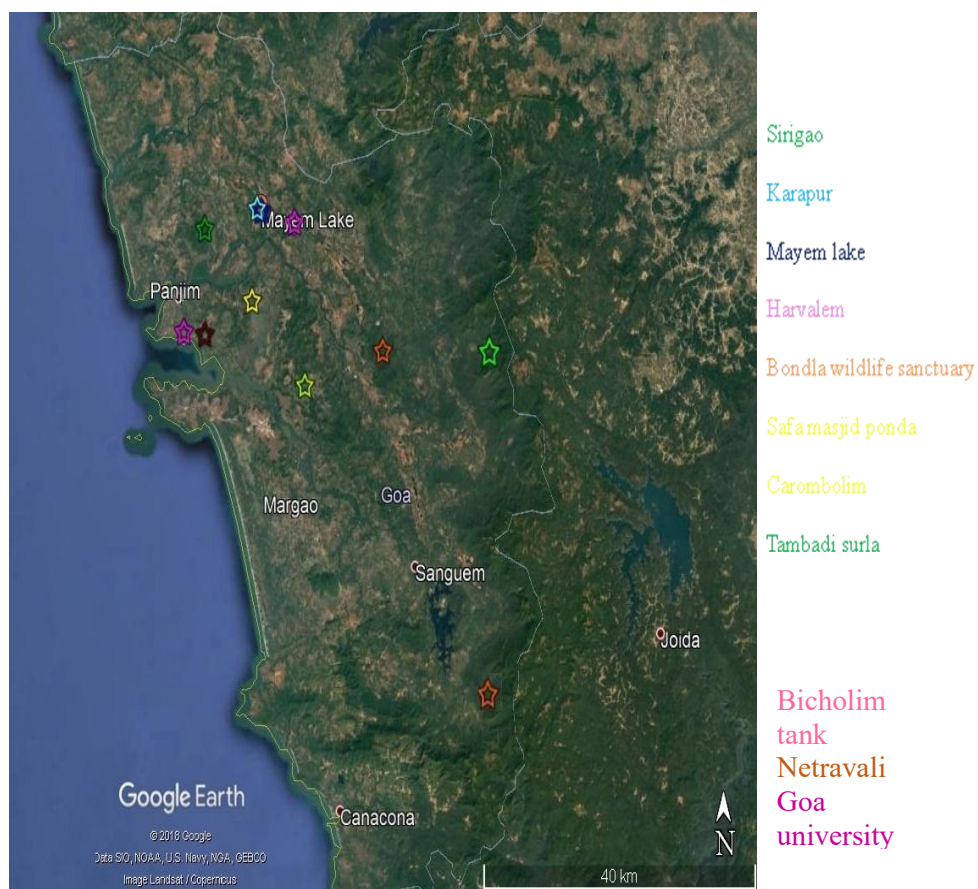
The seasonal pond in the Goa university campus showed macrophyte rich environment, at their surroundings. The pond near SafaMaszid had large number of fish population.

The Arvalem water fall's stream was turbid as human activities were going on, while the waterfall at chorla ghat and Netravali were clean and very less turbid. Mayem and Carombolim lakes were chosen as sampling sites as they are well known freshwater bodies in Goa's mining area. Macrophytes were absent near the Mayem lake Carombolim showed sufficient population of macrophytes. Both lakes had approximately identical turbidity

### **2.1.2. Water Sampling**

Freshwater bodies in Goa, including mining and non-mining areas, were chosen for water sample collection for the freshwater yeast isolation. As previously stated, the study locations were included temple tank, stream of water fall, and lake. Water samples were collected approximately 20 cm below surface in 1-L pre-sterile plastic bottles (Himedia). At the sampling sites, water parameters such as temperature (Multi Thermometer), electrical conductivity (Aquaro digital water conductivity meter), and total soluble solid (HM Digital TDS-3 meter) were measured. Freshwater samples were

taken in fifteen distinct locations from six talukas of Goa for the period of 2016 to 2018 monsoon season. Samples of water were brought to the laboratory, analysed and processed within 2-24 hrs of the collection of water samples in order to obtain yeast strains. Moreover, no preservative were used for the storage of the water samples.



**Figure. 2.1: Google locations of water sampling sites**



**Figure.2.2: Water sampling locations (Created with SimpleMappr,  
<https://www.simplemappr.net>)**

**Table.2.1 Water sampling sites**

<b>Water Body</b>	<b>Location (Talluka)</b>	<b>Latitude</b>	<b>Longitude</b>
Bicholim tank	Bicholim	15 °36 '6.7500"N	73 °57 16. 66 96"E
Bondla stream	Ponda	15°23'56.87"N	74° 0'44.59"E
Carambolim lake	Tiswadi	15.4888° N	73.9279° E
Chorla water fall's stream	Sattari	15°35'45.18"N	74° 3'33.79"E
Harvalem water fall stream	Sanquelim	15°33'29.24"N	74° 0'44.84"E
Karapur tank	Bicholim	15°33'23"N	73°58'35"E
Mayem lake	Bicholim	15.5760° N	73.9400° E
Netravali water fall	Sanguem	15.0955° N	74.2178° E
Netravali bubbling lake	Sanguem	15.0942°N	74.20919°E
Safamasjid pond	Ponda	15.4066° N	74.0000° E
Sirigao tank	Bicholim	15.6099° N	73.9036° E
Tambdi surla stream	Sanguem	15°12'38.96"N	74° 9'1.51"E
Goa University campus (pond 1,2,3,	Tiswadi	15.4588° N	73.8342° E



**Figure.2.3: Water bodies(a) Safamasjid Pond (b) Sirigao Pond**

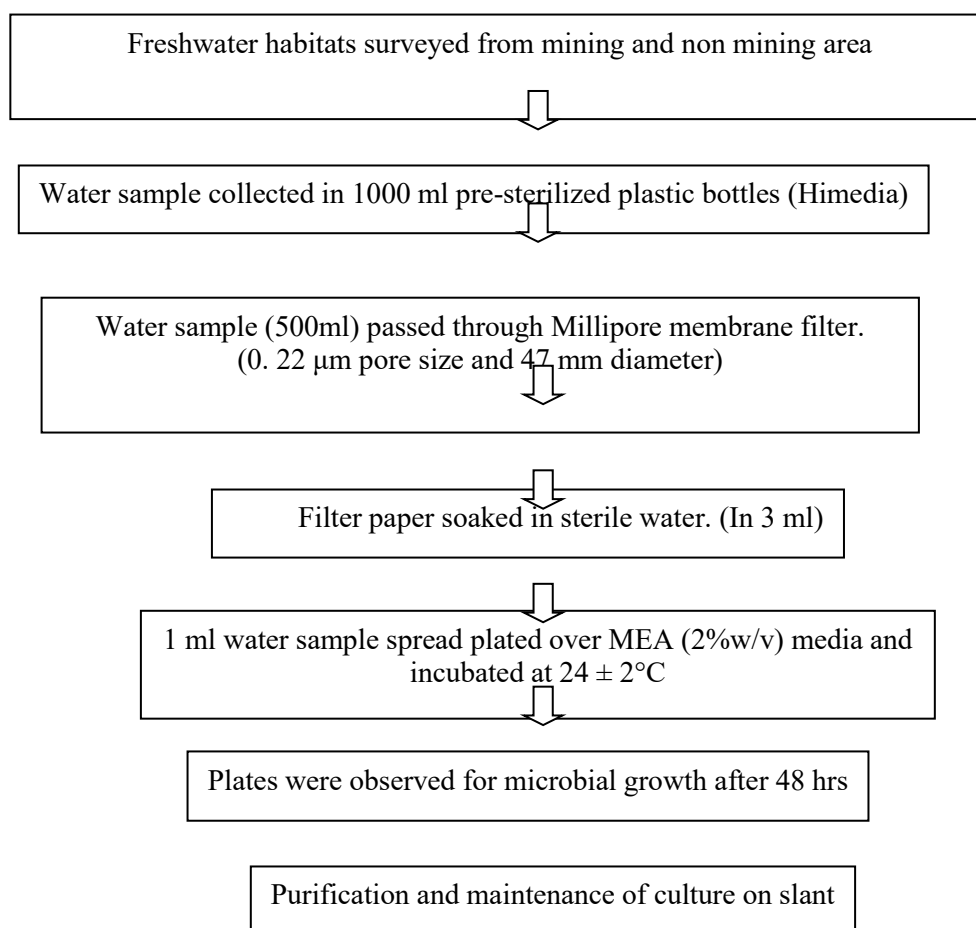


**Figure.2.4: (a) Bicholim temple tank, (b) Karapur tank, (c, d) Goa University seasonal pond (d) Bubbling Lake, (e) Netravali water fall**



## 2.2. Isolation of yeast cultures using membrane filter technique

Water samples were collected in 1000 ml pre-sterilized plastic bottles. 500ml water sample (in triplicates) was passed through a sterilized membrane filter (0.22  $\mu\text{m}$  pore size and 47 mm diameter, Millipore) in a sterilized membrane filter assembly, set up in the laminar air flow. After filtration the Membrane filter was removed and eluted in sterile 3ml water to remove the trapped material. 1 ml water sample was spread plated over MEA (2% w/v) media (incorporated with antibiotics 0.1mg/ml) and incubated at  $24 \pm 2^\circ\text{C}$ . (Kamat et al., 2010). Plates were observed for microbial growth, and morphological characteristics studied under the microscope, dissimilar colonies were picked from plates. Cultures were purified and maintained on malt extract agar slants.



**Figure.2.5: Scheme for freshwater yeast isolation**

### **2.2.1. Rainwater sample collection**

This work was done to check whether any rainwater yeast could enter the surface water body.

### **2.2.2. Assembly of sterile, dust- and contamination-free PVC containers**

Pre-sterile plastic bottles were used to collect the rain water. Six times in every monsoon from 2016 to 2018.

### **2.2.3. Selection of clear vegetation-free open spaces**

Water was collected at Goa University campus at the same place during every sampling. The area was vegetation free and open in order to avoid contamination.

### **2.2.4. Isolation of yeast, purification and maintenance**

Rain water samples were spread on MEA plates (containing 0.1mg/ml antibiotics) which were incubated for 48 hrs at 25°C. The plates were observed under a microscope for the growth of yeast and colony morphology.

### **2.2.5. Control plates**

Plates with nutrient agar and MEA (2%, with 0.1mg/ml antibiotics) were kept in the open air for one hour at the rain water collection site at the Goa university campus. Other plates were spread plated with water, which was used for rinsing the rainwater sampling apparatus. One plate with MEA (2%) was kept without spread plating rain water.

## **2.3. Taxonomic identification of the cultures tentatively**

Standard procedure for identification of yeast were employed and the yeast cultures were identified on the basis of their cultural and morphological characteristics.

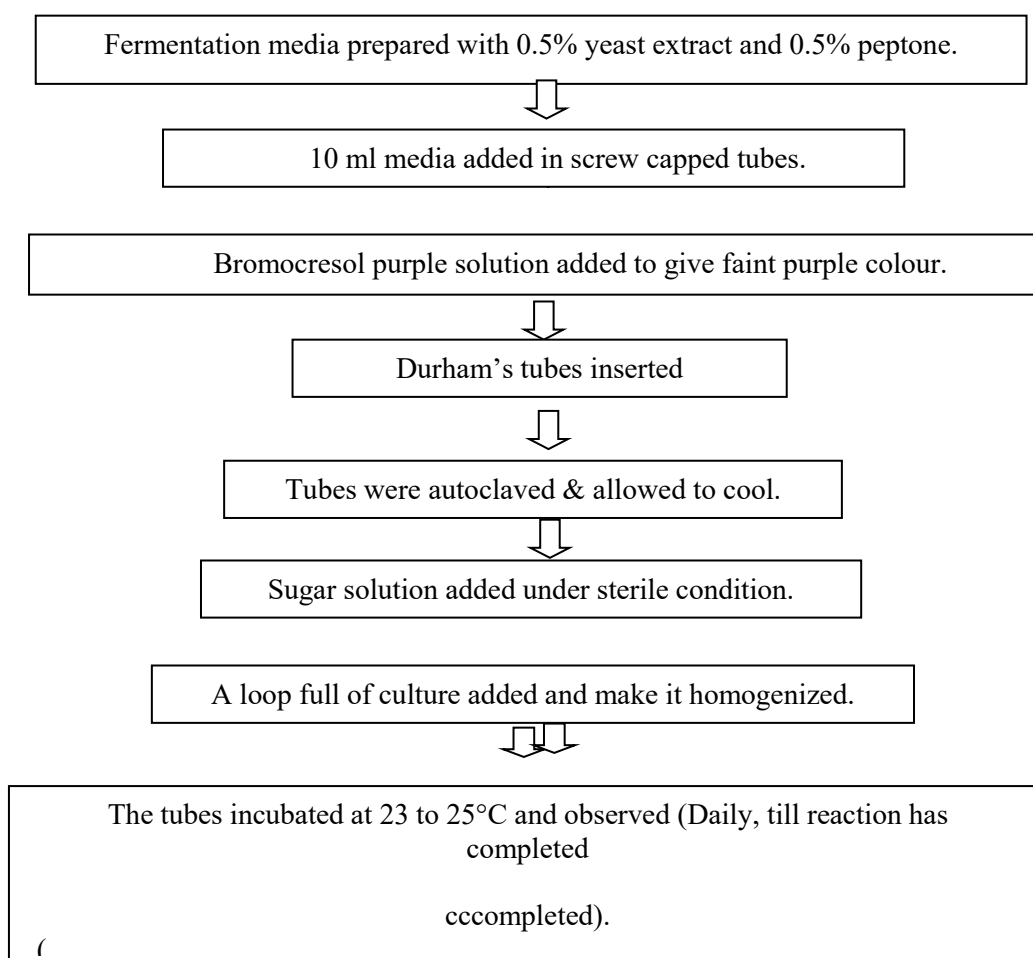
### **2.3.1. Morphological study of the cultures**

Strains were studied for its morphology under stereo microscope (Olympus SZ51, Tokyo, Japan). Micromorphology of cultures was studied under light microscope (Nikon Eclipse E200) by monochrome gram staining and photographed with NIS element microscope imaging software and confirmed the presence of yeast.

### 2.3.2. Biochemical tests

The yeast strains were identified on the basis of sugar fermentation, assimilation, ascospore production, formation of true mycelium test (Yarrow., 1998; Bhargva et al., 2015). Cultures tentatively identified upto the Genus level. (All chemical used, were of HiMedia Chemicals Ltd., India)

#### 2.3.2.1. Standard protocol for carbohydrate fermentation test(Dunford et al., 1999)



**Figure.2.6: Scheme of the of sugar test**

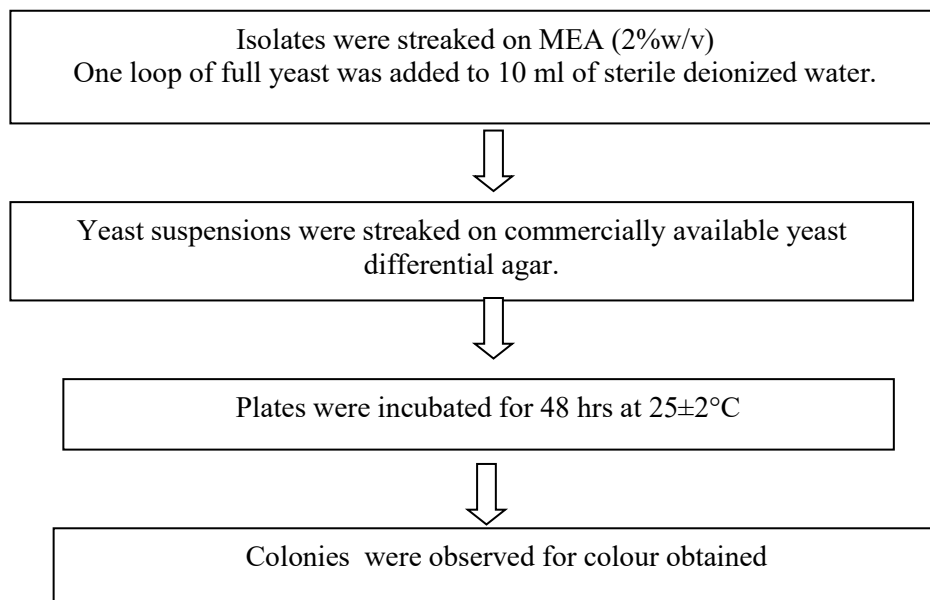
#### 2.3.2.2. Observed assay

- (a) Acid production as change in color of medium for purple to yellow.
- (b) Gas detection as bubbles of gas in durham tube.

### 2.3.2.3. Study of cultures on Candida differential agar

Candida Differential Agar is a selective and differential medium that enables for quick isolation of yeasts from mixed cultures as well as distinguishing Candida species such as *C.albicans*, *C.krusei*, *C.tropicalis*, and *C.glabrata* based on colony form and coloration. Results are obtained in 48 hours, making it ideal for the quick and presumptive identification of common yeasts in the mycology and clinical microbiology laboratories.

Nitrogenous, carbonaceous chemicals, and other critical growth nutrients are provided by peptone special and yeast extract. Phosphate acts as a medium buffer. Chloramphenicol lowers the bacterial flora in its immediate environment. *C.albicans* appears as light green smooth colonies, whereas *C.tropicalis* appear as blue to metallic blue elevated colonies. *C.glabrata* colonies have a cream to white smooth appearance, but *C.krusei* colonies have a purple fuzzy appearance.(Himedia manual information). (Bardkar et al., 2010)



**Figure.2.7: Scheme for cultures study on differential agar media**

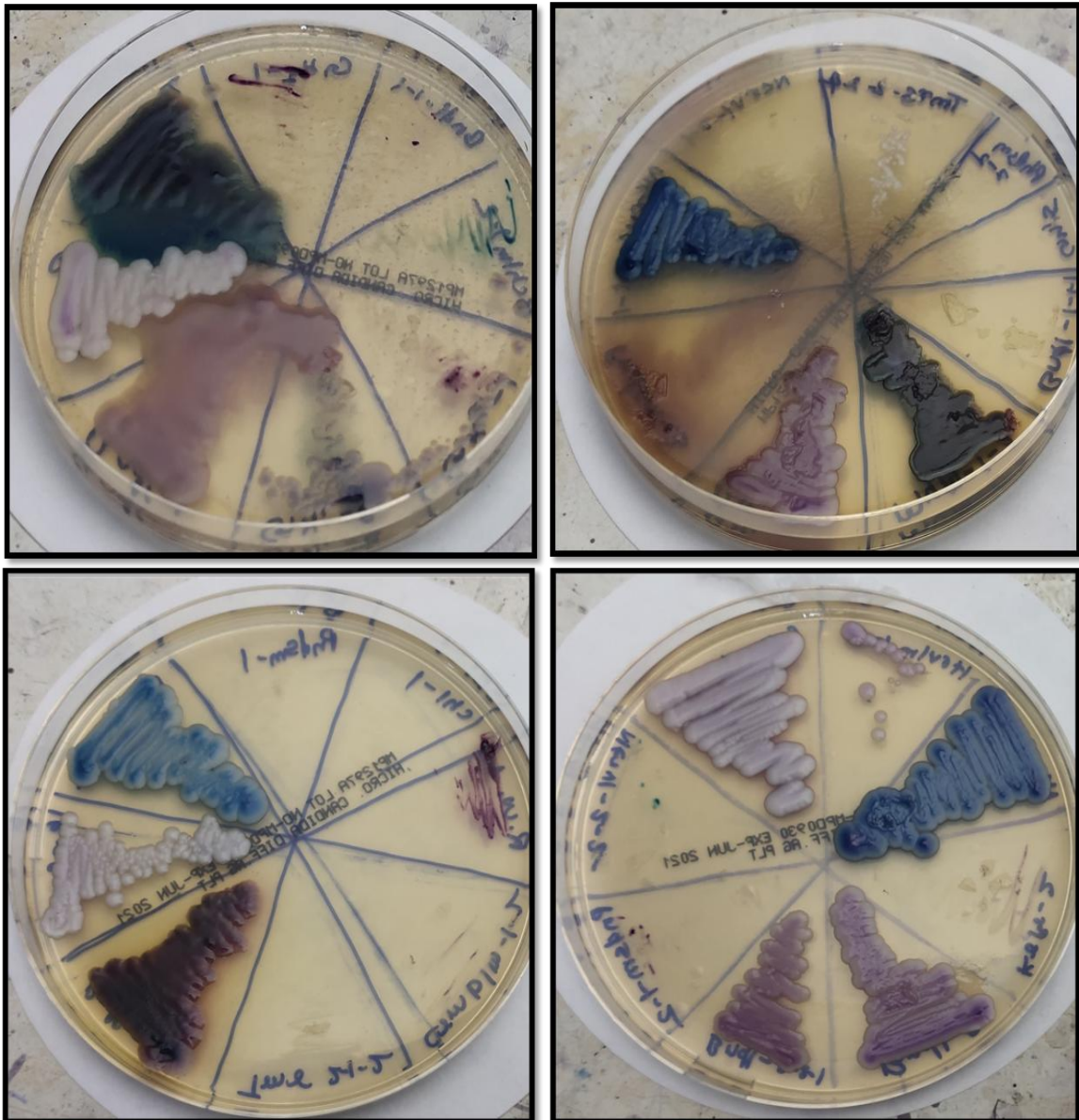
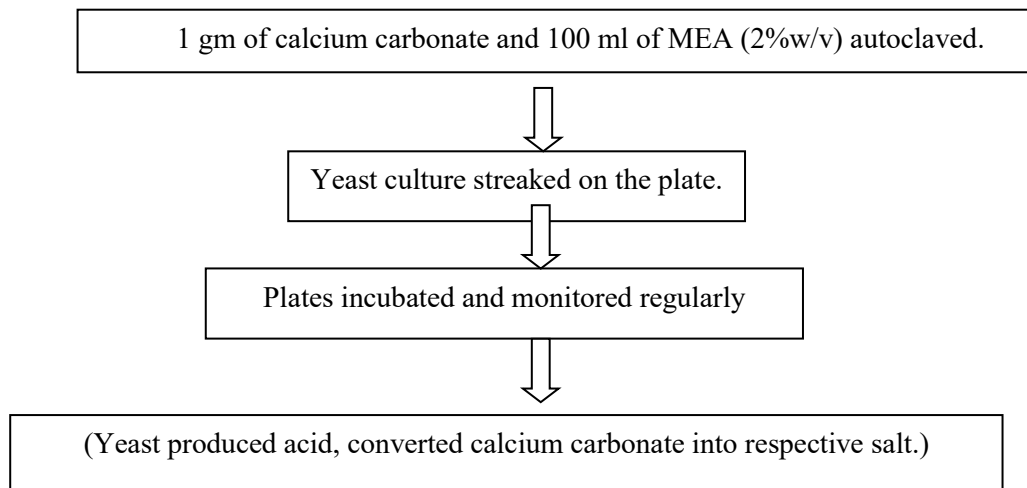


Figure. 2.8.: Cultures grown on differential agar media

#### 2.3.2.4. Organic acid production

During alcoholic fermentation, several important organic acids, such as oxalate, succinic, pyruvic, lactic and acetic acid, are produced by yeast. So test yeast evaluated for its organic acid production. (Campbell et al., 1996)

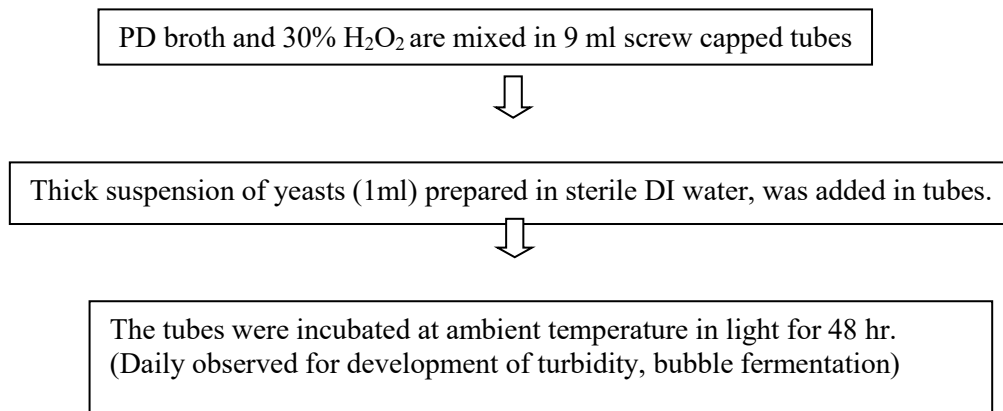
### 2.3.2.5. Standard protocol for organic acid production



**Figure.2.9: Scheme of acid production test**

### 2.4. Catalase production test

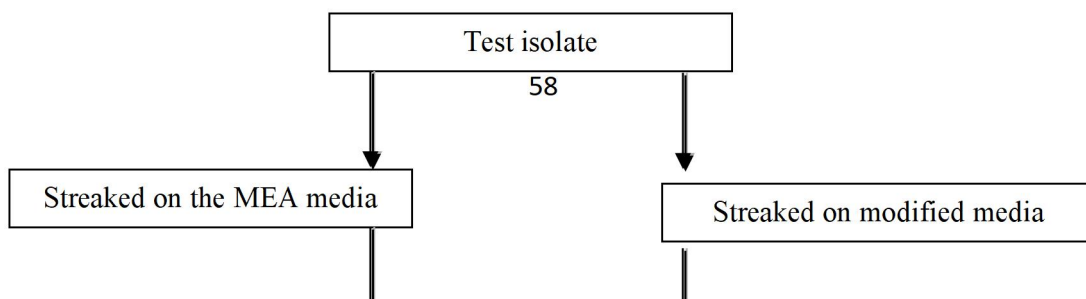
This test is based on the principle that yeast has high-powered peroxidases, such as catalase. Peroxidases are enzymes that act as catalysts in the oxidation of specific substances in living systems. Anaerobes do not produce catalase. (Dunford et al., 1999).



**Figure.2.10: Scheme of the catalase production test**

#### 2.4.1. Morphological study of promising yeast on modified media

Most promising strain was streaked on MEA media's plat and the plate prepared by sterile mine tailing soil and agar. Plates were incubated at 25±2°C for 48 hrs. Yeast grown on modified media were restreaked on MEA (2%w/v) media (commercially available) and observed under microscope.



### **Figure. 2.11.1: Scheme for promising strains's study in simulated environment**

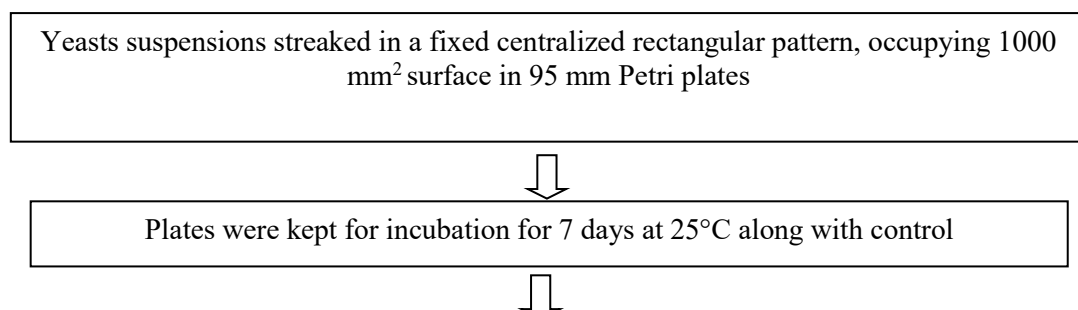
#### **2.4.2. Phosphate solubilization test**

Phosphate exists in both organic as well as inorganic forms in soil. Organic matter derived from dead and decaying plant debris is rich in organic sources of phosphorus. However, plants are able to utilize phosphorus from soil only in the free available form. Soil phosphates are rendered available either by plant roots or by soil microorganisms. Therefore, phosphate-dissolving soil organisms play a part in correcting phosphorus deficiency of crop plants. Turbidity is also caused by insoluble phosphate colloids including particle colloids.

The potential of wild ascomycetes yeast strains from polluted water bodies in the mining area to solubilize phosphate was tested.

#### **Phosphate solubilization efficiency of selected cultures**

Yeast culture streaked on the pikovskaya agar plates (Nautiyal., 1999) on predetermined pattern and incubate for 2-3 days at ambient temp (25-30°C). Colony growth and the phosphate solubilization zone were monitored on a daily basis.



**Figure .2.11.2: Scheme for phosphate solubilization efficiency test**

$$PSE = [(A_{BB} + A_{OB}) / 7089] \times 100$$

PSE= Phosphate solubilization efficiency

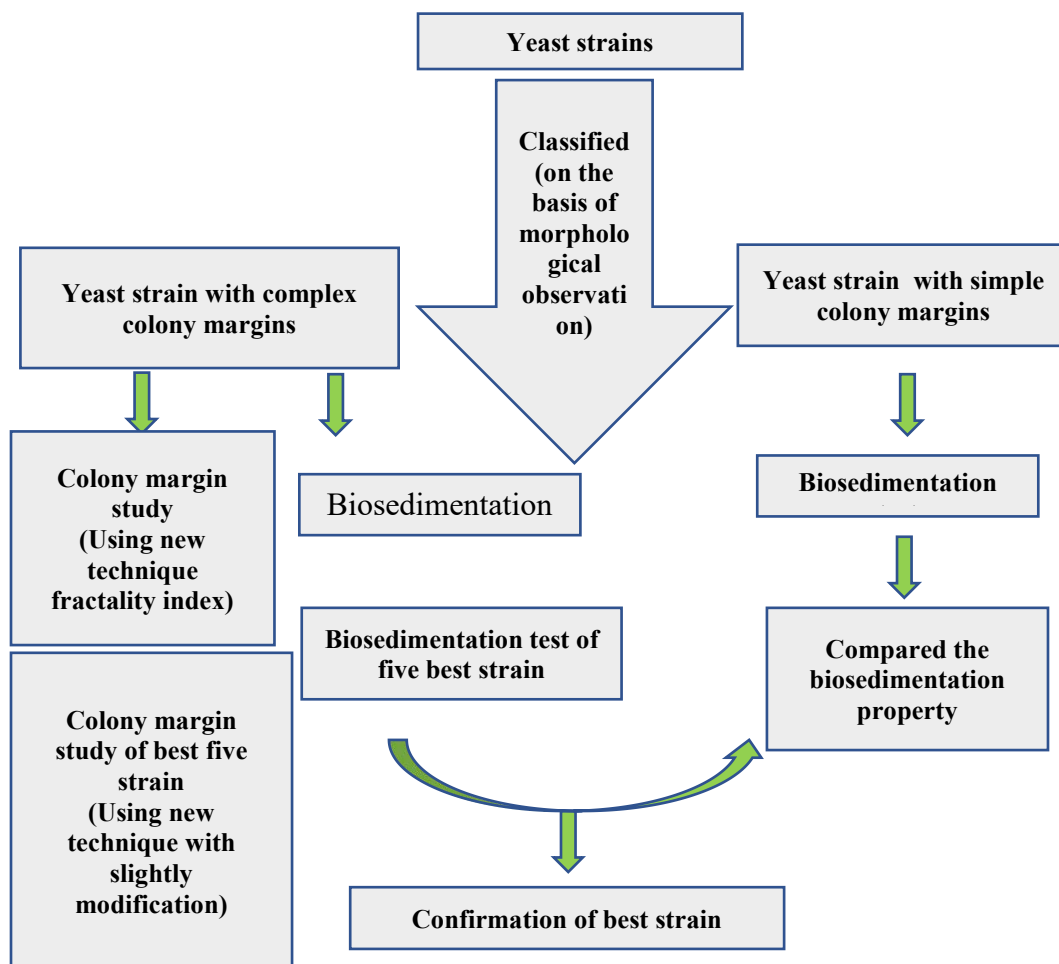
$A_{BB}$ =Area below biofilms

$A_{OB}$  =Area outside biofilm

**2.5. Screening of promising strain for efficient turbidity removal capacity**

During cultures's morphological study it has been observed that some strains were showed simple colony margins while some were complex colony margins. So, cultures were classified on the basis of their colony margins.





**Figure. 2.12: Scheme strategies for screening of promising strains for biosedimentation of SMTTW**

**Table.2.2 List of cultures with simple and complex colony margins**

<b>Cultures with complex colony margins</b>	<b>Cultures with simple colony margins</b>	
Bchlm-1	Bchlm-2	Sc-1
Bchlm-1-2	Srg-2-2	Srg-2-1
Gh1-1	Bndl1-1	Tmrs-2-2-3
Gh2-1	Pndsm-3	Pnd-2-2
Gh1-3	Gh1-2	Srg-1
Mym-1-2	Gh1-5	Tmrs-2-3
Pnd-2	Chl-1	Ntrvl-1-1-1
Pnd-3	Krpr-1	Ntrvl-1-2
Pndsm-1-2	Krpr-2	Ntrvl-2-2
Pndsm2-2	Bndl-1-1-1	Ntrvl 2-3
Tmrs-2-2-4	Bndl-1	-
Crblm-2	Bnd-2-2	-
Gh1-4	Crblm-1	-
Hrvlm-1-2	Srg-3-3	-
Bndl-3	Krpr-3-3	-

### **2.5.1. Screening of Strains**

Cultures were evaluated for their biosedimentation efficiency on the basis of visibility of tubes, biosediment compactness and settleability in 10 ml screw capped tubes by mixing 1 ml yeast cultures suspensions (same absorbance) and 9 ml simulated mine tailing water and scored according to table 2.3. Simple colony margins and complex colony margins key was used to classify strains

#### **(A) Visual screening**

All cultures were screened for efficient biosedimentation of clay colloids in turbid water by manual observation using biosedimentation efficiency scoring scheme as described in table 2.3.

Freshwater yeast strains were tested to see whether they could bind clay colloids found in SMTTW in 10ml screw test tubes and compared.

**Table.2.3. Scoring scheme for efficient biosedimentation**

<b>A</b>	Water column very clear no deposits on inner wall	100
	Water column very clear, some deposits on inner wall	80
	Water column partially clear no deposit on inner wall	50
	Water column Partially clear some deposits on inner wall	25
	Water column turbid	0
<b>B</b>	Sediment very compact	100
	Sediment just about compact	80
	Sediment loose	50
	Sediment very loose creates instant turbidity	25
<b>C</b>	After agitation of sediments	
	It breaks into fine particles takes long time to rain down to reform sediment	0
	Not as above breaks into large sediments and settles rapidly	100

### **2.5.2. (B) Screening of cultures by digital image analysis of colony margins**

Further, cultures with complex colony margins were screened with the help of JFRAD software, image analysis based technique. Puchkov (2016) has provided examples of using image analysis in the studies of both the macroscopic and the microscopic microbiological objects obtained by various imaging techniques. Fractality index of colonies margins was investigated, and an attempt was made to link the fractality index and biosedimentation efficiency. Those cultures with complex margins have been investigated for colony margin characteristics as a preliminary step and with modified method.

All cultures showing highest fractality index of colony margins were studied for bioclarification ability, sediments formation time and morphology of sediments were recorded and scored.

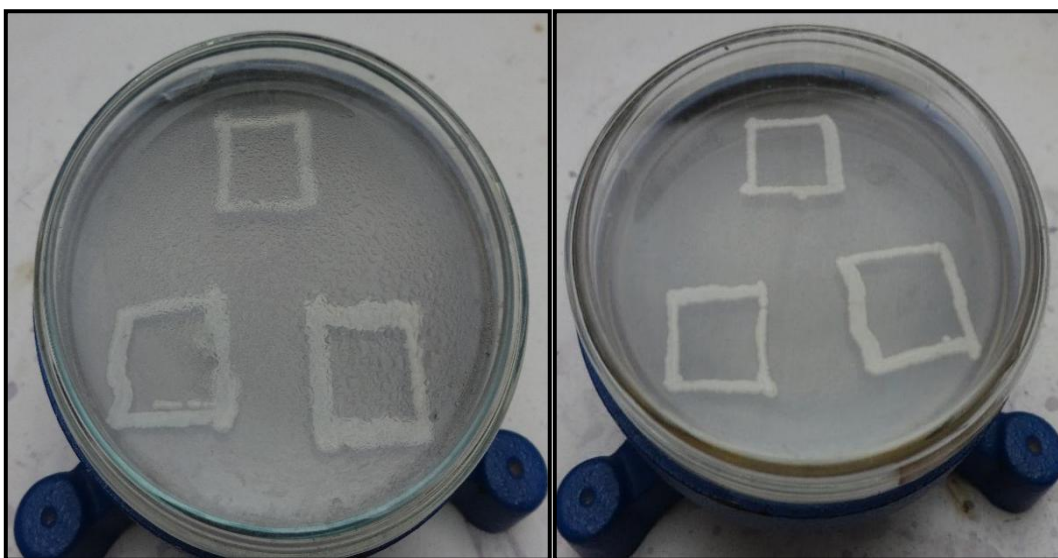
### 2.5.3. Fractality index

It is a four-digit number obtained by multiplying the score produced by the fractal analysis software like JFRAD by 1000 with last number being rounded up (Kamat, 2019). The detailed description of method is given below

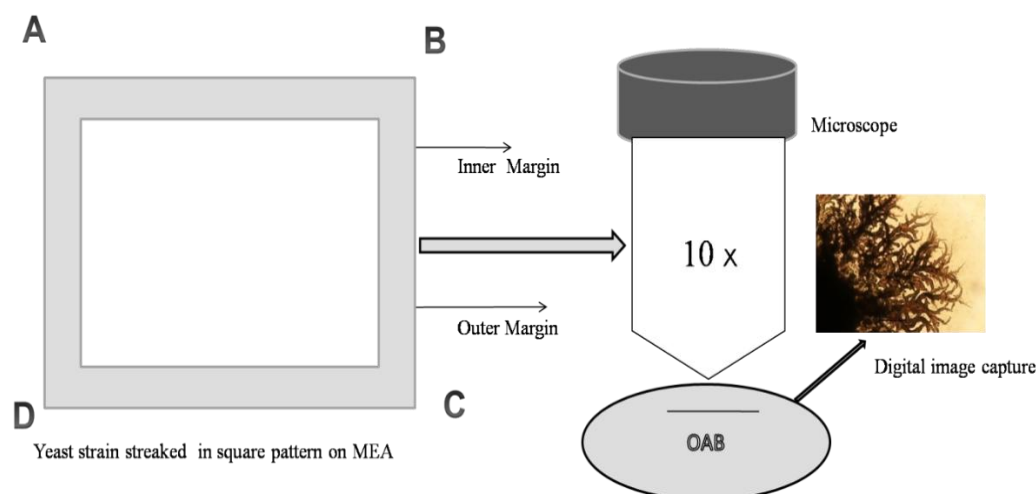
### 2.5.4. Primilary study of fractality Index of colon margins

Colonies were grown on 2% MEA in a square pattern to produce defined colonies (Fig.2.13), viewed under a 10X objective lense (Nikon microscope), and photographed the outer and inner colony margins (at same point) with NIS Element software up to 7 days, after the inoculation of strains on media (9 c.m. petriplates). Representative images were imported and processed to compute fractal dimension employing CMEIASJFrad version 1.0 software freely available at <http://cme.msu.edu/cmeias/> (Ji et al., 2015).

The output data of yeast colony fractal dimensions were saved as \*csv files and analyzed statistically using the SYSTAT 13. Fractality index of the margins were calculated and compared.



**Figure. 2.13: Streaking pattern of cultures for preliminary study**



**Figure.2.14: Scheme for colony image capturing**

### 2.5.5. Screening of cultures with complex colony margins for turbidity reduction

Colonies of selected cultures with complex colony margins were taken by nichrome loop, mixed to 10 ml of sterilised deionized water to make a yeast suspension with the same absorbance. In screw-capped tubes, 1ml yeast suspension was introduced to 9 ml SMTTW.

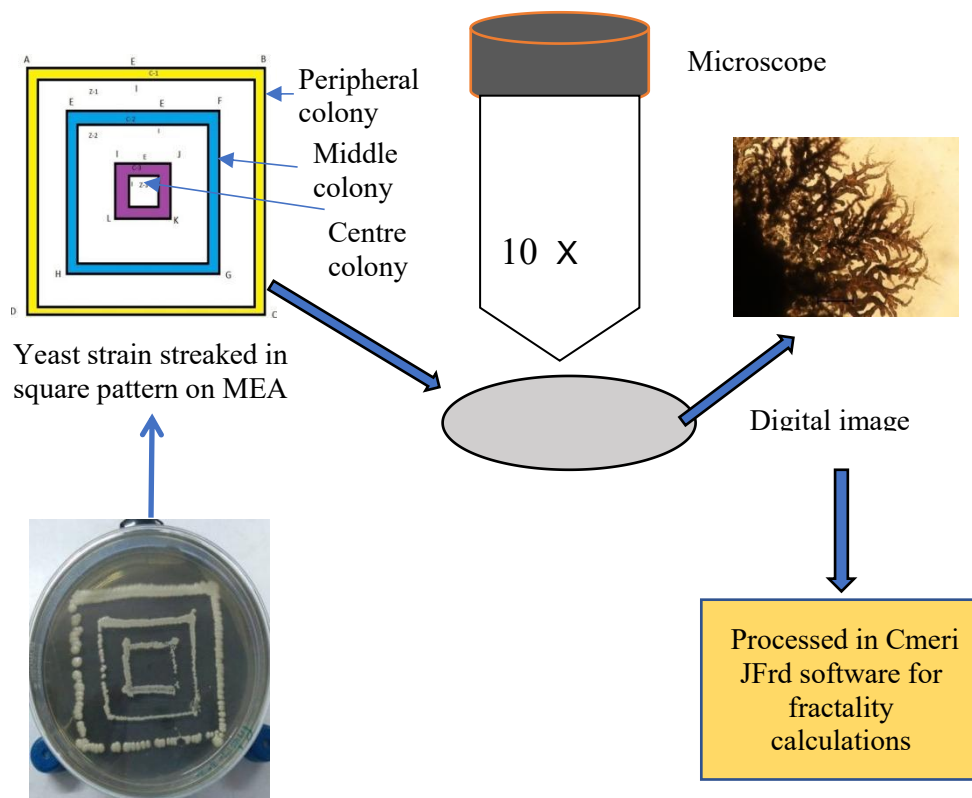
The fractality Index of biosediment has also been investigated, with the goal of correlating it with the fractality Index of strains colony margin for optimal biosedimentation of suspended solid in mine tailing water.

### 2.5.6. Study of fractality Index of colony margins with modified method

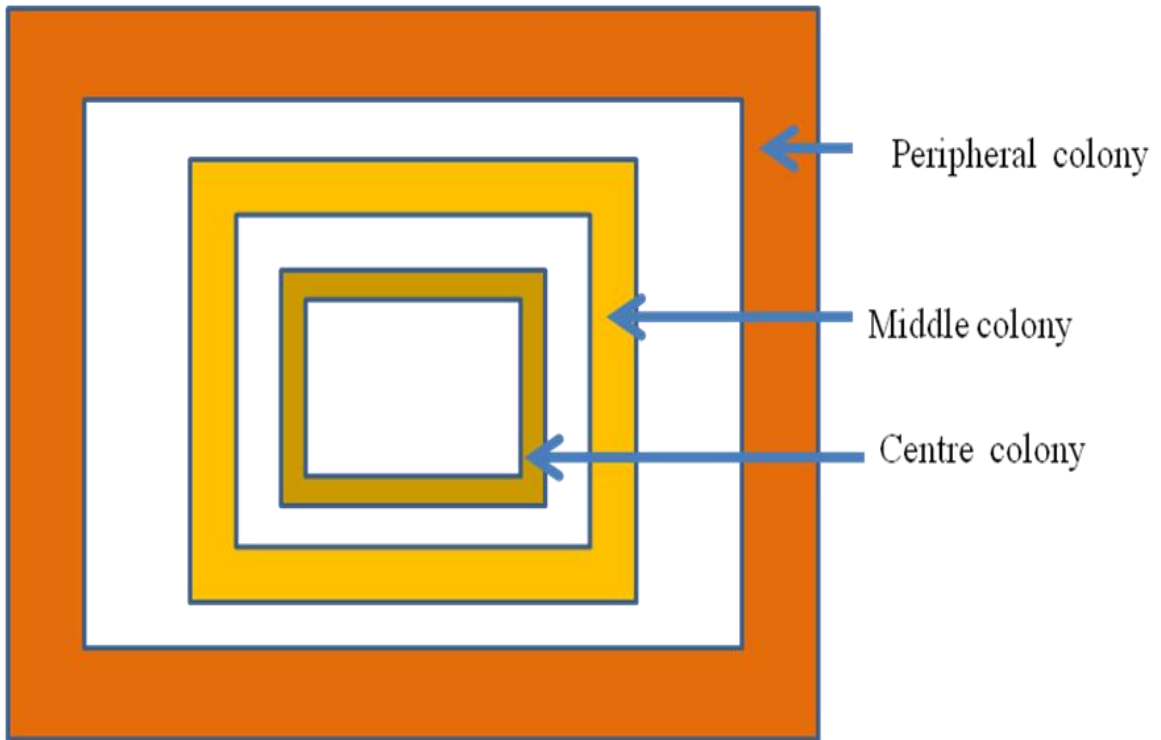
After a little change in approach, the best five strains with complex colony margin (among all preliminary tested strain) were studied again for colony features. Yeast strains were streaked on solid media in a nested colony square pattern (in 9cm Petri plates), allowing colonies to be observed under the microscope easily (Fig 2.16).

Images of colony margins were processed using CmeisJFrad software to determine the colony margins' Fractal dimension and fractality index calculated as described above.

Nested colony patterns allow for simultaneous observation of colony margin interaction on the same media on which cultures are injected. Furthermore, it has been discovered that the same method may be used to study strains adaptation to inadequate nutritional conditions.



**Fig.2.15: Scheme of colony image capturing after slightly modification**



**Figure.2.16: Predetermined pattern to obtain colonies amenable to digital analysis**

## **2.6. Assessment of drinking water quality of mining areas of Goa**

The sampling locations were chosen based on their location and intended purpose. Sampling was carried out on 6 stations during the monsoon and non-monsoon seasons to assess the quality of fresh surface water bodies. Water samples were taken and analysed for two seasons in a row. Surface water was sampled at every location. Furthermore, several of them were using for drinking purposes. A portable temperature and TDS meter was used to measure the temperature (Multi thermometer) and TDS (HM Digital TDS-3 meter) in the field. The physicochemical parameters of water samples were investigated.

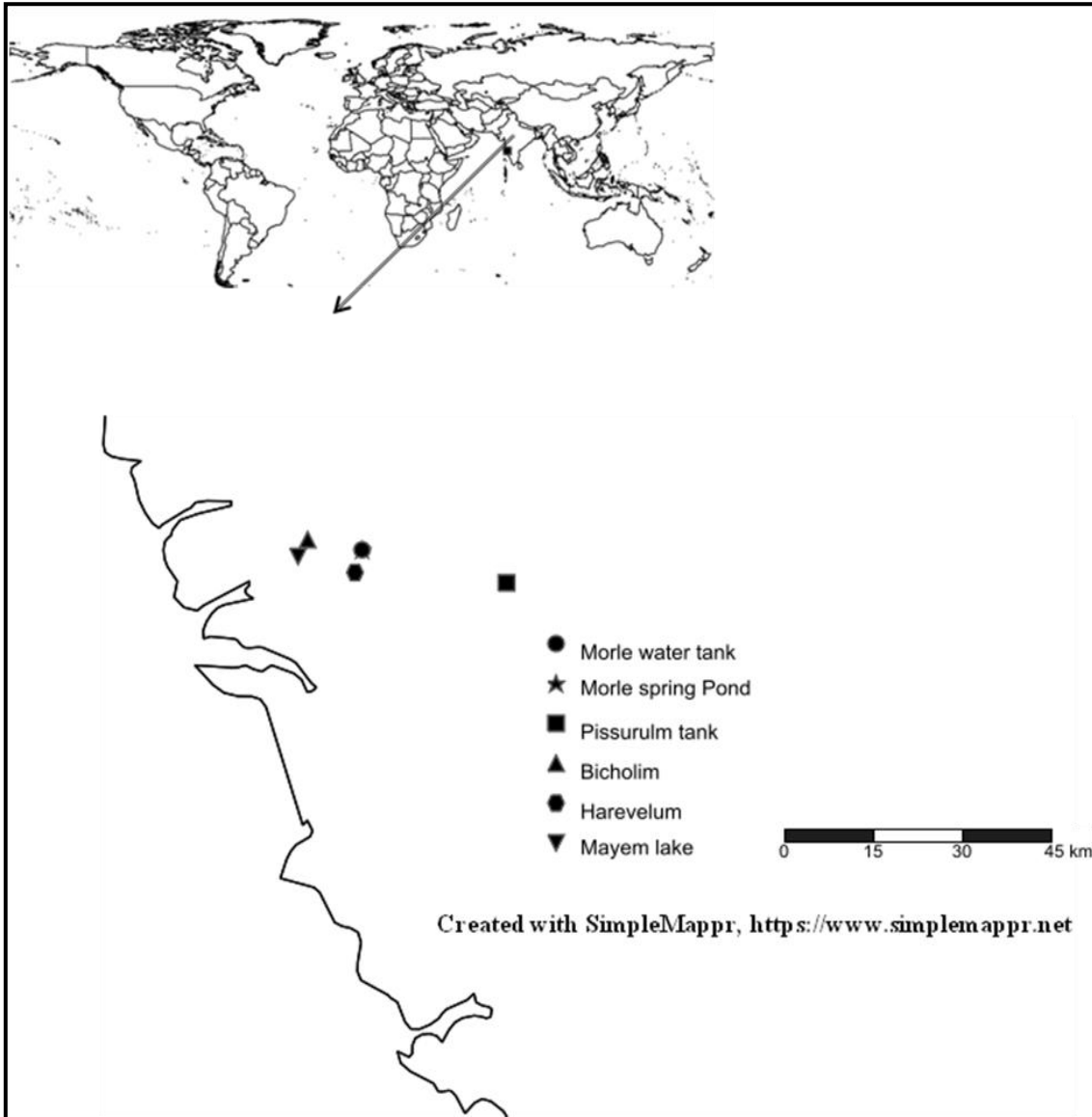
The goal of this study was to determine the physicochemical parameters of surface water in the Goa mining areas.

This research will include data on the quality of water bodies as well as their environmental status. The correlation coefficient between water quality metrics has been calculated in a systematic manner.

The APHA's (1998) standard prescribed protocol, which is a standard protocol for water quality characterization all around the world, was used to develop the methodology for water sample characterization. Plastic canes were used to collect the water samples. During the sampling, grab sampling was commonly used.

Himedia India procured analytical-grade chemicals for the current study were used. All of the glassware were standard, and it was made by Borosil India Ltd. Standard procedures were used to conduct the analysis.

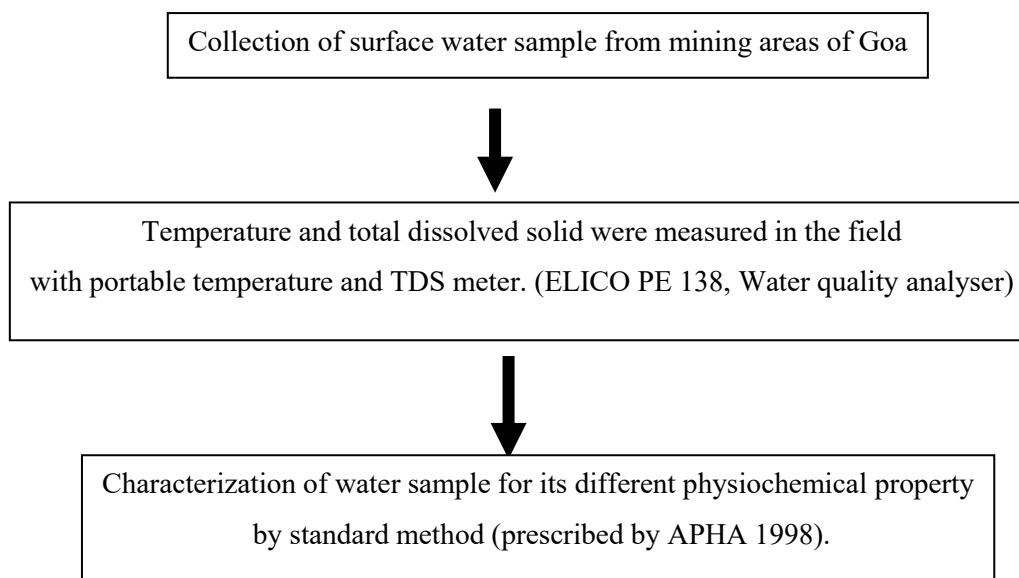




**Figure.2.17: Water sampling sites**

**Table.2.4. Geographical co-ordinates of locations**

Place	Longitude/ Latituted
MorlemWater tank	15.58611 °N,74.0375 °E
Morlemspring pond	15.5861 °N,74.0375° E
Pissurlem tank	15.5351 °N,74.2574 °E
Bicholim tank	15.6018 °N, 73.9546°E
Arvalem fall	15.5511° N, 74.0267° E
Mayem lake	15.5760° N, 73.9400° E



**Figure.2.18: Scheme for water analysis**

**Table.2.5 Water analysis methods and instruments**

Water Parameters	Instrument/ Methodology	Water Parameters	Instrument /Methodology
Turbidity	Thermo scientific EU Tech TN 100	Cl	Iodometric
TDS	HM Digital TDS meter	NO <sub>3</sub>	UV -100 SHIMAZDU
EC	ELICOPE138	SO <sub>4</sub>	UV -100 SHIMAZDU
pH	pH meter	PO <sub>4</sub>	UV -100 SHIMAZDU
Temperature	Multi-thermometer	Alkalinity	Titration
DO	ELICOPE138	Total hardness	Titration
Na	Flame photometer	K	Flame photometer

### 2.7. Biosedimentation test using promising strain

Following the screening of the most promising strain for biosedimentation of clay colloids in SMTTW, the strain was further investigated for its turbidity reduction quality on a wide scale using various experiments. Furthermore, prior to the large-scale trial, the biosedimentation technique was standardized to improve strain efficiency for larger-scale biosedimentation tests.

Process and conditions were optimized for the efficient biosedimentation of clay colloids. SMTTW was prepared by adding the mine tailing soil in distilled water. Yeast was grown in MEA broth and yeast suspension was prepared in sterile deionized water (all chemicals used were of HiMedia Chemicals Ltd., India).

Biomass's dose, age and pH of turbid water (4 to 9) estimation was performed for efficient biosedimentation, using sedimentation efficiency scored scheme (Table.2.3). Further, biosedimentation study was performed in the laboratory at the scale of 10 ml, 1L, 20L, 100 L. Simulated turbid water and yeast suspension (with desired concentration) were mixed in the ratio of 1:9 respectively at every scale. Water column stirred vigorously at 50 rpm for 1min smaller scale, 10 mins higher scale and water samples withdrawn with glass pipette at different time intervals ( $t_1=0$  min,  $t_2=10$  min,  $t_3=20$  min,  $n=120$  min. Turbidity were measured of withdrawn samples. (Thermo scientific EU Tech TN 100).

Biosedimentation scoring scheme is independent for screw capped tube and Imhoff cone.

Concentration of yeast in yeast suspension – 0.005gm/l Three yeast suspensions with 0.005gm/l were prepared.

Ratio of yeast suspension and SMTTW 1:9. (to attain more water treatment using less yeast biomass.)

### **2.7.1. Preparation of log phase Yeast suspensions by both method on plate and broth.**

During various trial experiments, standardized yeast suspensions of known concentration were generated in the laboratory. Diluted MEA broth (0.4%w/v) in deionized water was used to create yeast inoculums in Erlenmeyer flask. And then put it on the rotator shaker for 48 hrs at 25 °C and 150 revolutions per minute (Fang.T.J. et al., 1996) (Scigenics Biotech orbitek ). The inoculums were centrifuged for 10 minutes at 10,000 rpm. In the supernatant, sterilised water was added and centrifuged twice under the same conditions. Furthermore, yeast colonies were grown in bulk on MEA media in Petri plates incubated for 48 hr at 25±2°C, scratched with a sterile blade, and mixed with sterile deionized water to prepare a yeast suspension of the desired concentration.

Because this weight was specified for quick biosedimentation in previous investigations, yeast suspension (0.005gm/l) was made by dilution with sterilised water.

Turbidity reductions were investigated using cultures harvested using both procedures, and the turbidity reductions achieved by both yeast suspensions (cultivated by different methods) were compared .

### **2.7.2. Characterization of SMTTW.**

SMTTW has been characterized for its pH, temperature (Multi thermometer), SPM by standard method, electrical conductance, (Aquaro digital water conductivity meter), and TDS (HM Digital TDS-3 meter).

### **2.7.3. Addition of yeast in SMTTW and observation of the reduction in turbidity**

0.005gm/l (dry weight) of yeast suspensions was mixed into simulated mine tailings turbid water (1:9). Whole water was stirred with the help of glass rod. Water samples were withdrawn at different interval of time to measure the turbidity with turbidity meter (Thermo scientific EU Tech TN 100).

Moreover, biosedimentation procedure has been evaluated by observation of complex sediments formation and its rapid sedimentation. Further biosedimentation of clay colloids using yeast had been compared to control, (in which yeast suspension were not added in turbid water) under the laboratory condition.

#### **2.7.4.1. Biosedimentation of SMTTW using heat treated yeast cells**

The composition of the yeast cell wall is responsible for the attachment or adsorption of components. Clay colloids cause colloidal turbidity in mine water, and this notion has been utilised to remove it. As in a prior experiment, viable cells were used to reduce colloidal turbidity Yeast suspensions (0.005gm/l dry weight) were heated with a spirit lamp for various periods of time and then tested for sedimentation properties in screw capped tubes and the Imhoff cone. The biosedimentation patterns of live yeast and heat-treated yeast were also compared.

#### **2.7.4.2. Heat treatment of cells**

The most promising strain, Bchlm-1-2 suspension with known concentration, was heated with a spirit lamp for 60 seconds before being introduced to 900 ml of SMTTW in an Imhoff cone.

### 2.7.4.3. Strains combination study for efficient biosedimentation

Strains were studied for their combination to reduce turbidity of SMTTW. 0.5 ml of each strain suspension were mixed with same absorbance and added in 9ml SMTTW, scored for its biosedimentation efficiency according to table 2.25.

## 2.8. Biosedimentation test upto 10 ml to 1L

Mine tailing soil brought from mining area were dried in room temperature crushed properly using ceramic mortar pestle. Soil were sieved using sieve with different mesh size. 0.57 um was the last sieved. 5.6 gm of soil was added in 1000 ml of distilled water and stirred vigorously (50 rpm for 10min). After 45 min decanted soil water was used for biosedimentation study. Yeast suspension and SMTTW with desired concentration were used in 1: 9 ratio to maintain the less amount of yeast biomass. Compactness of sediment observed (according to table 2.25) and turbidity were measured in 1L of imhoff cone. Turbidity reduction in test cone were compared with control cone.

### 2.8.1. Turbidity removal efficiency

The data acquired after executing the biosedimentation test up to the scale 1 L and 1000 L were used to compute the adsorption efficiency of turbidity reduction of SMTTW. The adsorption efficiency of the biosedimentation test was calculated using the NTU reduction values of the control and test's reactors.

$$\text{Turbidity removal efficiency} = \frac{C_f - C_o}{C_o} \times 100$$

where  $C_o$  is the initial turbidity of the water sample (NTU) and  $C_f$  is the final turbidity of the water sample (NTU). The experimental studies were repeated thrice to check its repeatability and the average values only were discussed in the report.

$C_o$  = initial turbidity

$V$  = Volume in L

$C_f$  = Final turbidity

$m$  = mass in m

(Kumar P.S. et al 2016, Nirmala rani 2010)

### 2.8.2. Surface charges of Yeast cells

Alcian blue is a positively charged phthalocyanine complex that is absorbed by negatively charged yeast cell surfaces, particularly the mannosylphosphate. This dye has

been used to calculate changes in the charge on the cell surface. The degree of this dye's adsorption is proportional to the negative charge on the cell surface.

Yeast cells at a concentration  $5 \times 10^7 \text{ ml}^{-1}$  were washed twice in phosphate buffer (pH = 7.0) and harvested by centrifugation at 143Xg for 5 min at 4° C (Research centrifuge TC 4100F). Then yeast was suspended in 0.02 mol L<sup>-1</sup> sodium acetate buffer (pH = 4.0) and washed twice with the same buffer. Yeast was incubated with 1.8 ml of Alcian blue tetrakis-chloride solution (50 mg L<sup>-1</sup> in the buffer for 30 min at 25°C. After centrifugation at 10000 for 10 min at 20°C (Research centrifuge TC 4100F) the supernatant was decanted and its absorbance was measured at 615 nm (UV-100 SHIMAZDU) using uv-spectrometer. The Alcian blue retention ratio (ABR) was calculated according to the following formula, The ABR was expressed as the mean of three experiments. (Fukudome et al., 2003).

$$\text{ABR} = (\text{A}_{\text{ABSOLUTION}} - \text{A}_{\text{SUPERNATANT}}) \times 100 / \text{A}_{\text{ABSOLUTION}}$$

### **2.8.3.SEM EDX FTIR of sediments**

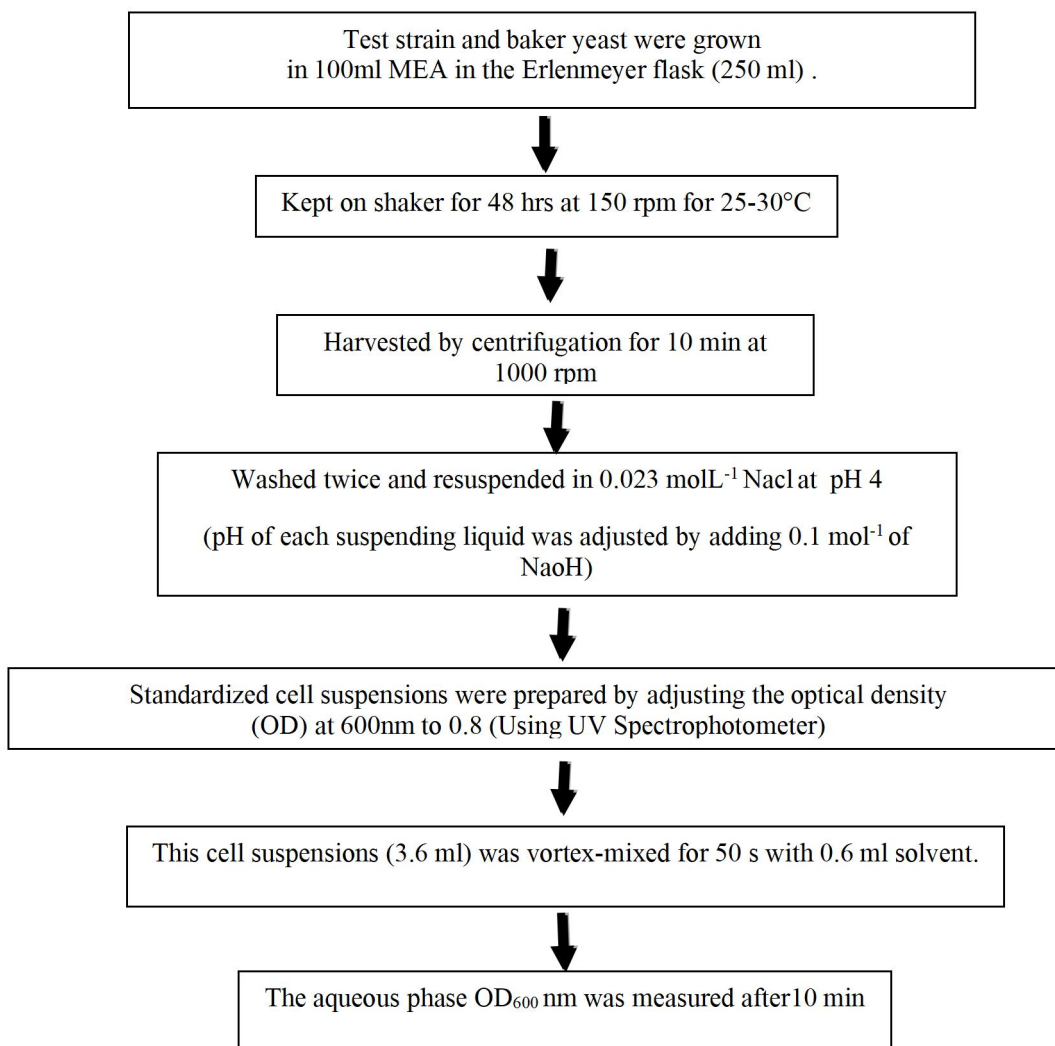
Scanning Electron Microscopy (SEM) was used to analyse the surface morphological structure of purified yeast, control sediment, and test sediment (Evo-18 Carls-Zeiss, India). EDS recorded with a JOEL-JSM 5800 LV was used to determine the elemental composition of the samples.

Cells were suspended in osmium tetroxide (2 %, w/v), incubated statically for 60 min at room temperature, and harvested by centrifugation (13000 r.p.m. for 1 min). The fixed and stained cells were dehydrated by 10 min incubations with gentle agitation in a graded ethanol series of 10, 20, 30, 50, 70, 100 and 100 % dried absolute ethanol. Cells and sediment were prepared for electron microscopy by air drying for 24 hrs room temperature. Sample were processed after coating, mounted on SEM sample stub, with double sided sticky tape (Barker and Smart, 1996, C.D. Powell et al. 2003, Das et al., 2016).

Control sediment and test sediment was subjected to Fourier Transform Infrared (FTIR) spectral characterization using, Shimadzu with FTIR grade KBr powder (Reddy et al, 2018).

#### 2.8.4. Hydrophobicity

Hydrophobicity was determined using a solvent approach described in a prior study (Solvent method) (Amar et al., 2006). The microbial cell adhesion to solvent test compares microbial cell affinity for a polar and nonpolar solvents. Baker yeast used as control.



**Figure.2.19: Scheme to calculate cell surface hydrophobicity**

Results are given as, where  $A_0$  and  $A$  are  $OD_{600}$  of the aqueous microbial suspension before and after mixing, respectively. The hydrophobicity in this method is defined as the cell adhesion to hexadecane. (Amaral et al., 2006).

$$\text{Percentages of bound cells: \% adhesion} = 1 - A/A_0,$$

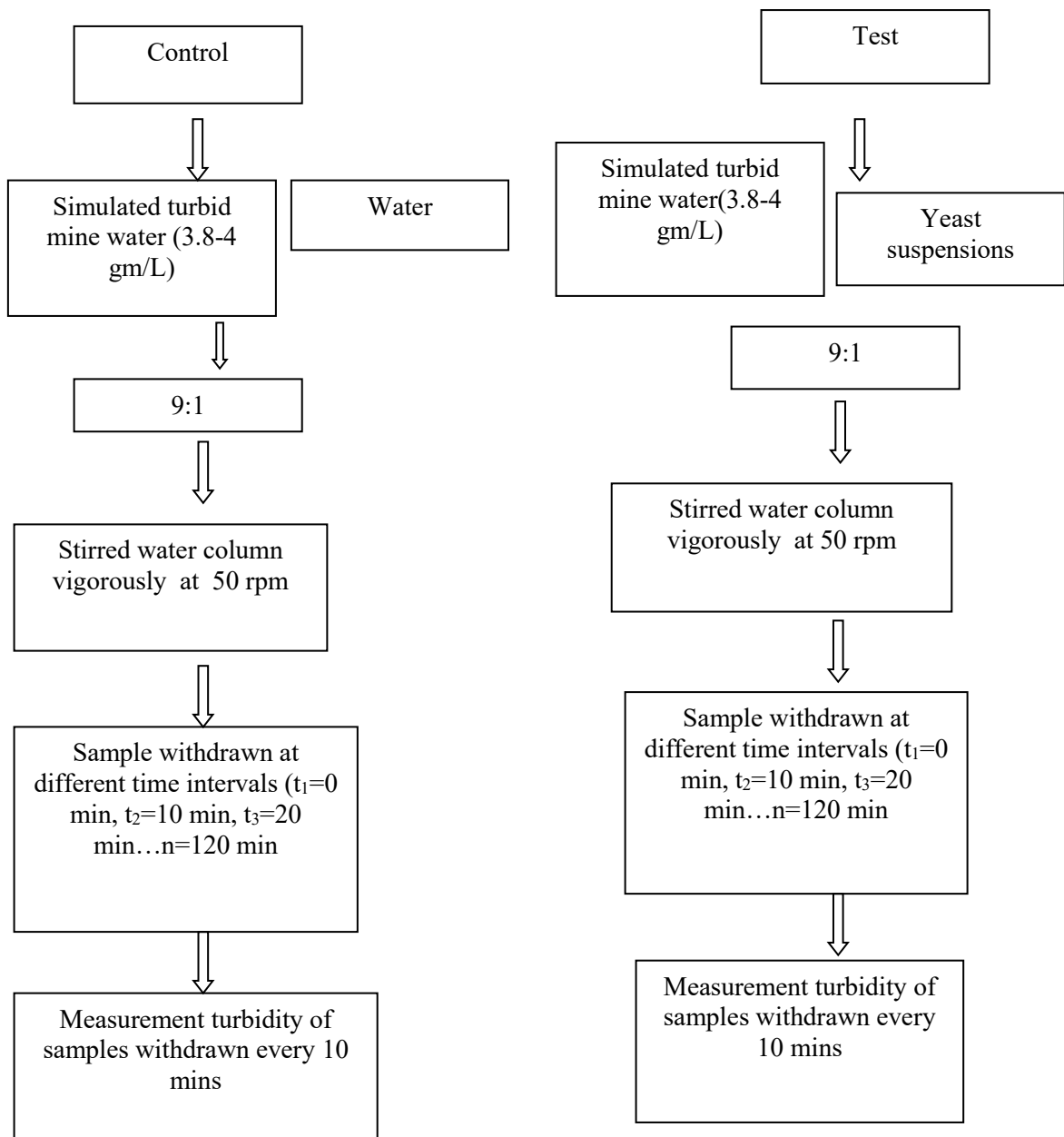
## **2.9. Lab scale biosedimentation using natural yeast cultures upto scale of 1 to100L**

Any treatment technique's progress or dependability can be tested on a bench scale in the laboratory. In the laboratory, the removal of turbidity from SMTTW using yeast cultures has been tested up to a size of 1-100 L. As a laboratory test, it may be used to optimize the treatment process in order to facilitate the treatment procedure on a wider scale. It saves money and time, and it can assist in the design of the treatment plant. Many biological methods for wastewater treatment have been effectively investigated at bench scale (Sheoran et al., 2005). The quality of SMTTW was measured before and after it was treated with yeast for turbidity removal. The results were compared in order to assess the biosedimentation capacity of the yeast strain.

The lab scale study provided insight on how to improve the method on a larger scale. The general goal of this research was to optimize the biosedimentation technique for application on a larger scale, as well as its behavior, such as sedimentation rate, HRT, sediment formation, and sediment wall deposition.

With the results of this research, a plan or approach for removing turbidity from SMTTW using yeast at a larger scale can be developed.





**Figure.2.20: Scheme for the biosedimentation test**

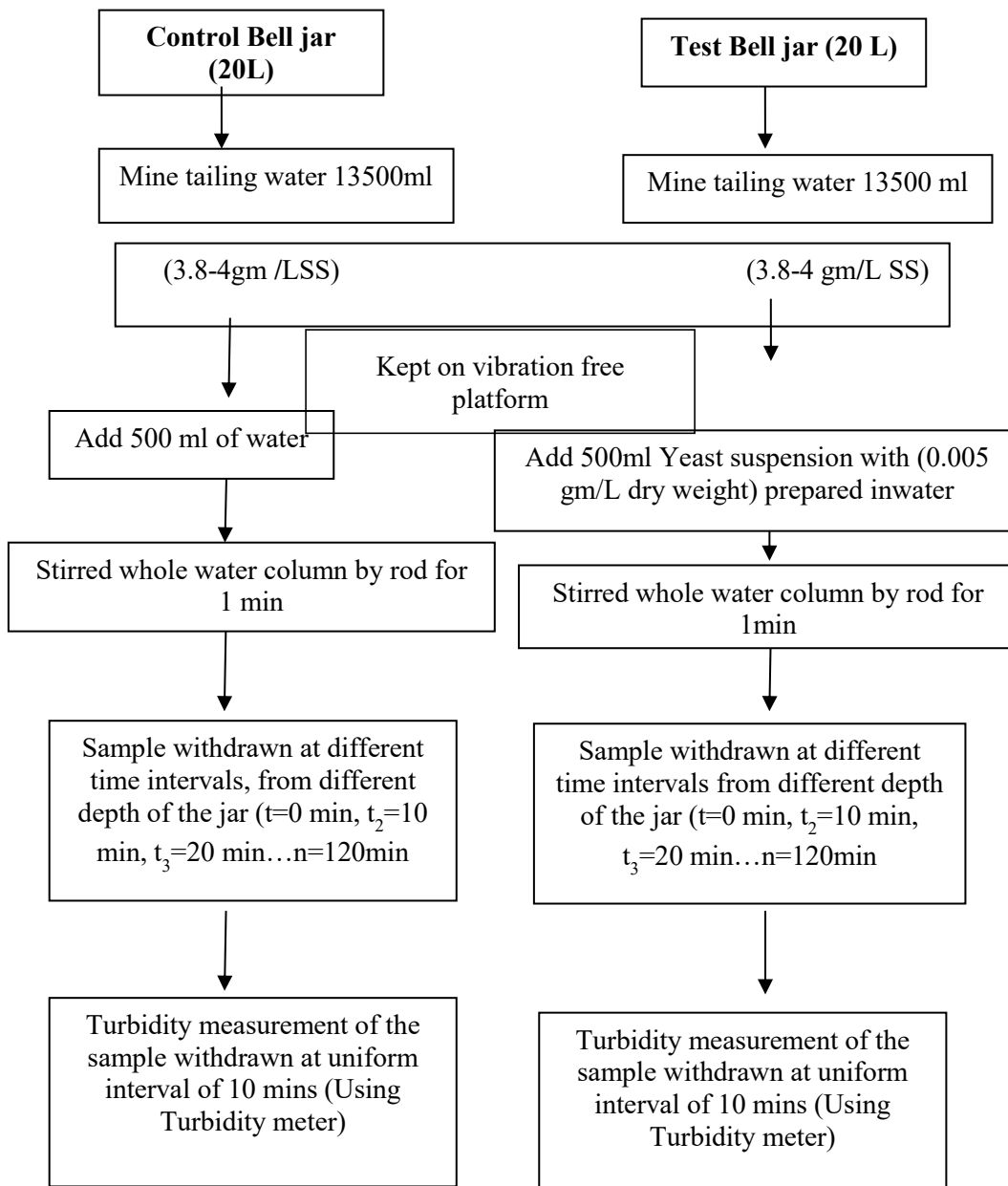
### **2.9.1. Turbidity reduction study of SMTTW in Bell jar using test cultures**

Biosedimentation of clay colloids in SMTTW by yeast cultures was done and standardized at a lower scale in prior research. Biosedimentation tests on a larger scale, ranging from 14 to 100 L, were carried out in a bell jar and a columnar plastic tank. Yeast were cultivated as discussed previous section, Yeast culture was employed after the fourth day of inoculation because this harvesting time had been standardised in prior experimental studies. And standardised yeast suspension was used. Experiments were carried out at room temperature with a pH of 6.8 in the medium throughout.

13500 ml SMTTW was placed to a 20 L bell jar, and 1500 ml yeast suspensions (0.005 dry weight) were added to the SMTTW (temperature 26.9 °C pH 6.4). Water samples were withdrawn with a glass pipette at varied time intervals and from different depths of the bell jar. The turbidity of the removed water sample was determined. The biosedimentation pattern and reduction in turbidity with time were investigated in a control bell jar in which yeast suspension was not added. The turbidity of water samples taken from the control tank at various depths was also assessed. Mean of the turbidity of both control and test jar were compared.

### **2.9.2. Microscopy of aliquots**

Microscopy of aliquots collected from different depth of bell jar was performed under 10x objective lense using light microscope.



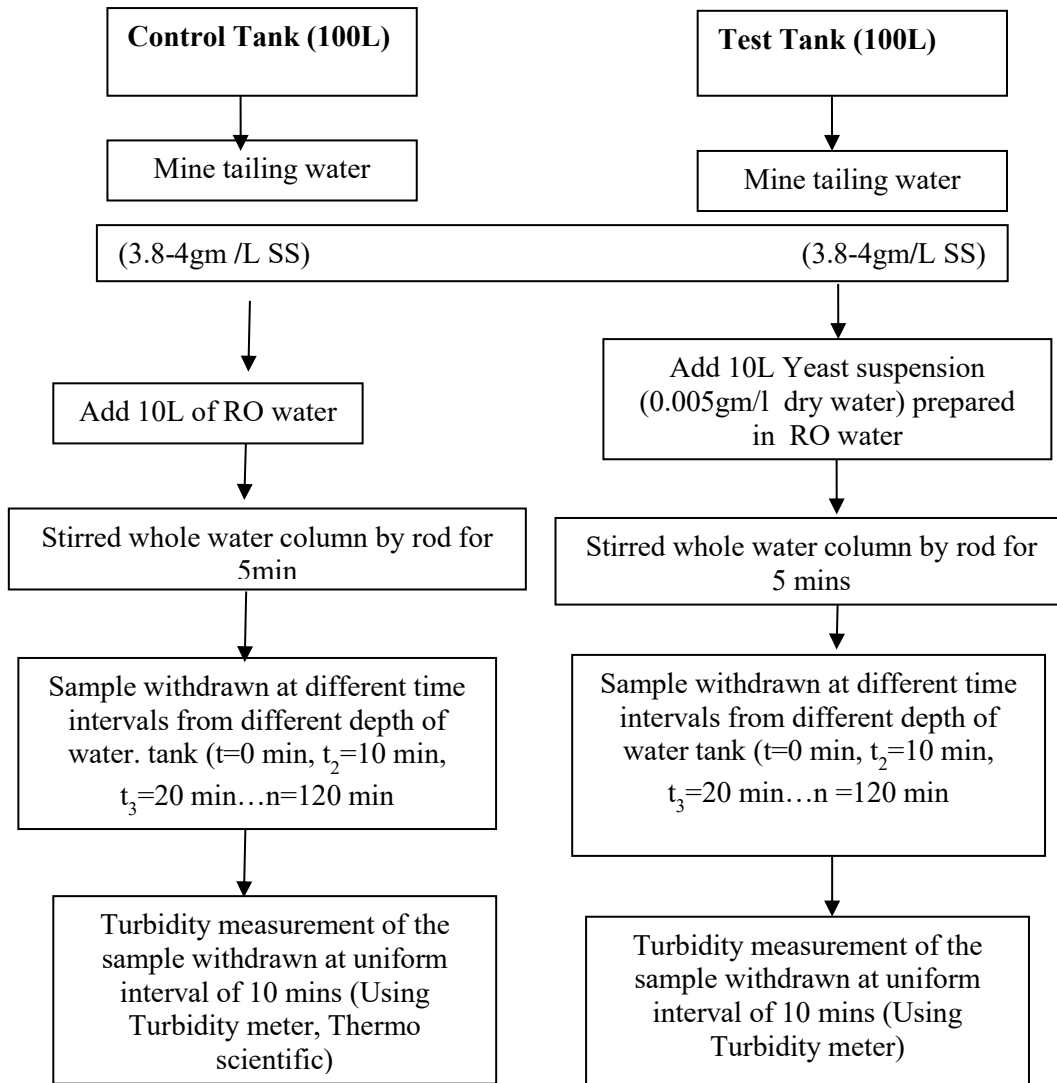
**Figure.2.21: Biosedimentation test in the Bell jar**

### **2.9.3. Biosedimentation test in 100 Litre of tank**

In the tank, 90 litres of mine tailing water (prepared in reverse osmosis water (Suspended solid 3.8) was added, along with 10 litres of yeast suspension with desired concentration. Furthermore, the turbidity reduction of a control tank in which yeast suspension was not added was investigated over time at various depths.

Whole suspensions were agitated thoroughly with a rod, and samples were taken from various depths in the tank, and turbidity was assessed across time intervals.

After two hours, the entire medium was agitated for two minutes and the sample was collected once more. Turbidity was also assessed in water samples collected at various depths in the tank and at varied time intervals.



**Figure.2.22 Biosedimentation test in 100 L tank**

#### **2.9.4. Sand bed filtration test at laboratory condition**

The sand bed filtration technique is a common method for removing turbidity from water. It usually reduces turbidity by 0.1-1 NTU. It does, however, necessitate regular cleaning or backwashing, as well as pump operation. The sand bed filtration procedure necessitates both pre-treatment (coagulations) and post-treatment (biosedimentation) (e.g. Disinfection chlorinations.). It works by percolating water via a sand bed. It is not suited for cleaning extremely turbid water directly; however, it can be utilised after the coagulation, biosedimentation, and sedimentation processes.

#### **2.9.5. Design of sand bed filter in laboratory**

A 0.60 m glass column with a 0.004 m diameter was filled with 1.47g/cm<sup>3</sup> sand. The sand bed column was allowed to pass treated water with NTU 190. Using a peristaltic pump, the flow rate was kept at 5 ml/min. In order to maintain constant turbidity and avoid particle sedimentation owing to gravity, the reservoir water was agitated continually using a magnetic stirrer set to 70 rpm.

#### **2.9.6. SMTTW treatment using Sand bed filter**

A sand bed filter with a flow rate of 5ml/min was used to filter mine tailings and treated water. To keep the turbidity of the entire medium constant, water was kept on a magnetic stirrer. The turbidity was measured at different rpms to normalize it. At 70 rpm, the turbidity of the entire medium remained constant. At various intervals of time, samples were collected and analyses for their properties. To observe microbial growth, samples were plated on malt extract agar (2%) and nutrient agar. The effluent of sand bed filtration was also tested for turbidity reduction, pH, Fe and Mn concentration, and other water parameters.



**Figure.2.23: Sand bed setup at lab scale**



**Figure.2.24: Petriplates spread plated with Sand bed effluent**

**(a) Nutrient agar (b) Malt extract agar**

### **2.9.7. Standardization of biosedimentation process for field level application scale 500 -1000 litres**

The observed laboratory results were encouraging and have inspired to scale up the volumes for relatively high-volume experimentation beyond laboratory scales. The strategy adopted was gradual scaling up. Hence initially a reactor volume of 500 liters was considered for experiment and it was tried for three times. Subsequently, the volume of 1000 litres were considered for reactor. The other benefit of this scaling up is that it may provide us with the possible deviation factor when scaled up.

The process of natural rate of biosedimentation of SMTTW's suspended solid under ambient conditions can be enhanced by using yeast as biosedimentaion agent to enhance the rate of biosedimentation and general properties of sediments.

#### **2.9.7.1. Environmental concerns**

The general environmental concern for any additives in reactors is that the product or additive should not leave behind residues in the stream especially when the life cycle assessment has not been carried out. Hence the microbial growth was observed in the outlets of control as well as reactor outlets and their respective sand filter outlets. These results are also encouraging.

Effective actual availability of yeast per gm or per ml of turbid water will definitely differ due to the hindrance of the particle co-hesive adhesive forces within liquid phase or the other chemical and physical compatibilities forces between the particles due to heterogeneity.

Area selected for performing the pilot study is based on representative condition. DM's college is located at plateau, not very far from mining area. Tailing soil sample is collected from Codlem mines of Vedanta Resources located in Dharbandora Taluka, was a part of representative sample collected.

Sediment concentrations cannot be determined easily or quickly in the field, and transportation to a laboratory for analysis is time-consuming and can be costly (Thackston and Palermo, 2000). As a result, these traditional methods are increasingly being replaced in favor of accurate, continuously-collected surrogate data for quantification of suspended solid that may be safer and (or) less expensive to obtain, such as turbidity measurements.



### 2.9.7.2. Preparation of SMTTW



**Figure.2.25: Mine tailing soil samples**

Total amount of soil was arranged in circular heap and divided into 4 parts (Fig.2.25). Soil samples were taken from the top, middle & bottom parts of the heap and mixed properly. Composite mine tailing soil was grinded with the help of ceramic mortar to get homogenous mixture of representative soil sample. 10 kg of mine tailing soil sample was added in the 100L water tank to prepare slurry. This suspension was stirred for ten min and left for 24 hr. After 24 hr suspension was restirred.

After 10 min of stirring, soil suspension was added in 800 L of water in the 1000 L of tank. (Turbidity was measured 860 (mean) and suspended solid (3.8-4 gm/L) of SMTTW). The total volume of 900 L was attained.

### 2.9.8. Preparation of Yeast Suspension (0.005mg/ml, Dry weight)

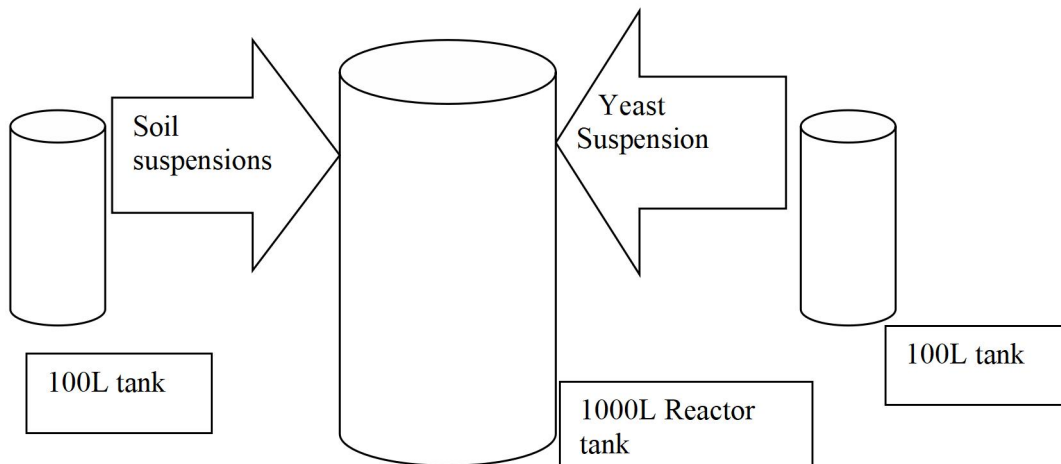
- Since 0.005mg/ml yeast suspension was required for turbidity reduction. (Based on standardization in previous experiment in lab)
- For 1000 L mine tailing (reactor volume) 100 L of yeast suspension required.
- Since, the desired strength of yeast is 0.005 mg per ml it will become 0.005gm in 1 Liter, and 0.5 gms in 100 L.
- So, from 20 gm /L(Dry weight) stock of yeast, actual volume required to be taken was calculated as below

$$\frac{1000}{20} \times 0.5 = 25 \text{ ml}$$

- Hence, it comes out to 25ml of yeast suspension in 99.975 L of water.

- 100 L suspension of yeast in water as prepared above was added in the 900L of turbid water. Making up total volume of reactor to 1000 liters.

(1 ml of antibiotics added in 1 L of yeast suspension to avoid bacterial contamination – the effective antibiotics concentration desired is 0.001 % - as standardized in previous experimental work)



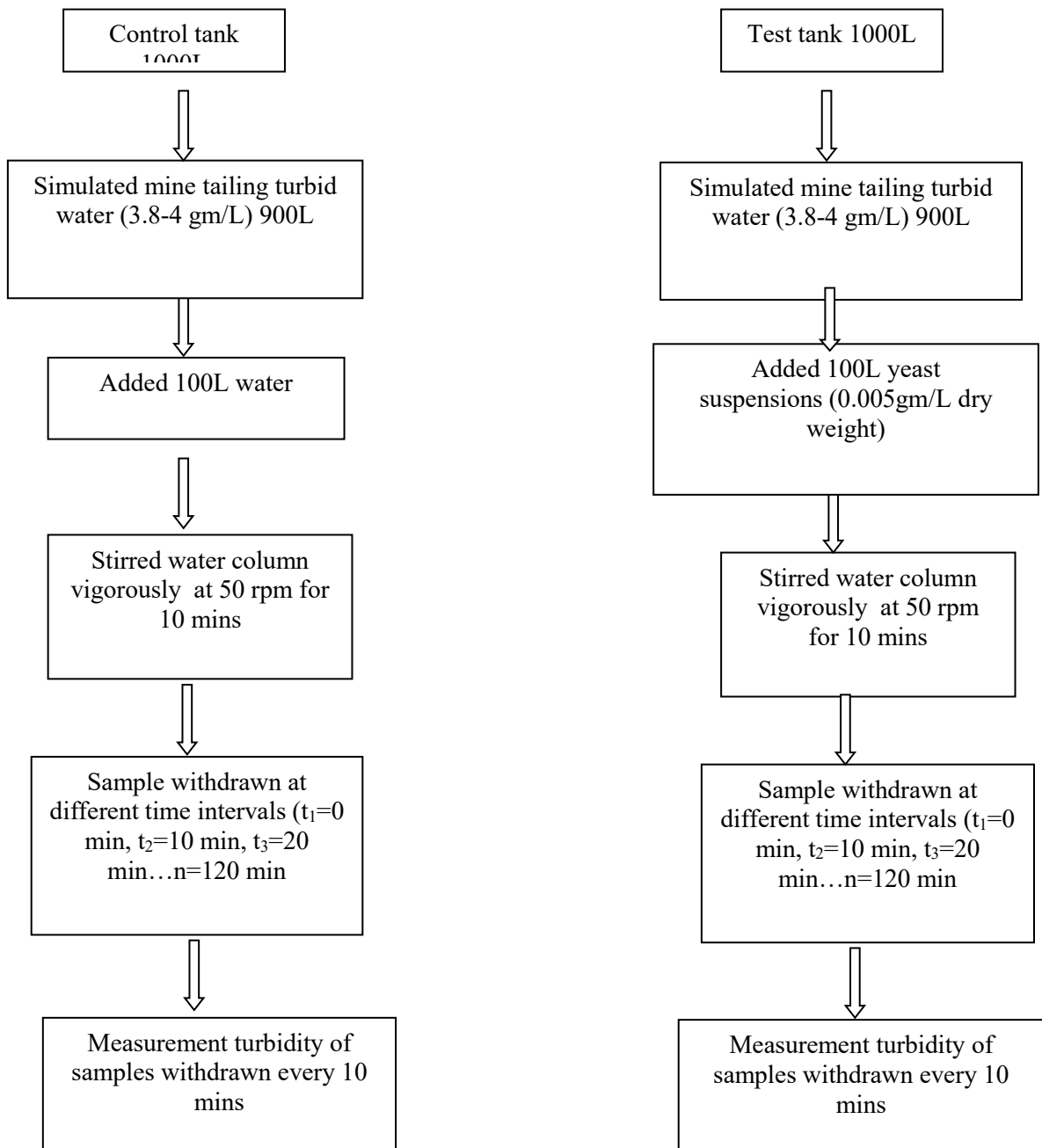
**Figure. 2.26: Scheme for assay preparation**

### **2.9.9. Sand bed treatment (From mid-point of the tank height)**

After 60 min of yeast addition in reactor tank & control tank and stirring, water sample was collected from both the tanks.

Sample from reactor tank was studied for physicochemical parameter of water – sample referred as (A)

Water sample collected from the outlet of sand bed was also studied for physicochemical parameter of water – sample referred as (B).



**Figure.2.27: Biosedimentation test in 1000 L tank**



**Figure. 2.28: Experimental setup for 1000 L test**

### 2.9.10. Sedimentation kinetic modeling aspects involved in the experiment

It is well known that the fundamentals of the rate of aggregation started from the classic work of Smoluchowski (1917). To describe the aggregation rate of particle count and sedimentation based on the Brownian controlled and Stock's law, the general differential equation can be shown below

$$dc/dt = kc^\alpha \dots \text{eq.1}$$

where  $\alpha$  is the order of coagulation process,  $k$  is the coagulation rate constant in (L/mg-min),  $c$  is the total concentration of constituent particles in (m) at time  $t$  (min).

To simplify and solve the Eq. (1), the theoretical values of the order of coagulation or sedimentation process are in the range of ( $1 < \alpha < 2$ ) (Chukwudi et al. 2009; WST 2003). Elimelech et al. (1995) proposed that the aggregation phenomenon followed second order by which the collision is proportional to the product of concentrations of two colliding species. Moreover, in real and empirical practice, extensive studies used ( $\alpha = 2$ ) (Ani et al., 2012) and found that it was more logical in representing primarily the aggregation rate of particles depend on both colloids and coagulants concentrations.

Based on the pervious information substituting ( $\alpha = 2$ ) in the Eq. (1) and integrating it with the following conditions, at initial condition ( $t = 0, c = c_0$ ) and at final condition ( $t = t_c = c$ ) to obtain Eq. (2):

$$1/c = (k_2 t + 1)/c_0 \dots \text{eq.2}$$

Rearranging the eq. (2) which is used to calculate the values of ( $k_2$ ) mathematically in the current research:

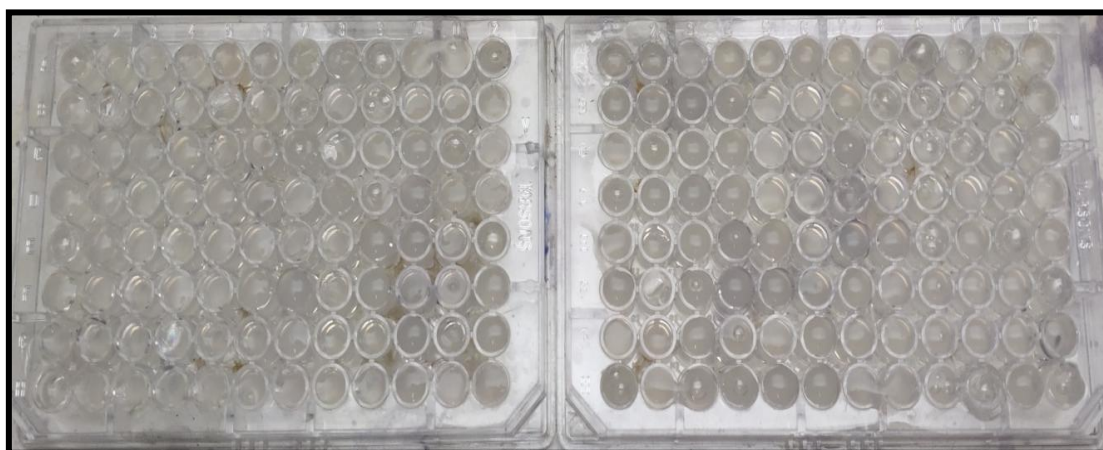
$$k_2 = (1/c - 1/c_0)/t \quad \text{eq.3}$$

Where  $k_2$  is the second order coagulation rate constant. The total concentration suspended constituent particles of three levels of synthetic turbid water as (blank) before treatment ( $c_0$ ) and after treatment ( $c$ ) used in eq. (3) could be expressed in r (m) and the unit of ( $k_2$ ) would be (L/mg-min) (WST 2003; Niam et al. 2007) or the unit of ( $k_2$ ) would be ( $\text{NTU}^{-1} \text{min}^{-1}$ ), (NTU) as turbidity reading.

### 2.10. Preparation of immobilized yeast in agar beads

Yeast suspension with known concentration was pipetted into micro wells of molten agar and allowed to cool before collecting beads. Under a microscope, beads were examined to see if yeast had been trapped. The control beads were made using the same

technique but without the addition of yeast. To increase the frequency of collisions between the beads and clay colloids, beads were placed in turbid water with known NTU and shaken at 100 rpm. After the experiment, the turbidity of the water was measured, and the clay colloids content of the beads was investigated.



**Figure.2.29: Beads preparation in microwells**

**Table.2.9. Yeast immobilization study**

Test	Positive control	Negative control
120 ml turbid water and beads with yeast.(20 ml yeast suspension with known absorbance+180 ml agar beads)	180 ml Turbid water	120 ml turbid water and beads without yeast
pH, 6.4, Tem (°C) 25.4, NTU 780 of SMTTW		

In 9 ml turbid water with known NTU, add 0.5 ml of the most promising yeast suspensions (known absorbance). In separate tubes, add 0.5 ml of each strains with the same absorbance, followed by 0.5 ml of Bchlm-1-2 suspensions. Observed and scored tubes for the removal of clay colloids, as it had been investigated in prior research. The sediment generation of the two most promising combinations was also investigated using the same methods up to a scale of 10-300 ml

### **2.10.1 Turbidity reduction study of natural mine tailing turbid water**

Natural SMTTW from mining area of Goa have been brought to the laboratory and characterized for its different physiochemical properties. Biosedimentation test of clay colloid using yeast cultures have been performed. Further biosedimentation fashions have been compared with mine simulated water

### **2.10.2 Characterization of natural mine tailing turbid water**

Natural mine tailing turbid water was characterize in terms of turbidity and pH.

### **2.10.3 Biosedimentaion test of natural mine tailing turbid water using yeast**

In 900 ml of natural mine water, 100 ml of yeast suspension (0.005 mg/ml, dry weight) was mixed. For 60 seconds, the mixture was vigorously stirred (50 rpm). Water samples were taken from different depths of the Imhoff cone at different intervals of time to measure turbidity with a turbidity meter. Complex sediments generation and quick sedimentation were used to evaluate the biosedimentation efficiency, which was compared to turbidity reduction in a control cone (which yeast were not added, under laboratory condition)



**Figure.2.30: Water body in mining area, turbid due to runoff of the mine tailing soil**

### 2.11.1. Molecular identification of most promising cultures for biosedimentation:

Culture were identified by PCR method, (Triyat Scientific Company (Nagpur) provided this service )

2.11.2. DNA Extraction → PCR → Purification of PCR Product →  
Sequencing → Bioinformatics

### 2.11.3. DNA Extraction

Cells grown in monolayer lysed by suspending 1-3 colonies aseptically and mixed with 450 µl of “B Cube” lysis buffer in a 2 ml micro centrifuge tube and lyse the cells by repeated pipetting.

2. Added 4 µl of RNase A and 250 µl of “B Cube” neutralization buffer.
3. Vortexed the content and incubate the tubes for 30 minutes at 65°C in water bath. To minimize shearing the DNA molecules, mix DNA solutions by inversion.
4. Centrifuged the tubes for 20 minutes at 14,000 rpm at 10 °C.
5. Following centrifugation, transfer the resulting viscous supernatant into a fresh 2 ml micro centrifuge tube without disturbing the pellet.
6. Added 600 µl of “B Cube” binding buffer to the content and mix thoroughly by pipetting and incubate the content at room temperature for 5 minutes.
7. Transferred 600 µl of the contents to a spin column placed in 2 ml collection tube.
8. Centrifuged for 2 minutes at 14,000 rpm and discard flow-through.
9. Reassembled the spin column and the collection tube then transfer the remaining 600 µl of the lysate.
10. Centrifuge for 2 minutes at 14,000 rpm and discard flow-through.
11. Add 500 µL “B Cube” washing buffer I to the spin column. Centrifuge at 14,000 rpm for 2 mins and discard flow-through
12. Reassemble the spin column and add 500 µl “B Cube” washing buffer II and Centrifuge at 14,000 rpm for 2 mins and discard flow-through.
13. Transfer the spin column to a sterile 1.5-ml micro centrifuge tube
14. Add 100 µl of “B Cube” Elution buffer at the middle of spin column. Care should be taken to avoid touch with the filter.
15. Incubate the tubes for 5 minutes at room temperature and Centrifuge at 6000 rpm for 1 min.
16. Repeat the above mentioned step 14 and 15 for complete elution. The buffer in the microcentrifuge tube contains the DNA.



17. DNA concentrations were measured by running aliquots on 1% agarose gel.

18. The DNA samples were stored at -20°C until further use.

#### **2.11.4 Polymerase Chain Reaction**

(PCR) is a method for amplifying certain cloned or genomic DNA sequences using primers and a special enzyme. On a single-stranded DNA template, the enzyme DNA polymerase guides the synthesis of DNA from deoxynucleotide substrates. When a custom-designed oligonucleotide is annealed to a longer template DNA, DNA polymerase inserts nucleotides to the 3' end. When a synthetic oligonucleotide is annealed to a single-stranded template with a region complementary to the oligonucleotide, DNA polymerase can use the oligonucleotide as a primer and elongate its 3' end to form a double-stranded region.

#### **2.11.5. Purification of PCR Production**

Montage PCR Clean up kit was used to remove unincorporated PCR primers and NTPs from PCR products (Millipore). The primers were used to sequence the PCR result. ABI PRISM BigDye™ Terminator Cycle Sequencing Kits with AmpliTaq DNA polymerase (FS enzyme) were used to execute the sequencing procedures (Applied Biosystems).

#### **2.11.6. Sequencing protocol**

Using 26s rRNA universal primers, single-pass sequencing was done on each template. Using an ethanol precipitation technique, the fluorescent-labeled fragments were separated from the unincorporated terminators. The samples were resuspended in distilled water and run through an ABI 3730xl sequencer for electrophoresis (Applied Biosystems).

# **CHAPTER III**

## **RESULTS**

In this chapter results of the performed experiment is presented, as the methodology presented in previous chapter. Freshwater sampling sites and water sample's physiochemical characters are reported. Furthermore, isolated cultures's biochemical characteristics is shown. After screening of the most promising yeast cultures biosedimentation kinetics of simulated mine tailing water using yeast, is presented. Furthermore, results of the experiment performed for the validation of biosedimentation experiment is presented eg.

1. Biosedimentation of natural mine tailing water using yeast culture.
3. Adsorption efficiency and biosedimentation or bioflocculation rate constant.
4. Cultures combination study for efficient turbidity reduction is presented.
5. Study of presence of the microbial in treated and nontreated water is presented.
6. Besides turbidity reduction other water parameter reduction result is presented.

### **3.1. Freshwater habitat from mining and non mining areas of Goa Surveyed**

The physico-chemical parameters of all under study waterbodies depicted in table 3.1 Temperature in selected water habitats from various locations ranged from 24.5 to 30 °C. The pH level of selected sampling sites ranged from 6.3 to 7.8. Netravali water bodies showed high pH due to ongoing activity. Karapur pond and Netravali recorded the highest electrical conductivity 140  $\mu$ s. Each sample's TDS were observed within a permissible level. As per Data on NTU of under study water sample locations ranged between 2 and 15 NTU.

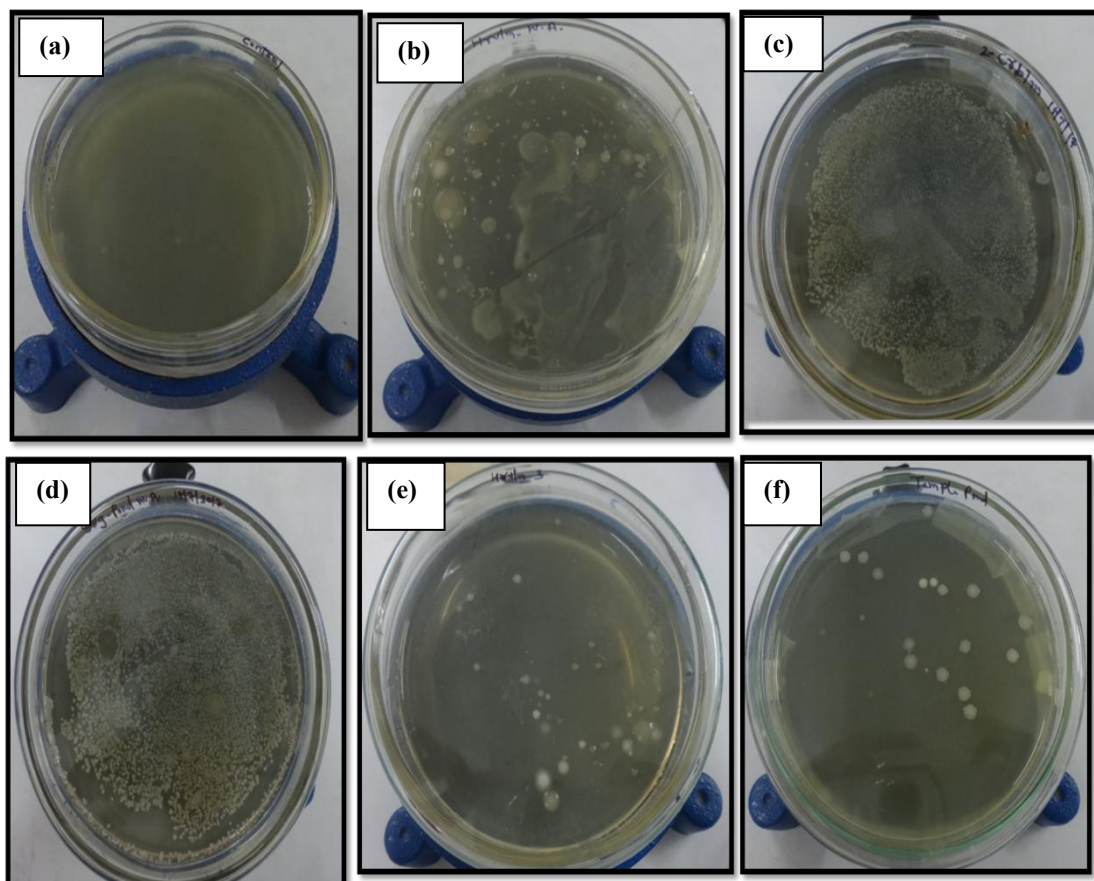
**Table.3.1 Water sampling sites and sample characteristics**

<b>Mining Area</b>					
<b>Name</b>	<b>Temp °C</b>	<b>pH</b>	<b>E.C. <math>\mu</math>s</b>	<b>TDS ppm</b>	<b>Turbidity NTU</b>
Sirigao tank	28.9	6.8	64	20	ND
Mayem lake	29	6.3	44	15	8
Bicholim tank	28	7.33-6.7	90	40	ND
Karapur tank	28	6.87	140	64	ND
Aravelum Water fall's stream	28	7.2	49	27	15
Tambdi surla stream	24.5	6.5	-	-	ND
Bondla stream	25.5	6.4	-	-	ND
Chorla stream	25.5	6.5	50	37	ND
Netravali waterfall's stream	26	7.47	40	15	2.35
Netravali bubbling lake	27	7.8	140	89	8.59

<b>Non Mining Area</b>					
<b>Name</b>	<b>Temp °C</b>	<b>pH</b>	<b>E.C. <math>\mu</math>s</b>	<b>TDS ppm</b>	<b>Turbidity NTU</b>
Safa masjid Pond	30	6.7	103	76	ND
Carambolim lake	29	6.9	128	48	ND
Goa University campus (Talegao plateu)					
Pond-1	25	6.9	102.5	51	6.2
Pond-2	25	6.9	-	-	6.1
Pond-3	25	6.33	92	45	6.2

### 3.2. Isolation of yeasts

A total 40 yeast colonies were isolated from various freshwater bodies of Goa. Control plates, on which water samples were not plated, no microbial growth were observed. On MEA plates without antibiotics, both bacteria and yeast cultures were appeared. On the antibiotic incorporated test plates, only yeast growth was observed.



**Figure.3.1: Representative isolation plates, (a) Control plate, microbial growth was not observed (b) Plate without antibiotics, mix culture observed. (c), Isolation plate carambolim, small tiny colony growth observed (d) isolation plate ,Sirigao, small tiny colony growth observed. Isolation plate of (e) Aravelum, two large dissimilar colonies observed (f) Isolation plate of temple pond, two dissimilar morphological colonies were observed.**

### 3.2.2. Colony Characteristics on the isolation plates

On a regular basis, isolation plates were observed for the microbial colony growth, and the following microbial growth pattern characteristics were observed (Table3.2). The colony of the some isolate had a waxy texture, some of them were shiny and smooth. The isolation plates have showed mixture of flat and elevated, elevations of the colonies. The colonies were white, beige, and cream colour in nature. Single colony showed pink colour Pnd-2-2. Moreover, plates without antibiotics showed mixed growth of microbial colonies.

**Table.3.2: Growth rate and colony morphology of cultures**

Strain Designation	Colour	Colony shape	Surface	Elevation	Texture
Bchlm-1	White	Circular	Shiny	Flat	Smooth
Bchlm-1-2	White	Circular	Shiny	Flat	Smooth
Bndl-2-2	Beige	Granular	opaque	Raised	Smooth
Bndl-1-1-1	Creamish	Circular	Shiny	Flat	Smooth
Bh-1	White	Circular	Shiny	Flat	Smooth
Bnd-1	Creamish	Circular	Shiny	Flat	Rough
Bndl-3	Creamish	Circular	Shiny	Flat	Smooth
Bndl-2-1	Creamish	Circular	Shiny	Flat	Smooth
Chl-1	Beige	Granular	opaque	Raised	Smooth
Crmbm-1-2	White	Circular	Shiny	Raised	Smooth
Gh1-1	White	Circular	Shiny	Flate	Rough
Gh2-1	White	Circular	Shiny	Raised	Rough
Gh1-3	White	Circular	Shiny	Raised	Rough
Gh-4	White	Circular	Shiny	Raised	Rough
Hrvml-2	White	Circular	Shiny	Flate	Smooth
Krpr-1	White	Circular	Shiny	Flate	Smooth
Krpr-2	White	Circular	Shiny	Flate	Smooth
Mym-1-2	Beige	Circular	Shiny	Flate	Smooth
Ntrvl-1-1-1	Beige	Circular	Waxy	Raised	Smooth
Ntrvl-1-2	White	Circular	Shiny	Raised	Smooth
Ntrvl-2-2	Beige	Circular	Waxy	Raised	Smooth
Ntrvl-2-3	White	Circular	Shiny	Raised	Smooth
Pnd-2	Pinkish	Granular	Shiny	Flat	Smooth
Pnd-3	Beige	Circular	Shiny	Raised	Smooth
Pndsm-1	White	Circular	Shiny	Flat	Smooth

Strain Designation	Colour	Colony shape	Surface	Elevation	Texture
Pndsm-1-2	Beige	Circular	Shiny	Flat	Smooth
Pndsm-2-2	Pinkish	Granular	Shiny	Flat	Smooth
Pndsm-3	White	Circular	Opaque	Flat	Rough
Pnd-2-2	White	Circular	Shiny	Raised	Smooth
Pnd-3	Beige	Circular	Shiny	Raised	Smooth
Sc-1	White	Circular	Shiny	Raised	Smooth
Rw.a	White	Circular	Shiny	Flat	Smooth
Srg-3-3	White	Circular	Shiny	Flat	Smooth
Srg-1	White	Circular	Shiny	Flat	Smooth
Srg-2-1	Cream	Grannular	Shiny	Flat	Smooth
Tmrs-1-2-4	White	Circular	Shiny	Flat	Smooth
Tmrs-2-2	Beige	Circular	Waxy	Flat	Smooth
Tmrs-2-2-4	Beige	Grannular	opaque	Raised	Smooth
Tmrs-2-3	Beige	Circular	Waxy	Flat	Smooth

### 3.2.3. Micromorphology of the cultures

Cultures were studied under the microscope; mix culture were grown on the plates which were not incorporated with antibiotics, bacterial cells and yeast cells were associated as typical bacilli and yeast. Most important point can be concluded is all yeast cultures are antibiotics resistance as they have grown in antibiotic incorporated media. Pure cultures have a variety of morphologies, as illustrated in the morphology of the cultures (Fig 3.2). Some of the cultures are spherical, few of them showed oval cell shape. Some of culture were in cylindrical shape and have pseudomycelium. Dense colony cells were compactly bounded to each other and strongly stained with crystal violet in the strain designated as Ntrvl-2-1, Pnd-3, and Ntrvl-1-1-1. The cells of the cultures Ntrvl-1-1, Ntrvl 2-3, showed scattered single cells and stained strongly with crystal violet. Tmrs-2-2-4 showed polymorphism as some cells were elongated, oval and spindle in shape, when the culture's smear stained in the crystal violet stain. Moreover, most of the culture's cells with different morphology formed pseudomycelium. The Pnd-2-2 cells were lightly stained in the crystal violet stain and showed small free cells. Chl-1 cells were small in size and highly stained in crystal violet. Some of the cultures showed single- budding independent cells eg. Bchlml-2, Ntrvl-2-2. Short branched chains were observed in many colonies eg. Gh-1-3 and long chains were observed in Ch-1 and Bnd-1-1-1.

### 3.2.3. Morphology of cultures in monochrome staining

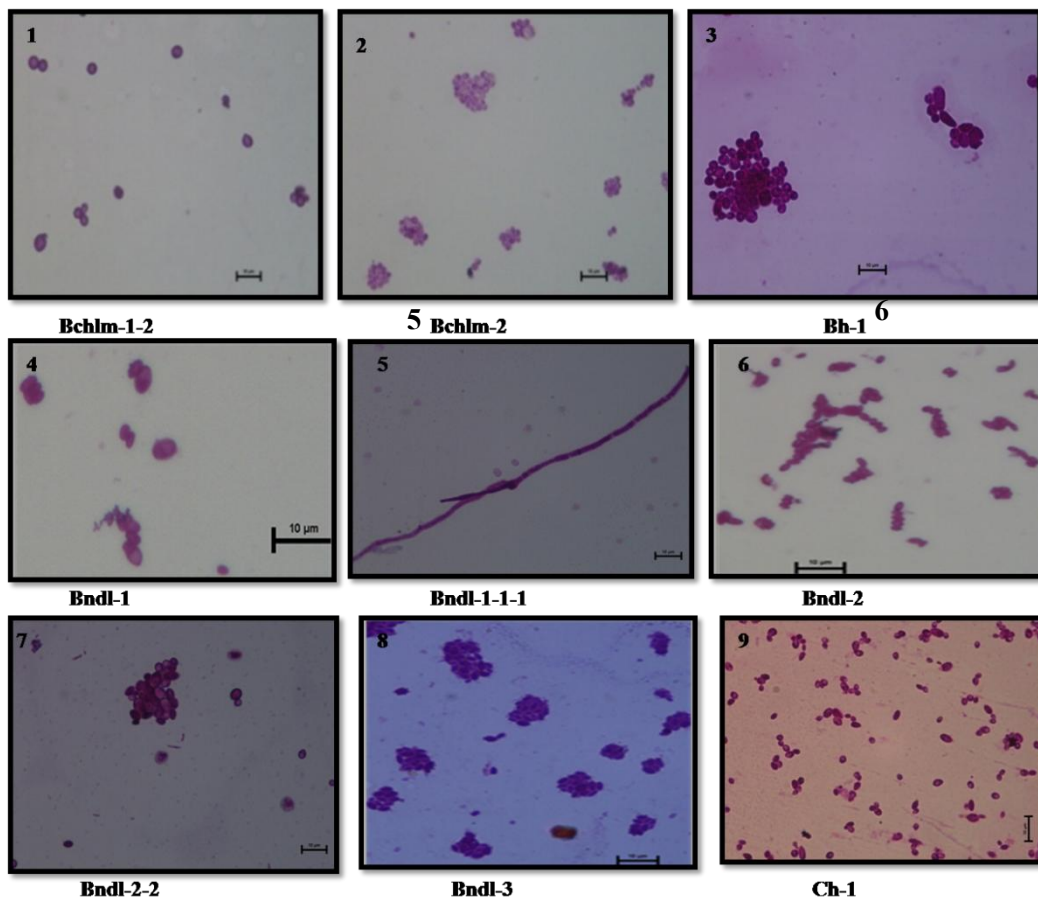


Figure.3.2: (1-9) Monochrome stained cells of yeasts



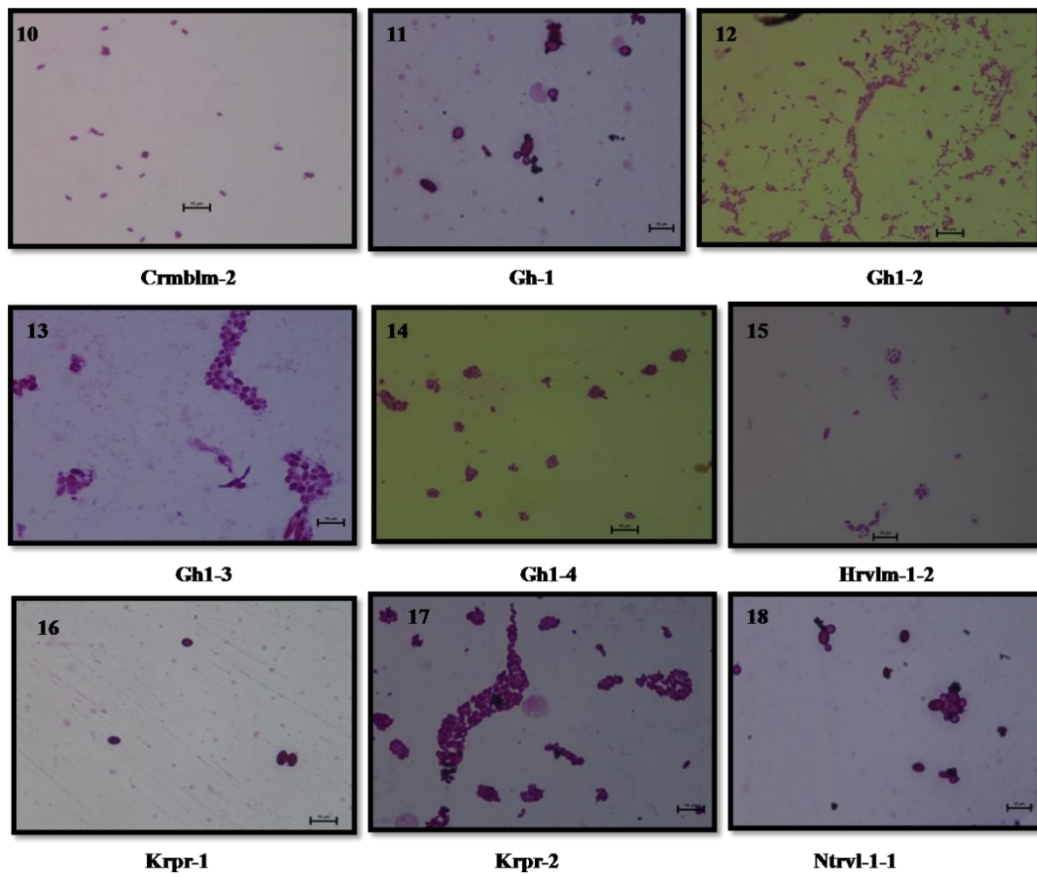


Figure.3.3: (10-18) Monochrome stained cells of yeasts

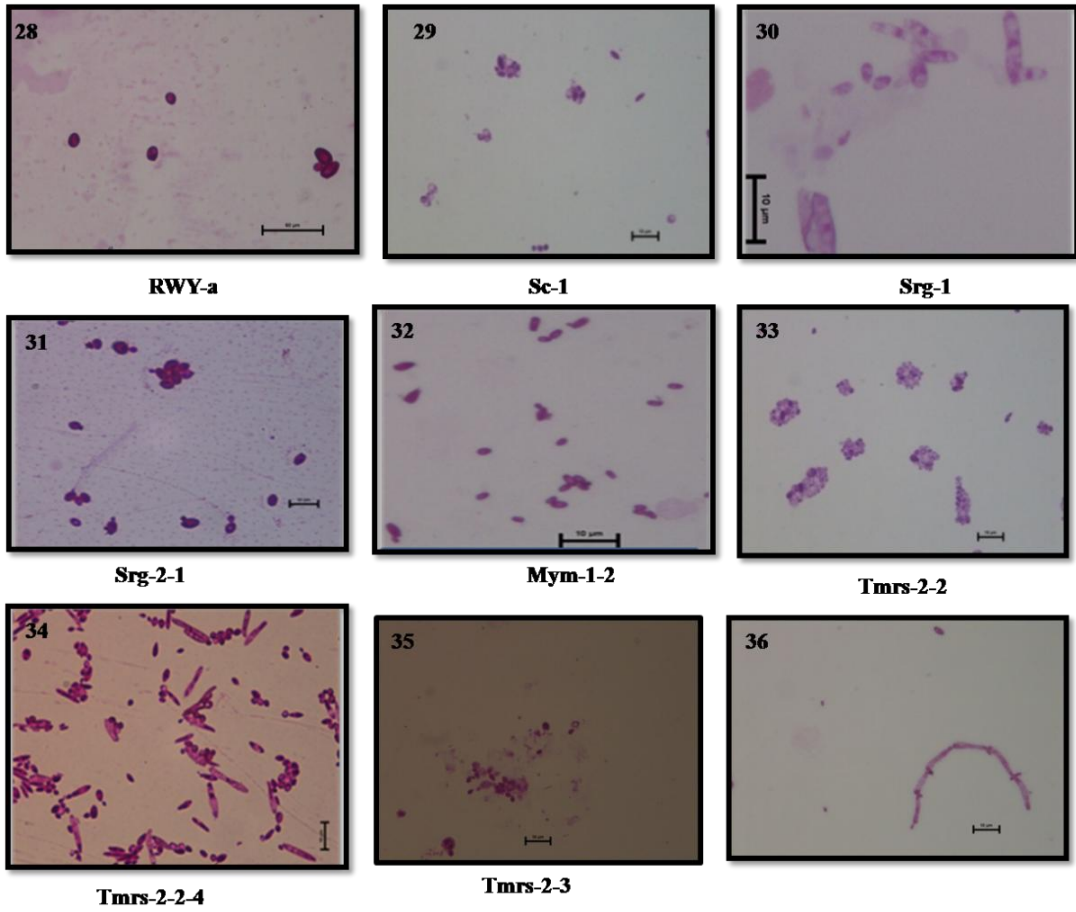


Figure.3.4: (28-36) Monochrome stained cells of yeasts

### 3.2. 4. Biochemical tests

Sugar fermentation tests of the cultures revealed a change in the medium colour from blue to purple pink, yellow, indicating the formation of acid (fig3.5). The reaction with sugar of cultures is showed in the table 3.3. Most of the strain showed positive test with glucose except Bchl1m-1, Crmb1m-2, Chrl-1, Hrv1m1-2, krpr-2, Mym-1-2. Ntrvl-2-3, Pnd-3. On the other hand Bchl1m-1, pnd-3, Pndsm-1, and Tmrs-2-3, did not ferment sugars at any stage. Bndl-1-1-1, Gh-4, showed positive test with glucose only. Some of the strain showed negative test with lactose and some of the strain showed negative result with maltose.

Yeast cells showed various sizes and shapes. Pseudomycilium and ascospores were examined on corn meal agar media and sodium acetate agar (Fig.3.10). The colour of the colonies and texture remained same on corn meal agar as on the isolation plate with MEA media and after streaking on the slant.

Bndl-1-1-1, Bndl-1-1, krpr-2, GH<sub>1</sub>-5, Pnd-2, Pndsm-1, Mym-1-2, Ntrvl-1-1-1 did not displayed the presence of mycelium in their colony after grown on specified media.

Moreover, after growing on Nitrate agar, strain designated as, Bchl1m-1-2, BH-1, Bndl-1-1-1, Crmb1m-2, Gh1-4, Hrv1m-1-2, Mym-1-2, Ntrvl-1-2, Ntrvl-2-3, Pndsm-1-2, showed positive growth.

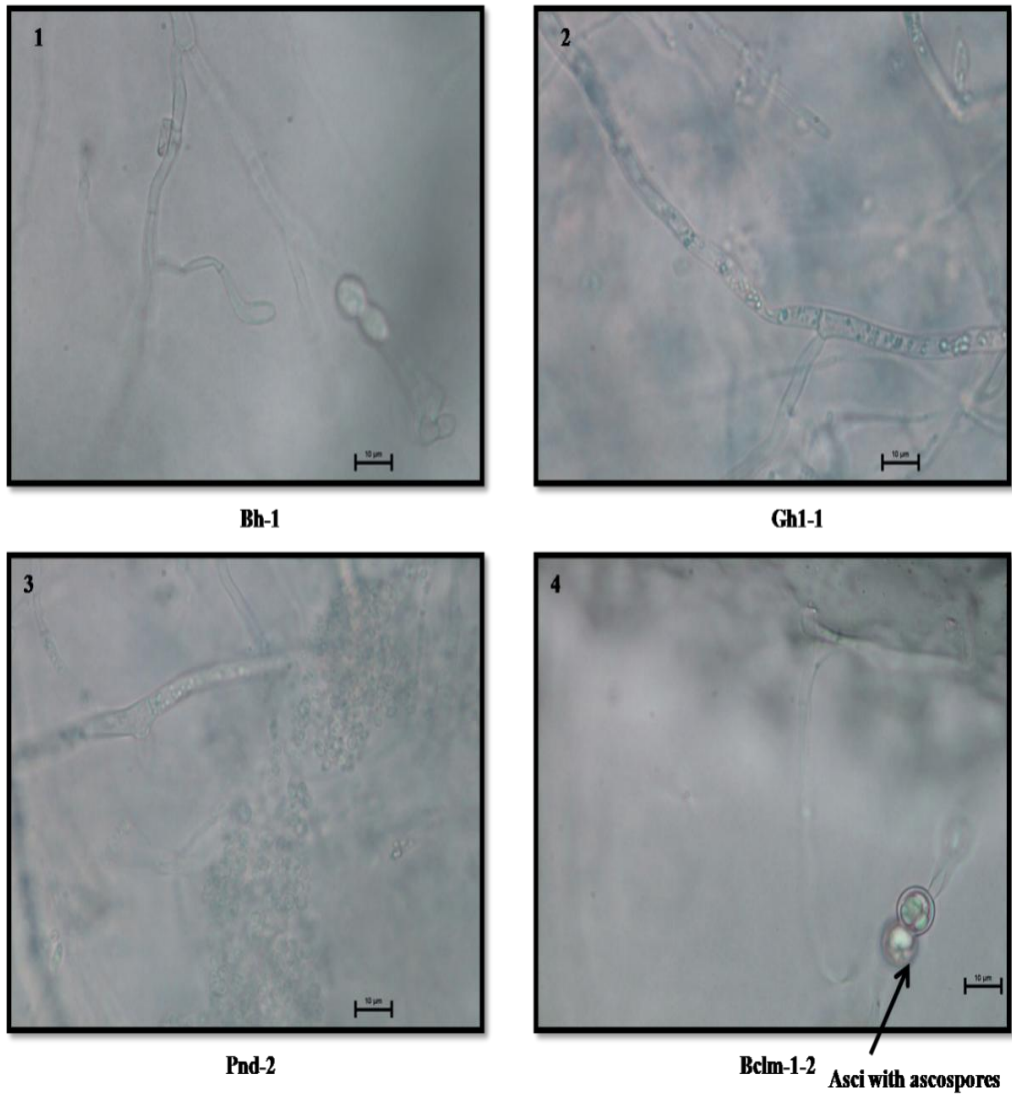
On the sodium acetate agar Bchl1m-1-2, BH<sub>1</sub>, Bndl-1, Bndl-1-1-1, Chr-1, Hrv1m-1-2, Mym-1-2, Ntrvl-1-1-1, Ntrvl-1-2, Ntrvl-2-2, Pnd-2, Pnd-3, Pndsm-1, Pndsm-1-2, Pndsm-2-2, Srg-1 showed positive result (Table. 3.4).

Culture streaked on the *Candida* differential agar media were showed different colour of the colony (Table. 3.4). Many strains did not grow on differential agar media can be predicted as belongs to saccharomyces group.

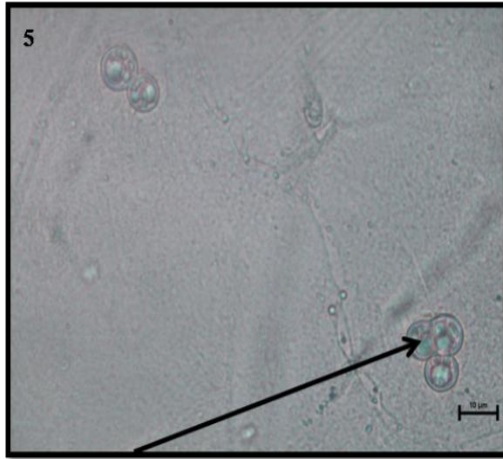
On the basis of colour development in the colonies on *Candida* differential agar, and using morphological features and biochemical tests, strains were tentatively identified, as (Table.3.5).



**Figure.3.5: Sugar fermentation test**



**Figure.3.6: (1-4) Various form of ascospores**



**Persistent,  
unconjugated asci with  
globose ascospores**

**Ntrv1-1-2**

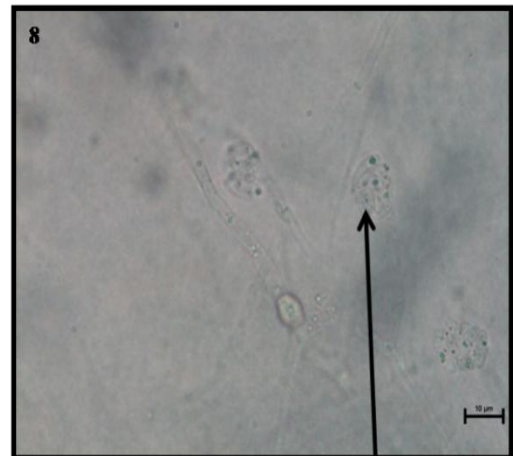


**Bh-1**



**Asci with globose  
ascospores**

**Srg-2-1**



**Bndl-1**

**Asci with ascospores**

**Figure.3.7: (5-8) Various form of ascospores**

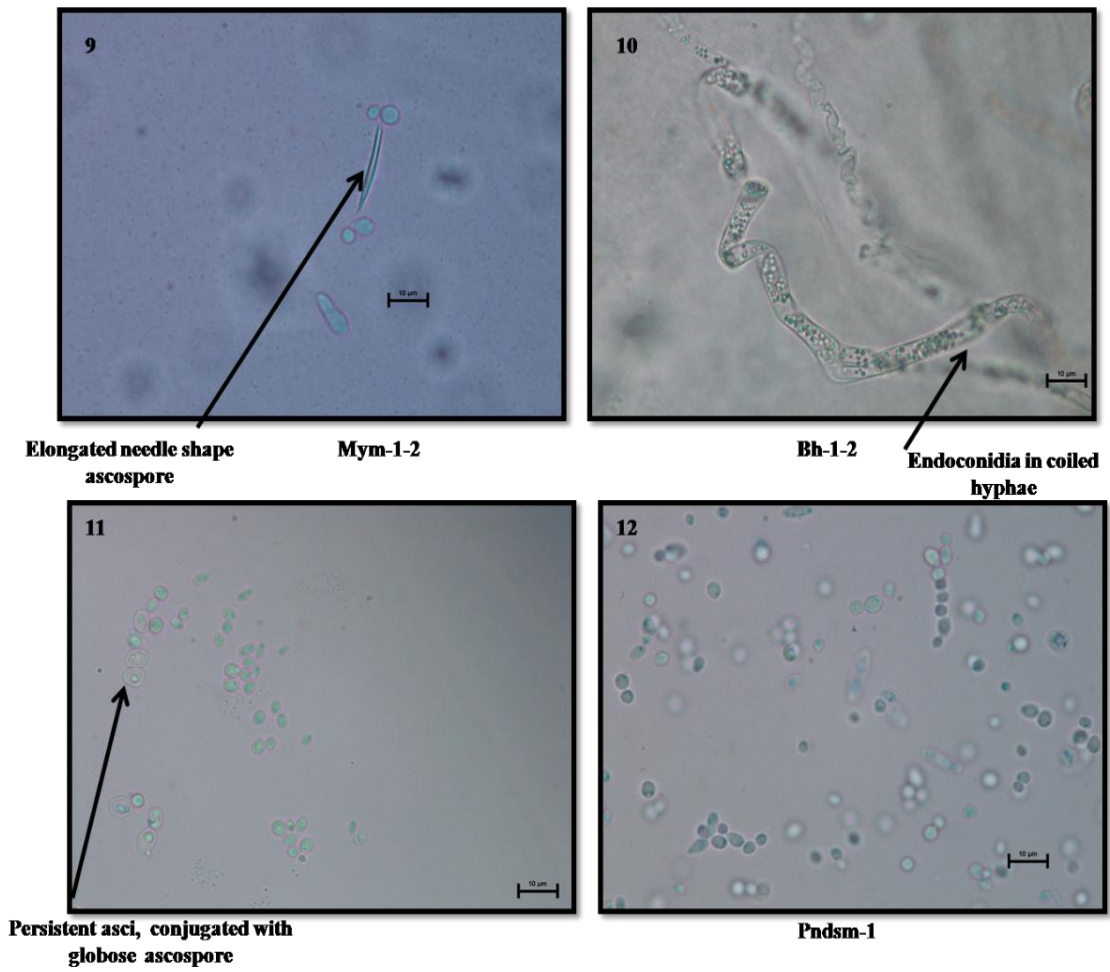
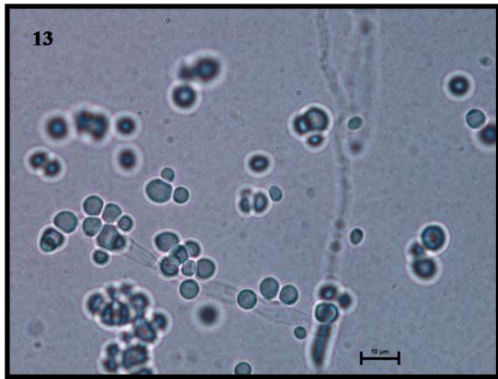
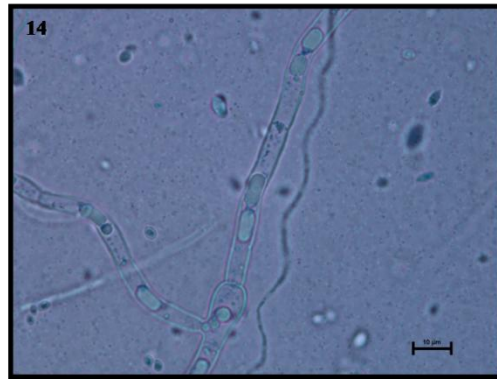


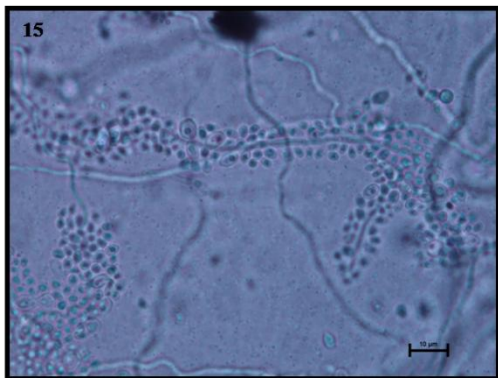
Figure.3.8: (9-12) Various form of ascospores



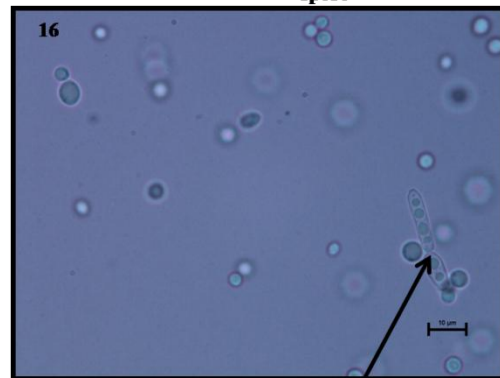
**Ntrvl-1-2**



**Gh2-1 Aseptate hyphae with spore**



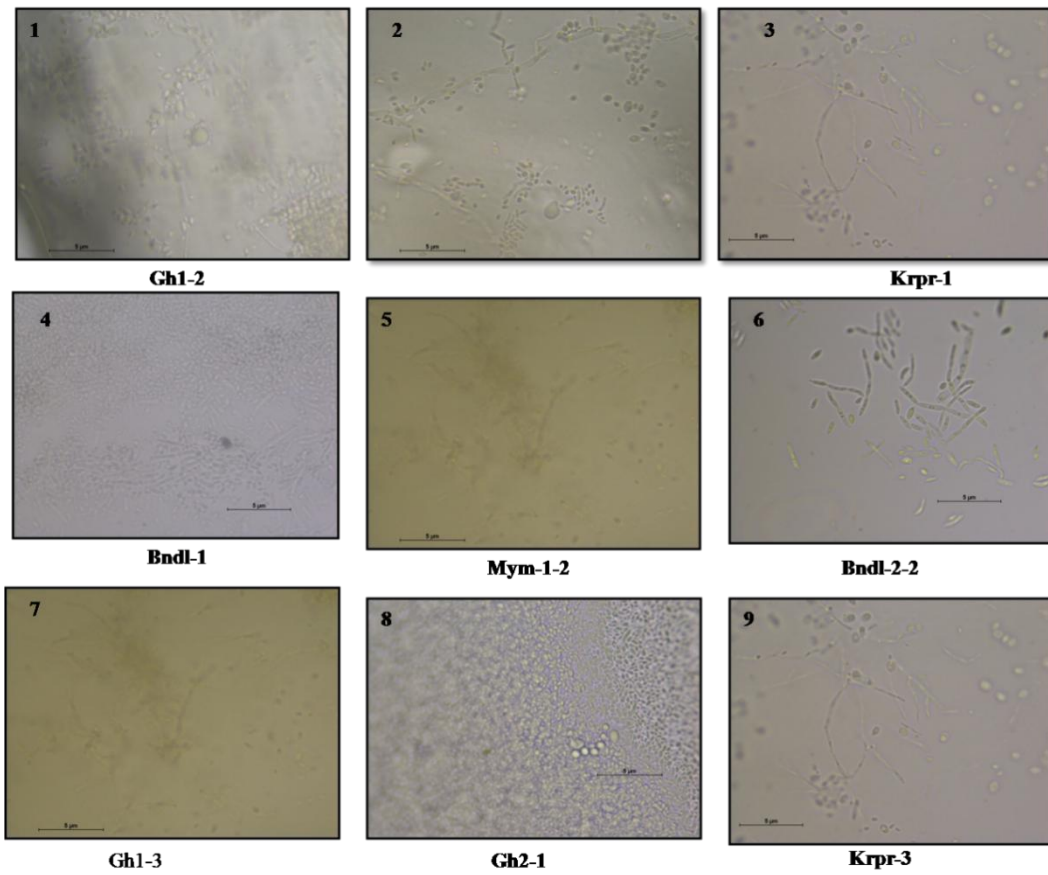
**Gh-1**



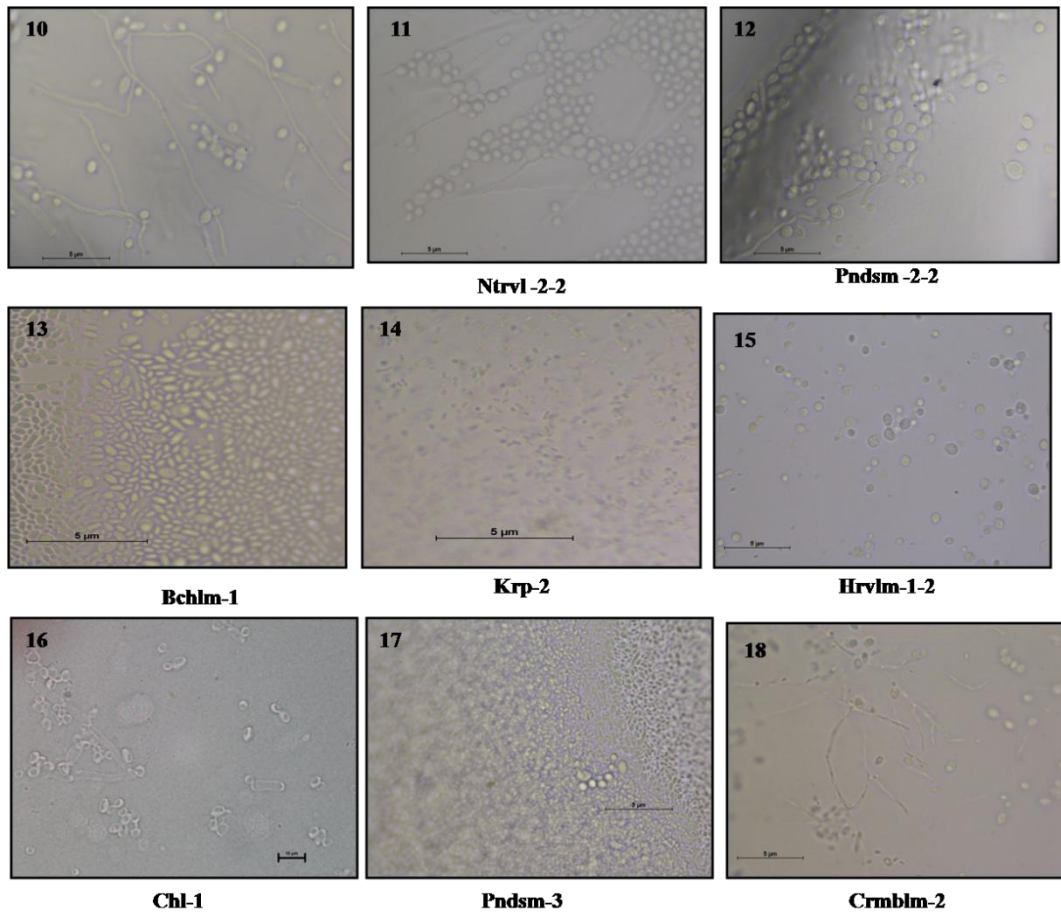
**Hrvlm-1-2 Globose ascospores**

**Figure.3.9: (13-16) Various form of ascospores**





**Figure.3.10: (1-9) Various form of pseudomycelium and non mycelia cells**



**Figure.3.11: (10-18) Various form of pseudomycelium and non mycelia cells**

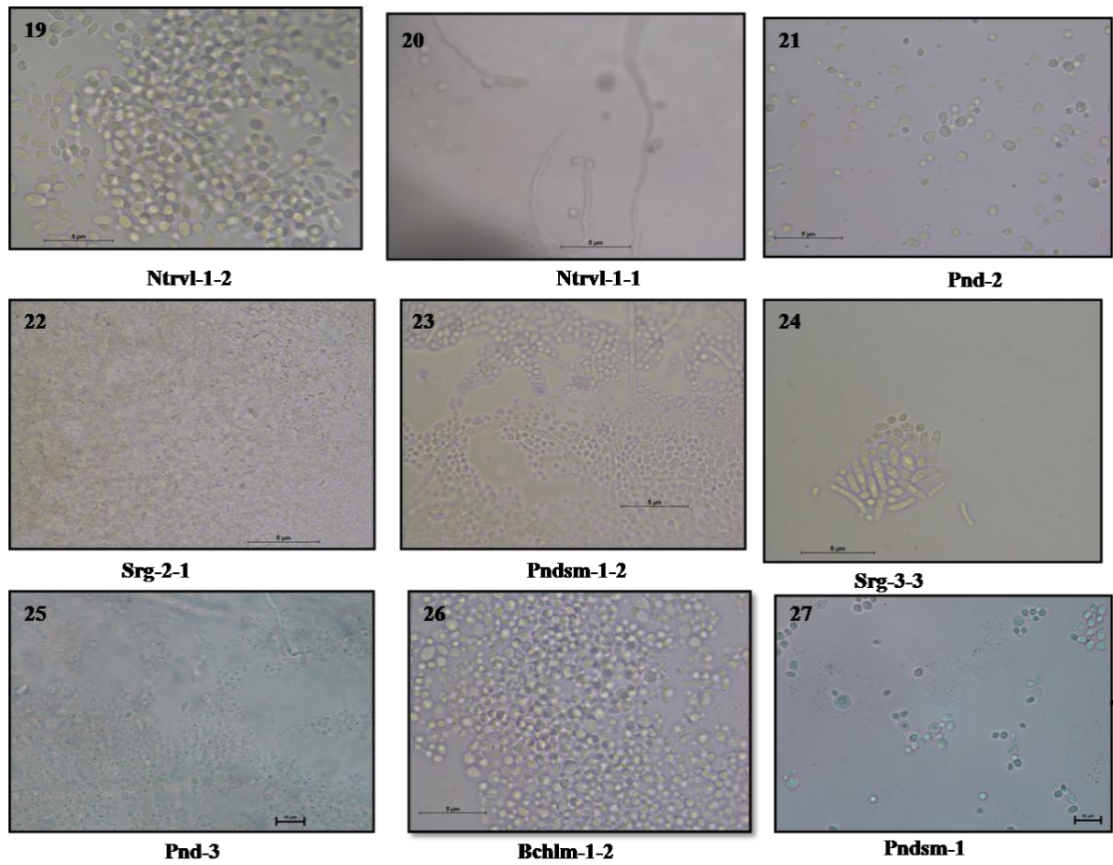


Figure.3.12 (19-27) Various form of pseudomycelium and non mycelia cells

**Table.3.3 Yeast's colony characteristics in various mediums**

Strain designation	Cielum production	Nitrate agar	Sodium acetate agar	Sugar test			
				Glucose	Sucrose	Lactose	Maltose
Bchlm-1-2	-	+	+	+	+	-	+
Bchlm-1	-	-	-	-	-	-	-
BH <sub>1</sub>	+	+	+	++	-	+	+
Bndl-1	+	-	+	+	+	+	-
Bndl-1-1-1	-	+	+	+	+	+	-
Bndl1-1-1	-	-	-	+	-	-	-
Bndl-2-2	-	-	-	+	-	-	-
Crmblm-2	+	+	-	-	NA	-	+
Chr1-1	+	+	+	-	+	+	+
Gh1-3	+	-	-	+	-	-	+

Gh1-4	+	+	-	+	-	-	-
Gh1-5	-	-	-	+	-	-	-
Hrvlm1-2	-	+	+	-	-	-	-
Krpp-2	-	-	-	-	-	+	+
Mym-1-2	-	+	+	-	+	+	++
Ntrvl	-	-	-	+	-	-	+
Ntrvl-1-1-1	-	+	+	+	+	+	++
Ntrvl-1-2	+	+	+	+	+	-	++
Ntrvl-2-2	+	-	+	+	+	+	+
Ntrvl 2-3	+	+	-	-	-	-	-
Pnd-2	-	-	+	+	-	-	-
Pnd-3	+	-	+	-	-	-	-

Pndsm-1	-	-	+	-	-		-
Pndsm-1-2	+	+	+	+	+	-	++
Pndsm-2-2	+	-	+	+	-	+	+
Pndsm-3	-	-	.	+	-	-	-
RW(2a)	+	+	+	+	+	-	+
Srg-1	+	+	+	-	+	+	+
Srg-2-1	+	-	.	+	+	-	-
Tmrs-2-2	-	-	-	+	+	-	+
Tmrs-2-3	-	-	-	-	-		-
Tmrs-2-2-4	+	+	Not grown	-	-	-	-

**Table.3.4 Colony characteristics on differential agar media ((The colour coding indicates colour observe in plates)**

Strain Designation	Colour in D.A.	Colour in MEA	Strain Designation	Colour in D.A.	Colour in MEA
Bchlm-1-2	Blue	White	Ntrvl-2-2	Green	White
Bchlm-1	Not grown	Beige	Pndsm-1	Not grown	White
Bndl-2-2	Brown	White	Pndsm 1-2	Shiny brown	White
Bndl-1-1-1	Not grown	White	Pndsm-2-3	Blue	White
Bh-1	Purple	Cream	Pndsm-3	Brown	Cream
Bnd-1	Pink	White	Pnd-2	Coke	White
Bndl-3	Purple	White	Pndsm-2-2	Not grown	White
Bndl-2-1	Purple	White	Pnd-2-2	Dark Purple	Red
Chl-1	Not grown	Beige	Pnd-3	White	White
Crmlm-1-2	Not grown	White	Sc-1	Not grown	-
Gh-4	Peach Pink	White	Rw.	Coke	Beige
Gh2-1	Coke	White	Srg-3-3	Pink	White
Gh1-3	Not grown	Beige	Srg-1	White	Beige
Hrvlm-1-2	Pink	White	Srg-2-1	Not grown	White
Krpr-1	White	White	Bchlm-2	Green	White
Krpr-2	Not grown	White	Tmrs-2-1-2	Not grown	White
Mym-1-2	Green	Brown	Tmrs-2-2-4	White	White
Ntrvl-1-2	Green	White	Tmrs-2-3	Not grown	White
Ntrvl-1-1-1	Blue	White	-	-	-
Ntrvl-2-1	Not Grown	White	-	-	-

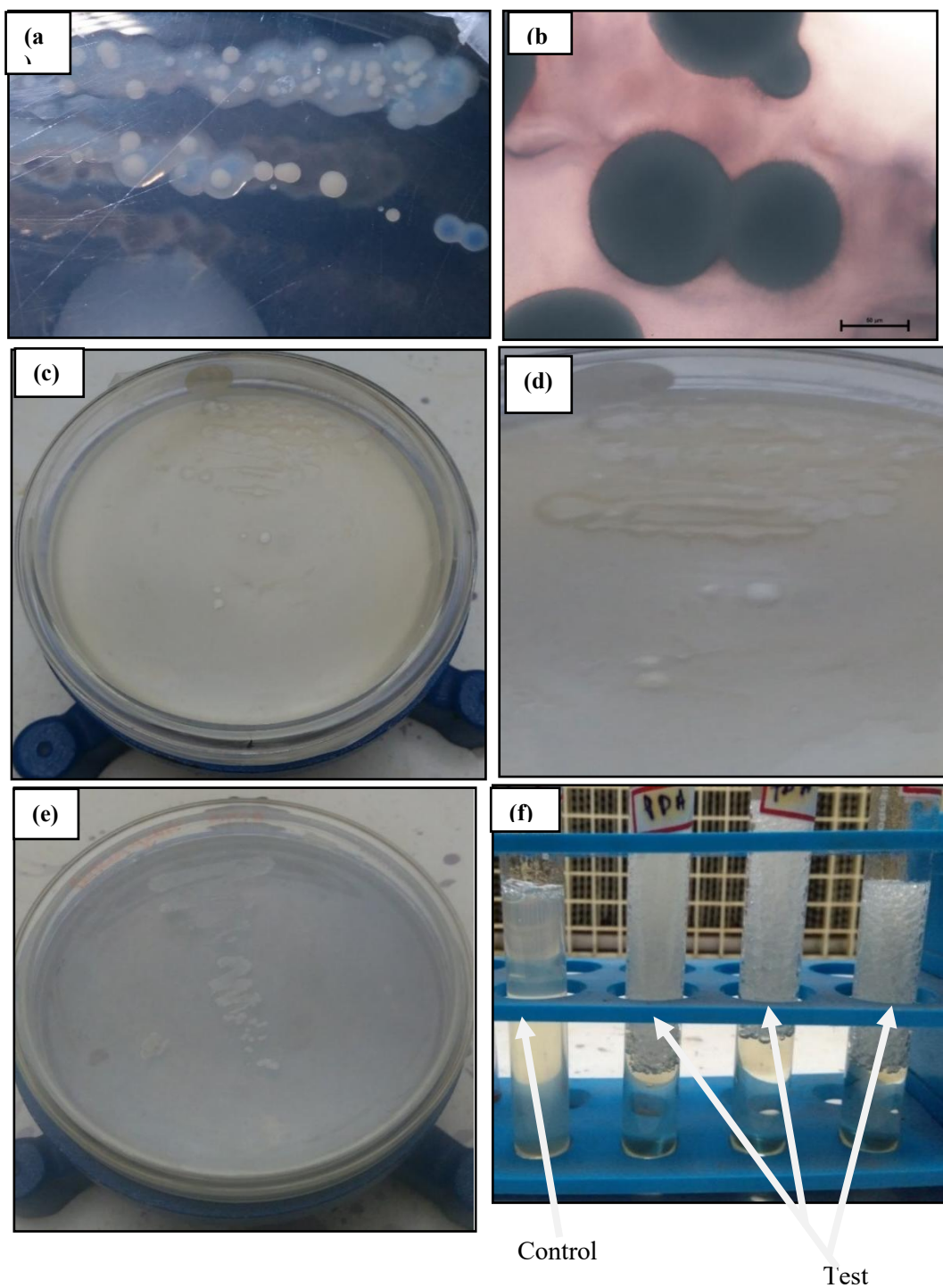
**Table.3.5 Tentative identified cultures**

<b>S. No.</b>	<b>Strain designation</b>	<b>Genus</b>
1	Bchlm-1-2	<i>Candida sp. I</i>
2	Bnd-1	<i>Candida sp. II</i>
3	Srg-3-3	<i>Candia sp. III</i>
4	Ntrvl 1-2	<i>Candida sp. IV</i>
5	Pnd-2	<i>Candida sp. V</i>
6	Ntrvl-1-1-1	<i>Candida sp. VI</i>
7	Pndsm-3	<i>Candida sp. VII</i>
8	Pndsm-2-3	<i>Candida sp. VIII</i>
9	Gh-4	<i>Candida sp. IX</i>
10	Bndl-3	<i>Candida sp. X</i>
11	Bndl-2-2	<i>Candida sp. XI</i>
12	Pnd-2-2	<i>Rhodotorula sp.</i>
13	Pnd-3	<i>Sacchromycopsis sp.</i>

### **3.2.5.1 Characterization of the most promising yeast**

The dye adsorption capability of the most promising yeast (for biosedimentation) was high (fig.3.13 (b)) whole colony turned blue on the media and zone of clearance observed. The occurrence of salt deposition on the plate can be attributed to calcium oxlate (fig 3.13 (c, d)). With catalase, the most promising strain produced a large gas bubble (fig.3.13 (f)) provides insight of hydrogen peroxide reaction





**Figure.3.13: Yeast isolate Bchlm-1-2 (promising for biosedimentation)**  
**(a) showing strong dye absorption (b) colony turned blue (c) acid**  
**(b) crystals (d). (f) Strain showed huge gas bubble formation with H<sub>2</sub>O**

### 3.2.5.2 Phosphate solubilization efficiency

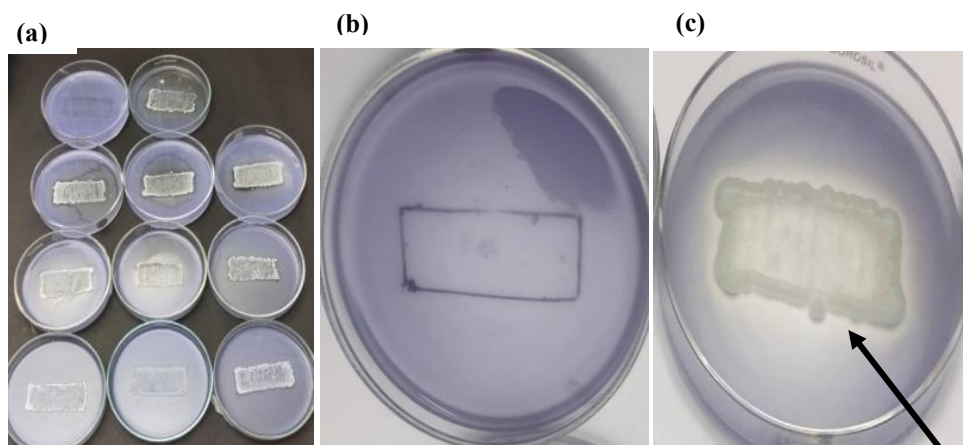


Figure.3.14: (a) Phosphate solubilization plate assay. (b) Control plate inoculated with BchlM1-2 strain showed clear zone around the inoculum. (c) Zone of clearance

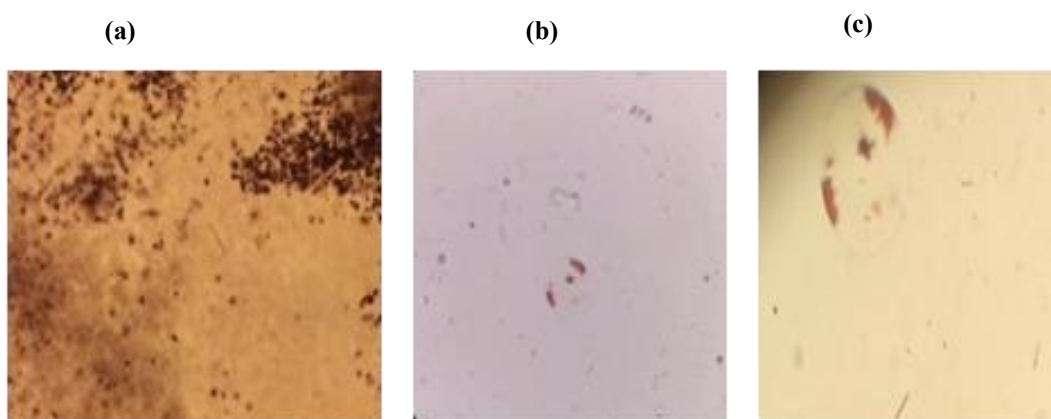
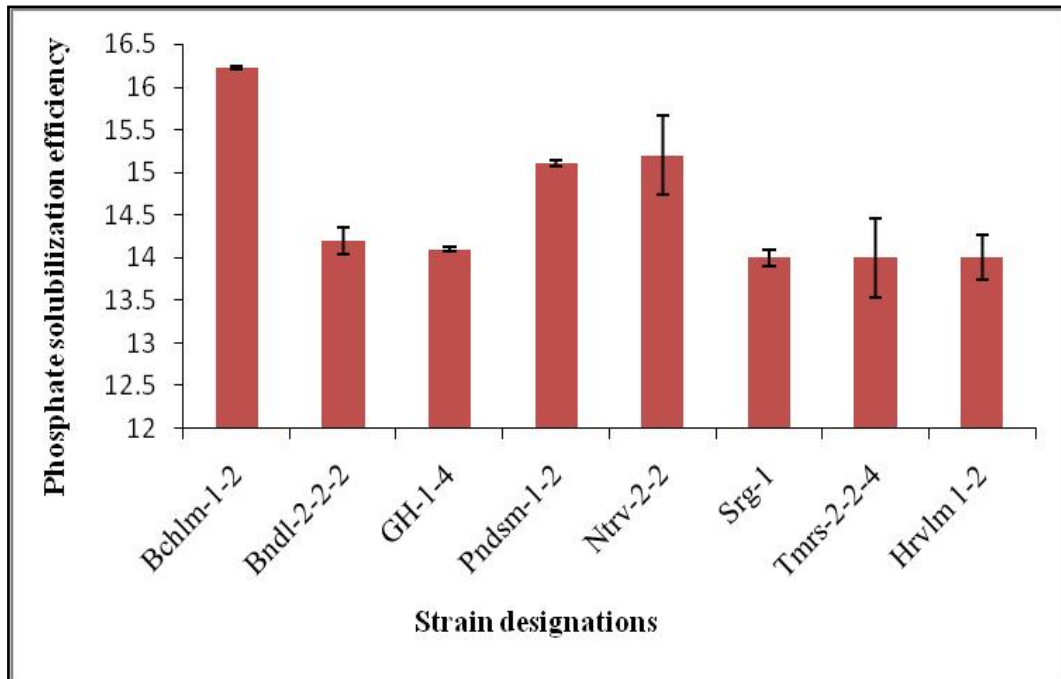


Figure.3.15: Inoculation plate for phosphate solubilization with crystal of calcium oxalate



**Figure.3.16: Phosphate solubilization efficiency**

Five strains with high scores were discovered. Among all studied strains, the Bchlm1-2 strain received a perfect score. Furthermore, the 150 mm<sup>2</sup> area surrounding each implanted rectangular biofilm was more transparent than the rest of the petriplate, owing to the high oxalic acid concentration along the boundary. The chemogradient of oxalic acid decreases as one moves away from the biofilm border.

The Pnd-3 strain scored poorly in terms of phosphate solubilization. The quantification provides a more consistent metric for comparing and selecting multiple strains.

### 3.2.5.3 Surface charge test

According to the results of surface charge test it is found that test yeast has a significant capacity to bind cations on its surface. In comparison to commercial baker yeast, test yeast showed a strong affinity with alcian blue dye.

**Table.3.6: Absorbance of test yeast, commercially available yeast, and dye**

	Absorbance			Mean
Yeast+Alcianblue dye Supernatant	0.002	0.002	0.002	0.002
Sachromyces cereveciae+Alcian blue dye supernatant	0.004	0.004	0.004	0.004
Alcian blue dye	0.009	0.009	0.009	0.009

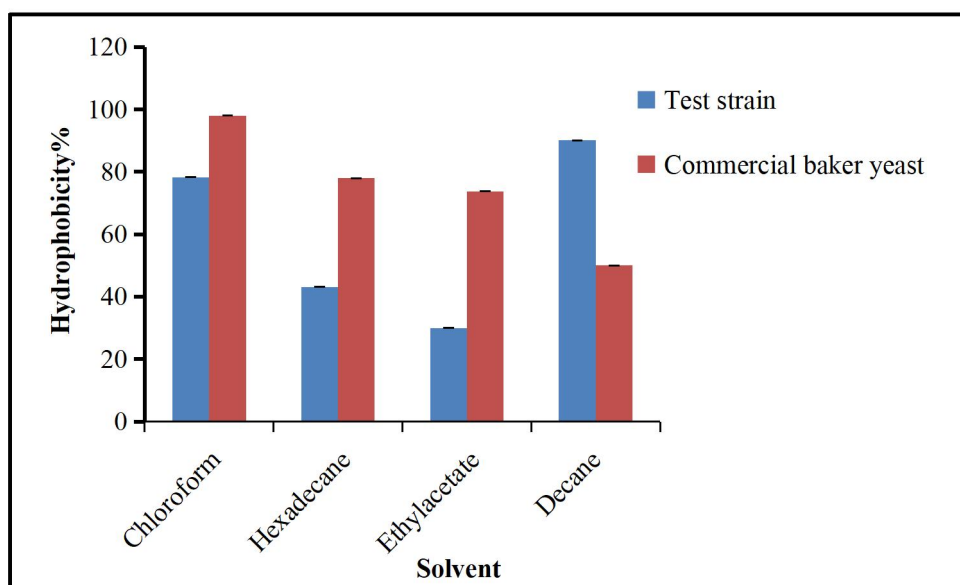
$$\text{ABR} = (\text{AABsolution} - \text{Asupernatant}) \times 100 / \text{AABsolution}$$

$$\begin{aligned} \text{ABR of test yeast} &= \frac{(0.009 - 0.002) \times 100}{0.009} \\ &= 77.77 \end{aligned}$$

$$\begin{aligned} \text{ABR of Baker yeast} &= \frac{(0.009 - 0.004) \times 100}{0.009} \\ &= 55.55 \end{aligned}$$

### 3.2.5.4 Hydrophobicity test

Baker yeast had a greater affinity with hexadecane, ethyl acetate, and chloroform. Moreover, the test strain's affinity for decane proved to be strong, it might be as culture was treated in buffer. Standard deviations of all replicates were negligible.



**Figure.3.17: Hydrophobicity percentage of yeasts**

### 3.2.5.5 Test for morphological stability of selected culture

The test culture turned black in the modified media, (prepared with mine tailing soil and agar), possibly due to metal adsorption, but it turned white again when restreaked on the nutrient-rich (MEA 2%) media. After seven days, colonies of both plate, MEA media and modified mine tailing soil medium turned slimy to protect themselves from infection and starvation. As shown in fig.3.18.

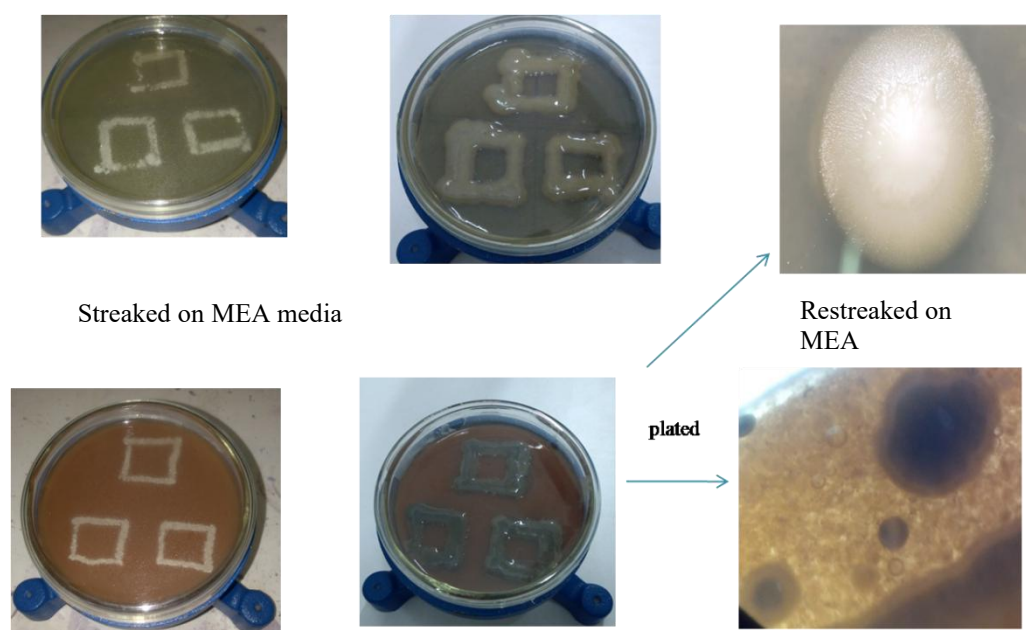


Figure.3.18: Bchl1-2 strain grown on MEA and modified media

### 3.3. Biosedimentaion efficiency test

The bioefficiency scores of selected strains is presented here as discussed in chapter two (table 2.25). Highest score is obtained by strain Bclm-1-2.

**Table 3.7 Biosedimentation efficiency test score**

<b>Strain designation</b>	<b>score</b>
Bchlm-1-2	280
GH-1-1	105
Pnd-2	130
Pnd-2	130
Pnd-3	105
Pndsm-2-2	130
Tmrs-2-2-4	160

### **3.3.1 Fractality Index of selected strains**

The results of Fractality Index of the colony margin (preliminary study) is presented in Fig.3.21, Further Fractality Index of best five strains, studied after slightly modification in method as showed in fig. 3.23 is presented. Strain designated as Bchlm-1-2 showed highest Fractality Index of colony margin (Fig 3.21;3.23)

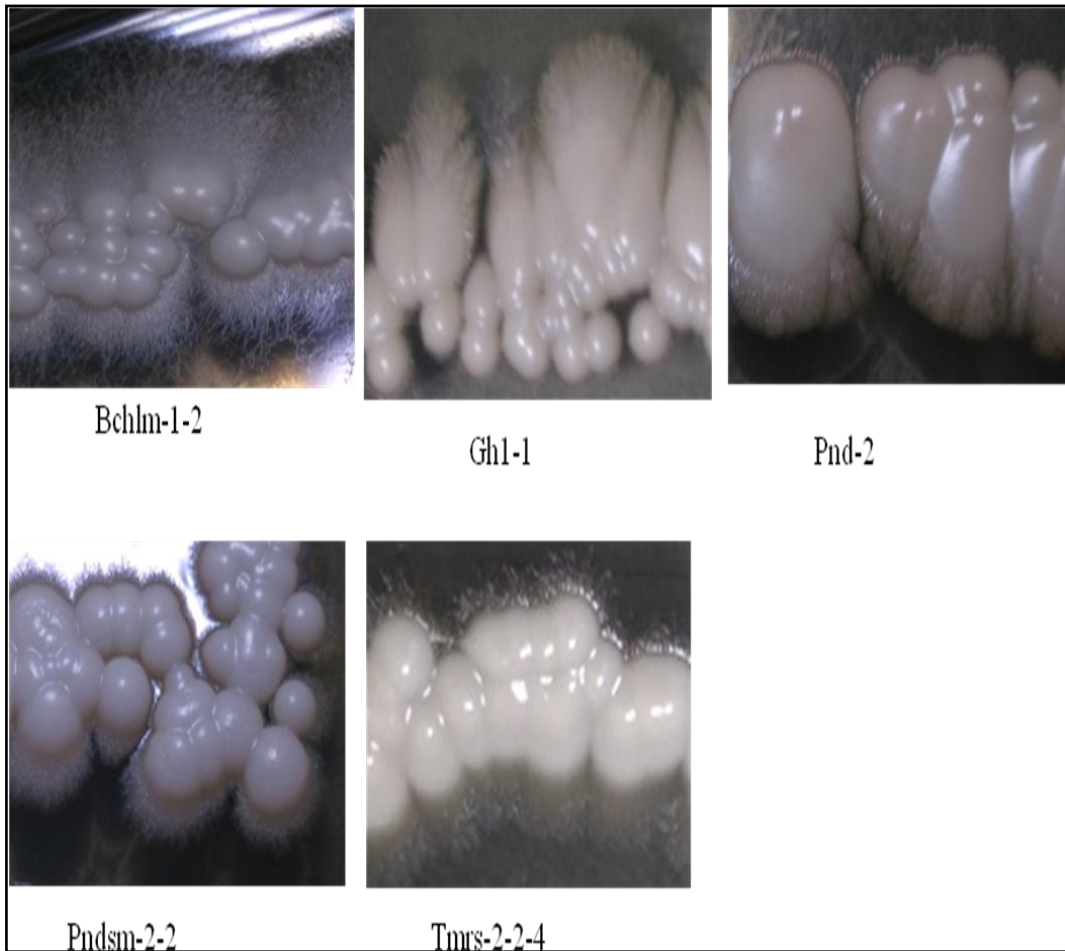
### **3.3.2 Fractality Index of sediment**

Strain Bchlm-1-2 showed highest Fractality Index of sediment (Fig. 3.22) formed by binding the suspended solid and yeast.

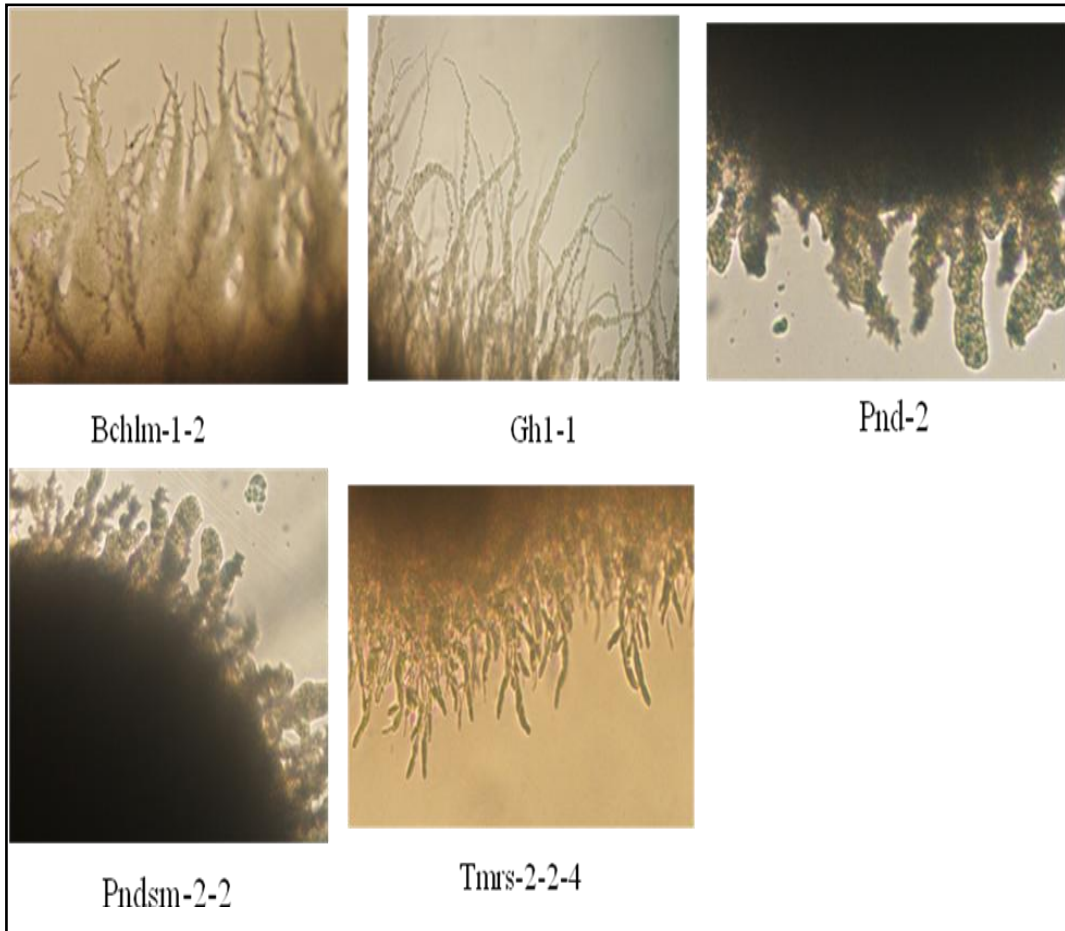
### **3.3.3 Preliminary study of Fractality Index of colony margin**

A preliminary analysis of the fractality Index of the thirteen strains (as discussed in Fig.3. 21 and Fig 3.23) revealed that the inner margins have a large fractality Index compared to outer edges. Besides Fractal dimension of colony margins the F.I. of the sediment is showed highest by same culture, designated as Bchlm-1-2. (Fig 3.22 and 3.24).

Moreover, negligible deviation was observed in the Fractal dimension of the sediment formed by yeast and respective suspended solids. (Fig.3.22 and 3.24). Almost similar Fractality Index of sediment in the every replicates reveals, sediment structure were same, formed by specific culture and its respective clay.



**Figure. 3.19: The complex margins of yeast strains**



**Figure. 3.20: The fractal margins of yeast strains under 100x magnification**



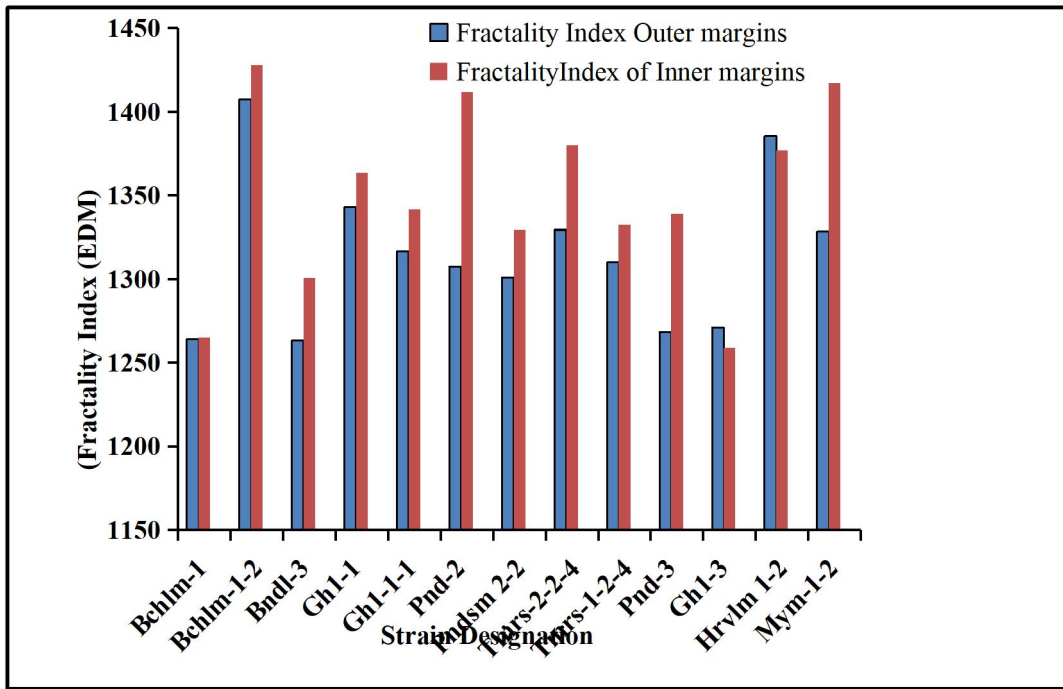


Figure.3.21: Fractality index of strains' inner and outer margins

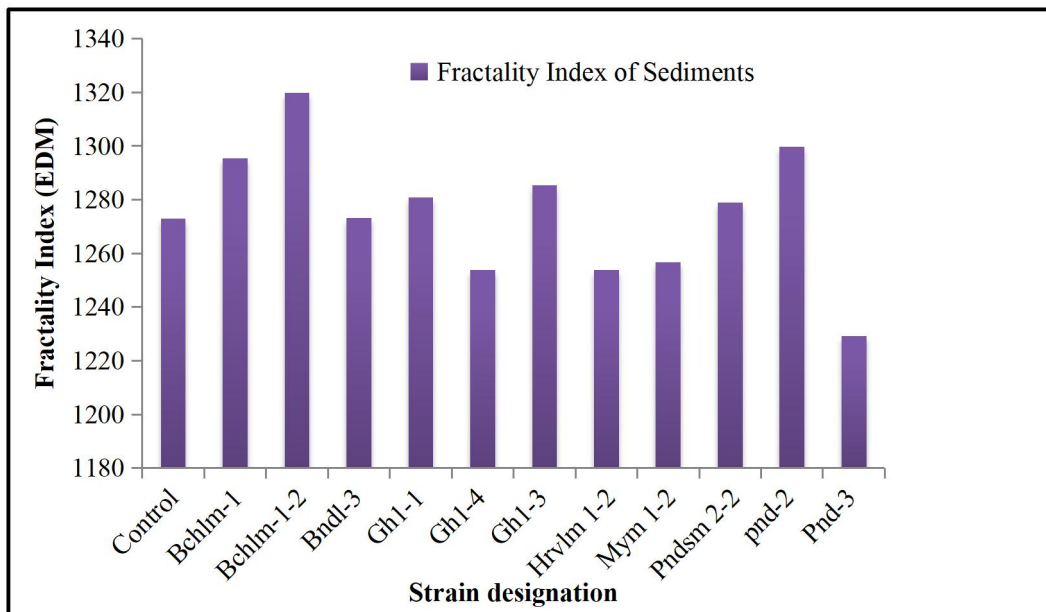
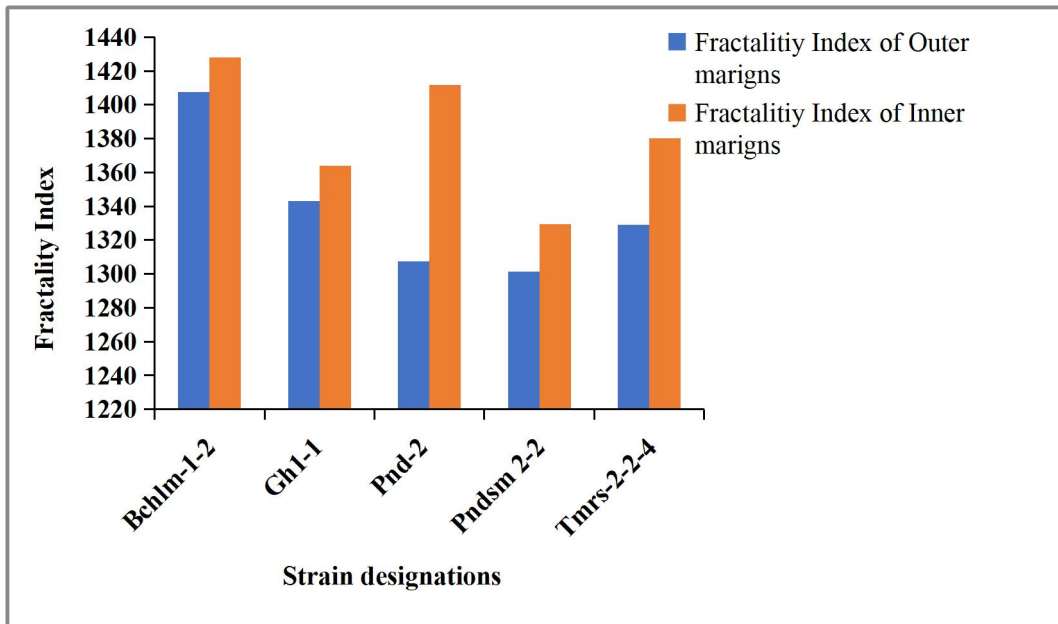
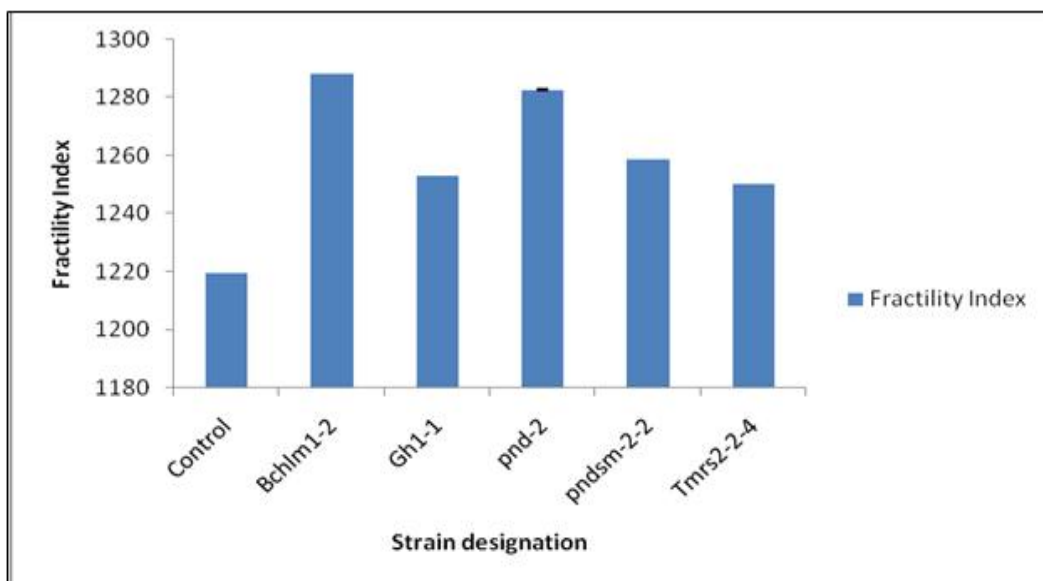


Figure.3.22: Fractality Index of sediments



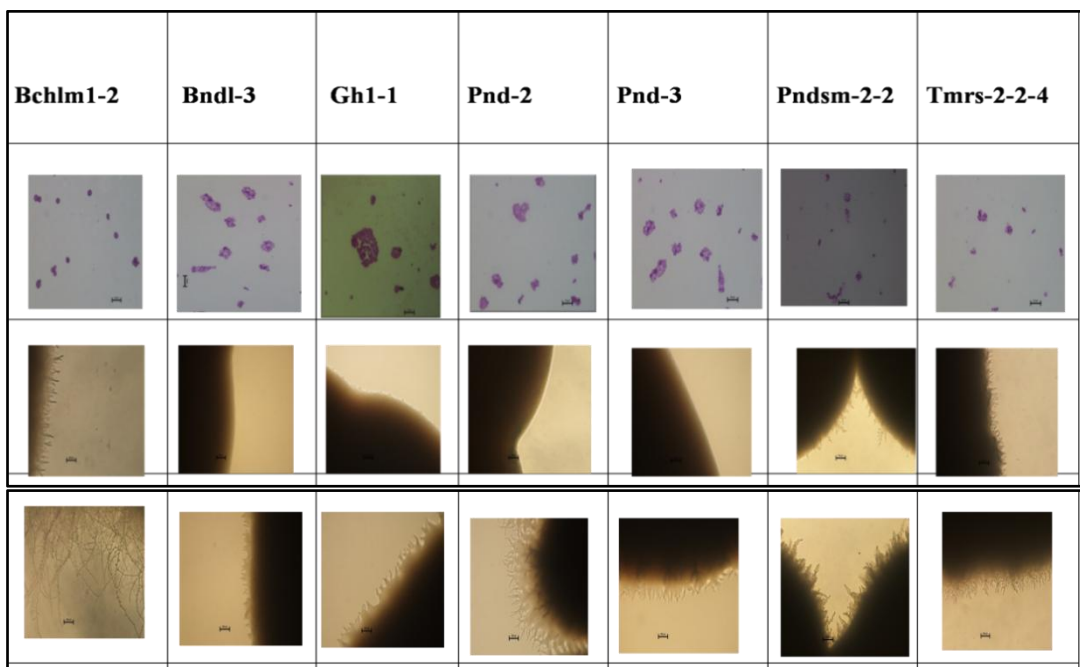
**Figure3.23: Fractality Index of inner and outer margins**



**Figure.3.24: Fractality Index of sediments**

### 3.3.4 Complex colony margins









Colony margins in cultures were complicated. The colonies of representative cultures are shown in Fig.3.25. The inner borders of all strains were more complex than the outer margins. Each colony has a unique pattern of fractality structure. The micromorphology (under 100x magnification) of selected cultures were showed unique surface morphology, in detail, photo among all cultures with complex colony borders showed in (Fig.3.25).



**Figure 3.25: Outer inner margins of yeasts strains**

#### 3.3.4.1 Morphology of sediments

With clay colloids, all tested cultures formed solid sediment (Fig.3.26). Sediment showed compactness, compared to control sediment. Furthermore, among all the strains tested, Bchlm-1-2 exhibits compact sediment as it remained unchanged after disturbing the settled sediment (under 40x magnification).

<b>Control</b>	
<b>Bchl1-2</b>	
<b>Bndl-3</b>	
<b>Gh1-1</b>	
<b>Pnd-2</b>	
<b>Pnd-3</b>	
<b>Pndsm-2-2</b>	
<b>Tmrs-2-2-4</b>	

**Figure.3.26: Micromorphology of sediments**

The amplified picture of yeast strains taken on 6<sup>th</sup> day by 10x objective lens is shown in Fig.3.19 to show cause the fractal margins of the five strain designations. Micromorphology of colonies margins shows variation (Fig.3.20) in the colony margin among the cultures. The observations for different strains were as follows:

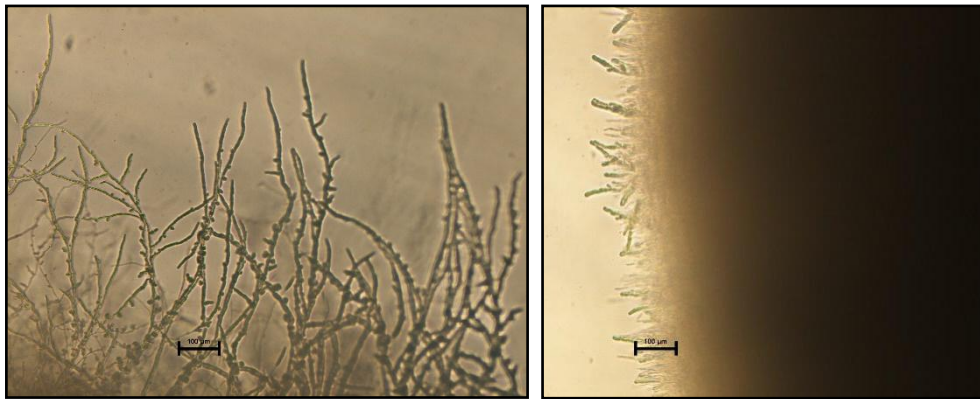
1. Strain designated as Bchlm-1-2 shows high growth rate of the complexity of colony margins and complexity on its edges within 24 hr of inoculation. Moreover, this strain shows more complex growth of margins with respect to time.
2. Strain designated as Gh1-1 shows less complexity on the surface of colony margin on the second day. However, its colony margin shows very complex structure after 3 days.
3. Strain designated as Pndsm-2-2 showed smooth edges at the beginning but after three days it exhibits complex growth of colony margins with respect to time.
4. Strain designated as Pnd-2 showed slow growth rate of its complexity of margins and after three days we observe irregular and complex colonies with increasing complexity with respect to time.
5. The cultures Tmrs-2-2-4 shows complex colony edges within two days.

**Table.3.6 Fractality Index of colony margins at different time interval**

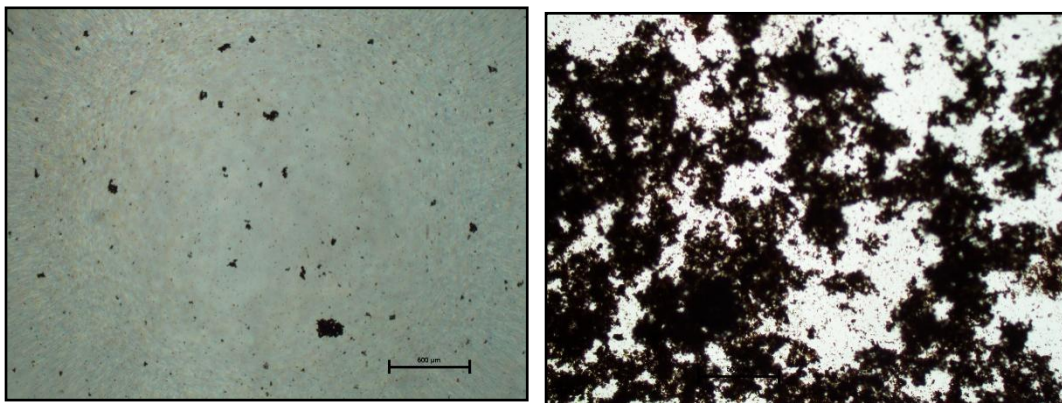
Day	Strains designation	Peripheral		Middle		Centre	
		Outer	Inner	Outer	Inner	Outer	Inner
2	Bchlm-1-2	1194	1192	1194	1225	1224	1281
	Gh1-1	1185	1184	1186	1181	1185	1212
	Pnd-2	1191	1186	1207	1219	1184	1238
	Pndsm-2-2	1184	1191	1183	1190	1177	1186
	Tmrs-2-2-4	1174	1180	1178	1190	1054	1179
4	Bchlm-1-2	1287	1309	1302	1309	1300	1311
	Gh1-1	1191	1231	1227	1231	1262	1205
	Pnd-2	1213	1179	1226	1289	1231	1226
	Pndsm-2-2	1216	1235	1230	1249	1235	1311
	Tmrs-2-2-4	1182	1187	1180	1258	1207	1183
6	Bchlm-1-2	1410	1422	1357	1408	1425	1397
	Gh1-1	1304	1268	1288	1230	1280	1326
	Pnd-2	1204	1231	1242	1278	1221	1256
	Pndsm-2-2	1193	1215	1305	1216	1269	1353
	Tmrs-2-2-4	1247	1281	1269	1366	1331	1277

### 3.3.5 Colony margins

The colony border of the most promising culture had a complex shape. As shown in fig. 3.27, the inner layer of the colony margin's was more complicated than the outer layer under 1000X magnification.



**Figure. 3.27: (a) Inner (b) outer layer of promising yeast's colony margin**



**Figure 3.28: (c) Pinch of sediments of control tube (d) Pinch of sediments mixed with yeast**

### 3.3.6. Particle size analysis of the test and control sediments

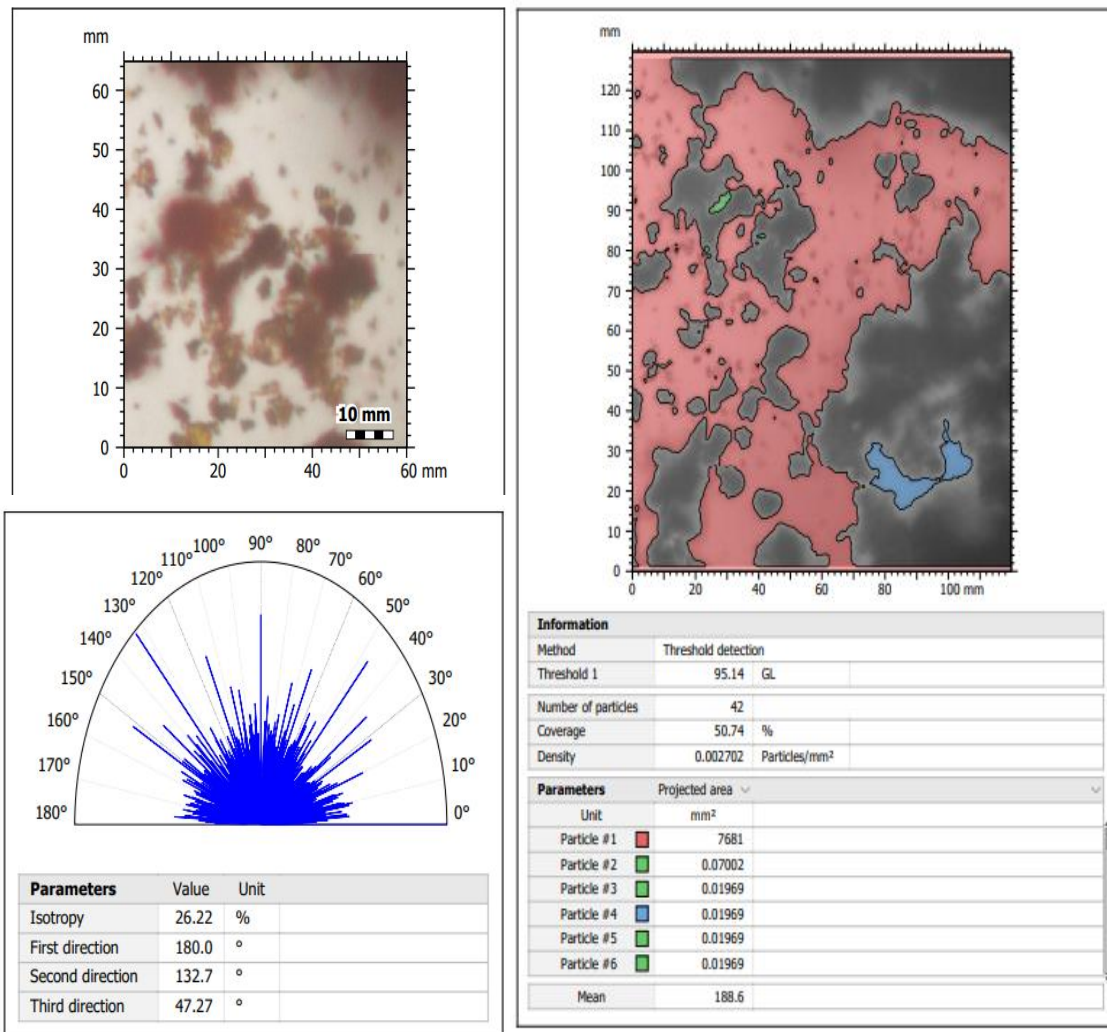


Figure 3.29.: Particle size analysis of control sediments



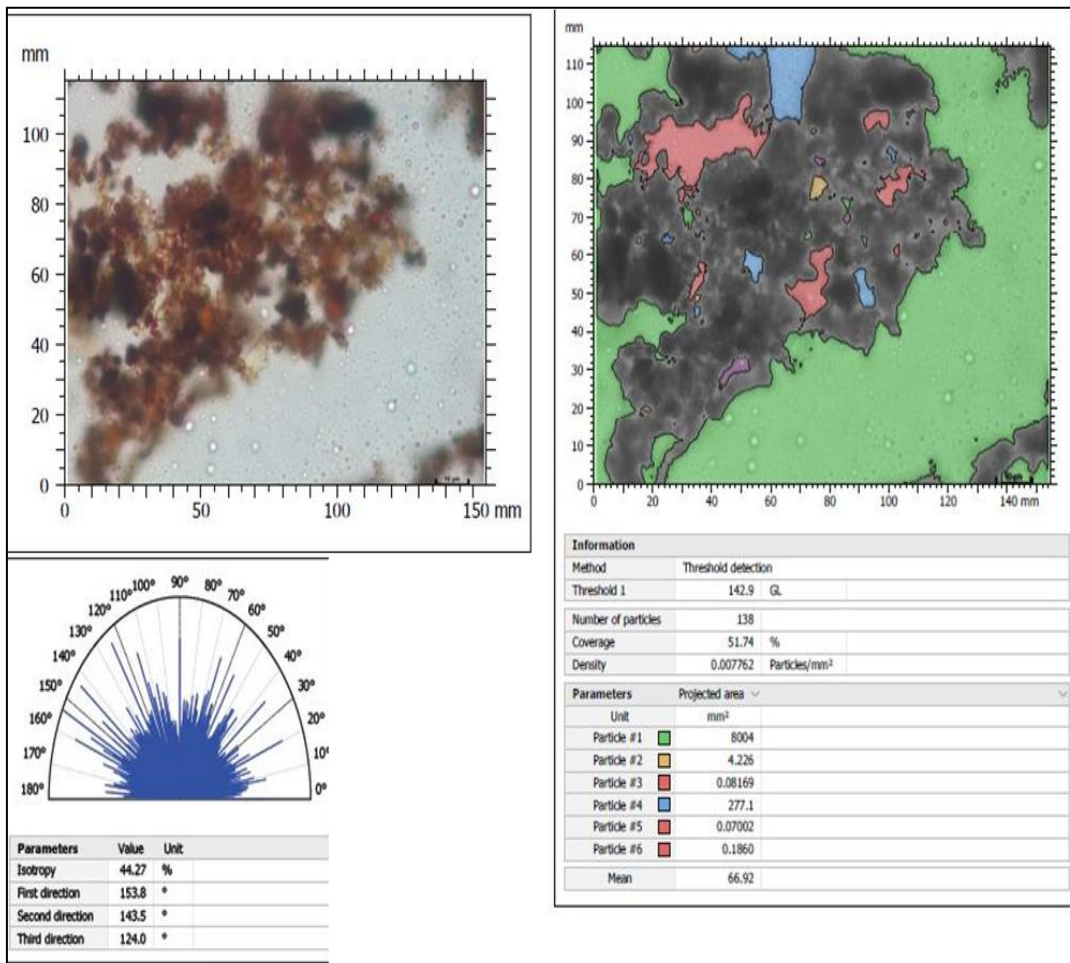
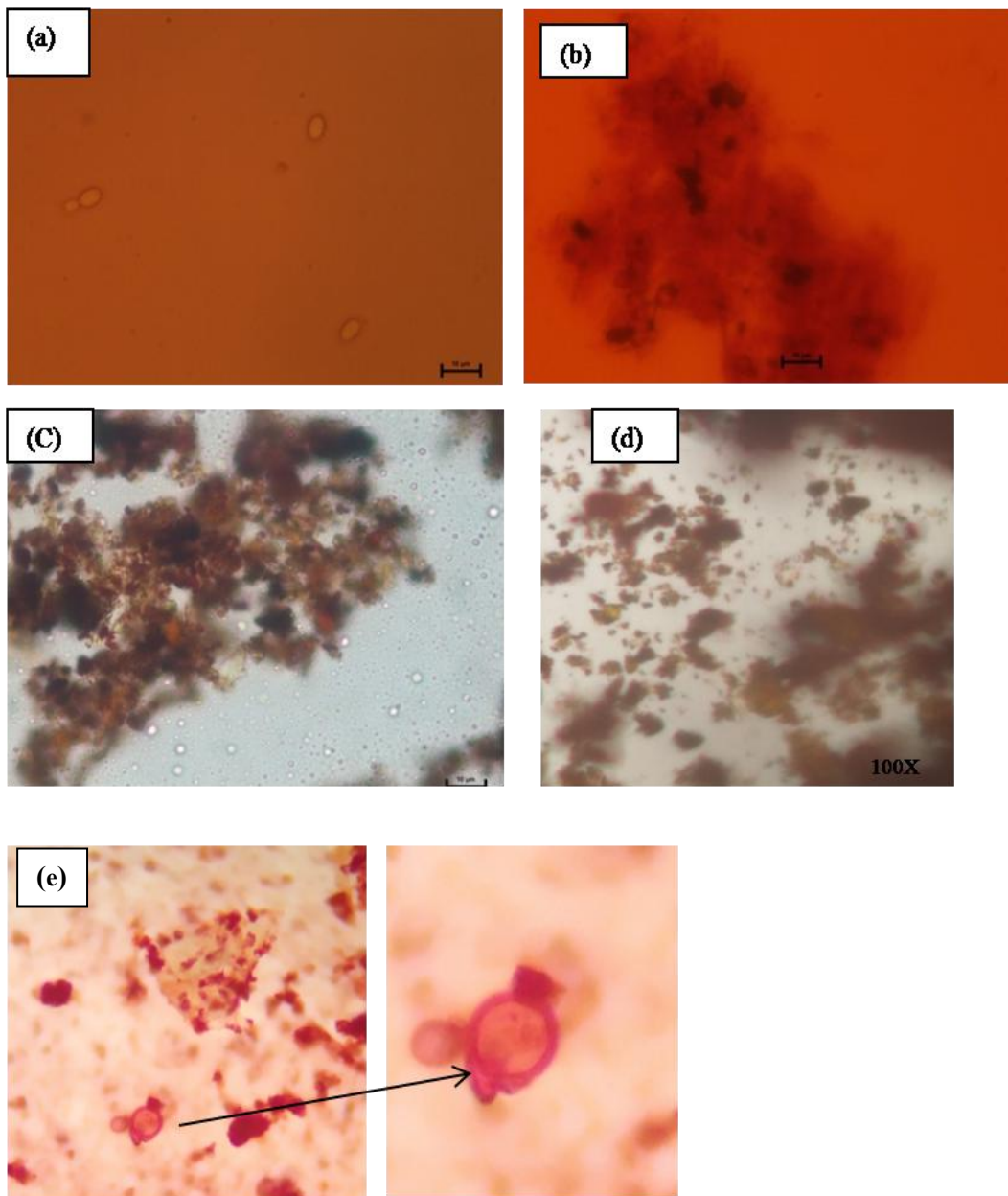


Figure 3.30: Particle size analysis of test sediments

### 3.4. Control sediment, test sediment and yeast



**Figure.3:31 (a) Micromorphology of yeast cells (b) sediment mixed with yeast cells stained in Congo red (c).Test sediment (d) Control sediment mounted in DPX Sediment of Kaolin clay and yeast, under100X objective lens, stained in Congo red.**

### 3.5.1 Water quality in mining region of Goa

**Table.3.7 Water quality parameters of samples, mining areas, non monsoon season**

Parameter (Average)	Arvalem water	Mayem lake	Morlem tank	Morlem spring	Bicholim tank	Pissrullum Pond
<b>Turbidity (NTU)</b>	26 ±0.57	9±0.1	10±1	15±0	22±1.547	22±0.230
<b>TDS</b>	29±	31±1.15	76±1.527	77±1.15	40±0	40±0
<b>EC µs</b>	50±0	55±0	105±5.291	113±2.156	90±5.506	90±0.057
<b>pH</b>	7.2 ±0	6.8±1.73	7.4±1.09	7.2±0	7.4±0.05	6.87±0.0
<b>Temp</b>	28 ±0	29.7±0.1	27±0	27±0	28.9±1.6	26.9
<b>D.O</b>	7.25±0	7±0	7.9±0	7.2±0	7.2±0	7.1±0
<b>Fe</b>	<b>0.6542</b>	<b>0.8056</b>	<b>1.4812</b>	-	-	<b>0.560</b>
<b>Mn</b>	<b>0.5915</b>	<b>0.6460</b>	<b>0.6565</b>	<b>0.7262</b>	-	<b>0.898</b>
<b>K</b>	0.77	1.7	1.4	1.2	1.3	1.7
<b>Na</b>	7	7	7	7	7	7
<b>Cl<sup>-</sup></b>	51.8±0.2	33±0.058	51.6±0.79	35.92±0.4	33±0.208	44±0.1
<b>NO<sub>3</sub><sup>-</sup></b>	0.29±0.16	0.65±0.0 01	0.129±0.0 01	0.65±0.00 1	0.55±0.0 1	0.54±0.01
<b>PO<sub>4</sub><sup>-3</sup></b>	0.0176	0.0917	0.0787	0.0398	0.07863	0.04095
<b>SO<sub>4</sub><sup>-2</sup></b>	0.2±0	0.7±0.03	0.7±0.053	0.1±0.001	0.57±0.0	0.24±0.08
<b>Alkalinity</b>	62.33±1	68.3±1.5	88±3.6	96±2.65	86.33±2.	86.66±2.
<b>Total Hardness</b>	68±1.7	48.66±3. 5	129.66±0. 5	119.66±2.	111.66±2	112±1.73

\*Temp=Temperature ,°C All values are in mg/L except temperature, electrical conductivity,pH

**Table. 3.8 Water quality parameters in mining areas in monsoon season**

Parameter (Average)	Arvalem water fall	Mayem lake	Morlem tank	Morlem spring	Bicholim tank	Pissurlum Pond
<b>Turbidity (NTU)</b>	18±0.577	8±0.1	12±0.577	23±2	20±0.1	18±0.577
<b>TDS ppm</b>	28±1.12	30±1.154	76±0	77±0	40±0.577	40±0
<b>EC µs</b>	49±0.577	53±1.527	105±0.57	113±1	90±0.577	89±0.57
<b>pH</b>	7.2±0.057	6.7±0.05	7.4±0.057	7.2±0.05	7.4±0.1	6.87±0.5
<b>Temp(°C)</b>	27±0.57	29.7±0.1	27±0.058	27±0	28±0	26±0.57
<b>DO</b>	6.4±0.252	6.9±0.25	7.1±0	7.25±0.1	7.2±0.58	7±0.116
<b>K</b>	0.72±	0.7±0	0.74±	0.73±0	0.77±0	0.77±0
<b>Na</b>	7±0	7±0	8±0	8±0	6.9±0	7±0
<b>Cl<sup>-</sup></b>	54.9±0.23	30±0.088	50.31±0.6	40.7±0	45±0.17	44±0.08
<b>NO<sub>3</sub><sup>-</sup></b>	0.56±0.00	0.85±0.0	0.28±0.00	0.56±0.0	0.54±0.0	0.53±0.0
<b>PO<sub>4</sub><sup>3-</sup></b>	0.02±0	0.02±0	0.01±0	0.01±0	0.01±0.0	0.01±0
<b>SO<sub>4</sub><sup>2-</sup></b>	1.6±0.1	1.5±0.06	0.9±0	0.9±0	0.87±0.0	0.8±0.06
<b>Alkalinity</b>	62±0	60.33±1.	96±3.61	89±1	83.33±.1	86.66± 2
<b>Total Hardness</b>	59±1.185	48±2.645	130.66±0.08	122.3±0.52	111.66±02	112.09±2

\*Temp=Temperature ,°C All values are in mg/L except temperature, electrical conductivity,pH

### 3.5. Correlation coefficients among various water quality parameters

Table. 3.9 Arvalem waterfall (Non monsoon)

	Total Hardness	Alkalinity	SO <sub>4</sub> <sup>-2</sup>	PO <sub>4</sub>	NO <sub>3</sub> <sup>-</sup>	DO	Temp (°C)	pH	EC $\mu$ s	TDS ppm	Turbidity (NTU)	Turbidity (NTU)
												1
										1	0	0
									1	0.11	0	0
								1	0	0	0	0
							1	1	0	0	0	0
						1	0.69	0	0	0	0.27	0
					1	0.69	1	-0.69	0.69	-0.32	-0.5	-0.5
				1	0.5	-0.28	0.5	-1	1	0.46	-1	-1
			1	0	0	0	0	-0.5	0.9	0.9	0	0
	1	0	0	1	-0.94	-0.89	-0.94	0	0	0	0.18	0.18
	1	-0.18	0	1	0.5	-0.27	0.5	-0.5	0.9	0.9	-1	-1

\*Temp=Temperature All values are in mg/L except temperature, electrical conductivity, pH

**Table.3.10 Arvalem water fall (Monsoon)**

	Turbidity (NTU)	TDS	EC $\mu$ s	pH	Temp	DO	NO <sub>3</sub> <sup>-</sup>	PO <sub>4</sub> <sup>-3</sup>	SO <sub>4</sub> <sup>-2</sup>	Alkalinity	Total Hardness
	1										
	-0.9	1									
	-0.5	0.46	1	1							
	0.5	-0.46	-1	1							
	-0.5	0.46	1	-1	1						
	-0.46	0	0.43	0	0	1					
	0.11	0	-0.72	0	0	-0.94	1				
	0	0	0	0	0	0	0	1			
	0.44	0	0.98	0	0	0.59	-0.83	0	1		
	0	0	0	0	0	0	0	0	0	1	
	-0.02	0	0.78	0	0	0.9	-0.9	0	0.8	0	1

\*Temp=Temperature All values are in mg/L except temperature, electrical conductivity, pH

**Table.3.11 Morlem pond (Non monsoon)**

Total Hardness	Alkalinity	SO <sub>4</sub> <sup>-2</sup>	PO <sub>4</sub> <sup>-3</sup>	NO <sub>3</sub> <sup>-</sup>	DO	Temp (°C)	pH	EC μs	TDS	Turbidity (NTU)	
									0	1	Turbidity (NTU)
									1	0	TDS ppm
								1	-0.12	0	EC μs
							1	0	0	0	pH
						1	0	0	0	0	Temp
					1	0	0	0.27	-0.93	0	DO
				1	0.13	0	0	0.92	-0.5	0	NO <sub>3</sub> <sup>-</sup>
				1	0.95	0	0	0.02	0.9	0	PO <sub>4</sub> <sup>-3</sup>
		1	-0.4	-0.42	0.7	0	0	-0.88	0.37	0	SO <sub>4</sub> <sup>-2</sup>
	1	0.35	0.24	-0.6	0.7	0	0	-0.98	0.32	0	Alkalinity
1	0.96	0.59	0.240	-0.98	0.06	0	0	0.8	-0.69	0	Total Hardness

\*Temp=Temperature °C, All values are in mg/L except temperature, electrical conductivity, pH

**Table.3.12 Morlem pond (Monsoon)**

Total Hardness	Alkalinity	SO <sub>4</sub> <sup>-2</sup>	PO <sub>4</sub> <sup>-3</sup>	NO <sub>3</sub> <sup>-</sup>	DO	Temp (°C)	EC µs	TDS	Turbidity (NTU)	Turbidity (NTU)
									1	Turbidity (NTU)
									0	TDS
								1	0	EC µs
								-	0.86	pH
						1		-	0	Temp (°C)
					1	0.5		-	0.86	DO
				1	0.86	0.86		-	1	NO <sub>3</sub> <sup>-</sup>
			0	0	0	0		-	0	PO <sub>4</sub> <sup>-3</sup>
		0	0	0	0	0		-	0	SO <sub>4</sub> <sup>-2</sup>
	1	0	0	1	0.87	0.87	--	-	1	Alkalinity
1	-0.32	0	0	-0.32	0.86	0.18	-	-	-0.3	Total Hardness

\*Temp=Temperature °C All values are in mg/L except temperature, electrical conductivity, pH



**Table 3.13 Morlem tank (Non monsoon)**

Hardness	Alkalinity	SO <sub>4</sub> <sup>-2</sup>	PO <sub>4</sub> <sup>-3</sup>	NO <sub>3</sub> <sup>-</sup>	DO	Tem (°C)	pH	EC µs	TDS	Turbidity (NTU)	Turbidity (NTU)
										1	Turbidity (NTU)
									1	0	TDS
								1	0	0.5	EC µs
							1	0	0	0	pH
						1	0	0	0	0	Temp
					1	0	-0.95	-0.95	0	0	DO
				1	0.76	0	0.5	0.5	0	0.18	NO <sub>3</sub> <sup>-</sup>
			1	0	0	0	0	0	-0.5	-0.5	PO <sub>4</sub> <sup>-3</sup>
			0	0	0	0	0	0	0	0	SO <sub>4</sub> <sup>-2</sup>
	1	0	0	0	-0.9	0	0	0	0	0	Alkalinity
1	0	1	0	0.96	0	0	0	0.97	0	-0.24	Total Hardness

\*Temp=Temperature °C All values are in mg/L except temperature, electrical conductivity,

**Table 3.14 Morlem tank (monsoon)**

Total Hardness	Alkalinity	SO <sub>4</sub> <sup>-2</sup>	PO <sub>4</sub> <sup>-3</sup>	NO <sub>3</sub> <sup>-</sup>	DO	Temp (°C)	pH	EC µs	TDS	Turbidity (NTU)	Turbidity (NTU)
										1	Turbidity (NTU)
									1	-0.98	TDS ppm
							0	1	0.62	-0.76	EC µs
							1	0	-3.80	0	pH
						1	0	0	0	0	Temp (°C)
					1	0	-9.5	-2.19	-0.79	0.66	DO
				1	-0.76	0	1.96	-0.66	0.19	0	NO <sub>3</sub> <sup>-</sup>
			1	0.5	-0.95	0	-7.85	0.33	0.94	0.9	PO <sub>4</sub> <sup>-3</sup>
		1	-0.40	0.59	-0.91	0	-2.03	-0.99	-0.67	-0.87	SO <sub>4</sub> <sup>-2</sup>
	1	0.35	0.34	0.91	-0.91	0	0	-0.41	0.45	-0.27	Alkalinity
1	0.96	0.59	0.5	1	-0.76	0	-2.	-0.65	0.18	0	Total Hardness

\*Temp=Temperature °C All values are in mg/L except temperature, electrical conductivity,

**Table.3.15 Mayem lake (Non Monsoon)**

Total Hardness	Alkalinity	SO <sub>4</sub> <sup>-2</sup>	PO <sub>4</sub> <sup>-3</sup>	NO <sub>3</sub> <sup>-</sup>	DO	Temp	pH	EC μs	TDS	Turbidity (NTU)	
										1	Turbidity (NTU)
									1	-0.8	TDS
								1	0.1	0	EC
							1	0.9	0.5	-2	pH
						1	0.5	0.18	1	-0.8	Temp
					1	0	0.9	-0.8	-0.8	0.8	DO
				1	0.	0	-0.5	-4	-4	-3	NO <sub>3</sub> <sup>-</sup>
				0	0	0	0	0	0	0	PO <sub>4</sub> <sup>-3</sup>
		1	0	4.	0.8	0.8	0.8	-1	-1	1	SO <sub>4</sub> <sup>-2</sup>
	1	0.1	0	0.9	-0.4	-0.3	-0.3	-0.2	-0.2	0.2	Alkalinity
1	0.6	-0.6	0	0.7	0	0	-0.9	0.6	0.6	-0.6	Total Hardness

\*Temp=Temperature °C, All values are in mg/L except temperature, electrical conductivity, pH

**Table.3.16: Mayem lake (Monsoon)**

	Turbidity (NTU)	TDS ppm	EC $\mu$ s	pH	Temp ( $^{\circ}$ C)	DO	NO <sub>3</sub> <sup>-</sup>	PO <sub>4</sub> <sup>-3</sup>	SO <sub>4</sub> <sup>-2</sup>	Alkalinity	Total Hardness
	1	1	1	1	1	1	1	1	1	1	1
	-1	-1	-1	-0.8	0	-0.5	-0.5	-0.8	0.5	-0.9	0.2
	-1	1	-0.8	0	0	0.5	0.5	0.8	0.5	0.9	0.3
	0	0	0	0	1	-0.5	-0.5	-0.8	0.5	-0.9	0.23
	-0.5	0	-0.5	0.5	0.5	1	1	0.8	0.5	0.3	-0.9
	-0.5	0	-0.8	0.5	-0.5	0.8	0.8	0.5	0.5	0.3	-0.9
	-0.8	0	-0.8	0.5	0.5	0.5	0.5	0.5	0.5	0.3	-0.9
	0.5	0	0.5	0.5	0.5	0.5	0.5	0.5	0.5	0.3	-0.9
	-0.9	0	-0.9	0.9	-0.9	0.3	0.3	-0.6	-0.6	1	-0.03
	0.2	0	0.2	0.3	0.23	-0.9	-0.6	-0.6	-0.6	-0.03	1

\*Temp=Temperature All values are in mg/L except temperature, electrical conductivity,pH

**Table.3.17: Bicholim tank (Non Monsoon)**

	Turbidity (NTU)	TDS	EC $\mu$ s	pH	Temp (°C)	DO	NO <sub>3</sub> <sup>-</sup>	PO <sub>4</sub> <sup>-3</sup>	SO <sub>4</sub> <sup>-2</sup>	Alkalinity	Total Hardness
	1										
	0	1									
	0.51	0	1								
	0.5	0	0.11	1							
	0.55	0	0.99	0.9	1						
	0.18	0	0.94	0.9	0.92	1					
	0.5	0	-0.49	-0.5	-0.4	-0.75	1				
	0.67	0	0.97	0.97	0.98	0.84	-0.29	1			
	0.51	0	0.7	-0.49	-0.44	-0.75	0.99	-0.28	1		
	0.97	0	0	0.69	0.73	0.42	0.28	0.8	0.29	1	
	-0.27	0	0.7	0.7	0.6	0.9	-0.97	0.5	-0.9	-0.03	1

\*Temp=Temperature All values are in mg/L except temperature, electrical conductivity, pH

**Table.3.18: Bicholim tank (Monsoon)**

	Turbidity (NTU)	TDS	EC $\mu\text{s}$	pH	Temp ( $^{\circ}\text{C}$ )	DO	$\text{NO}_3^-$	$\text{PO}_4^{-3}$	$\text{SO}_4^{-2}$	Alkalinity	Total Hardness
	1										
		1									
	-1										
		1	1								
	-1										
		-0.8	-0.8	1							
	0.8										
		0	0	0	1						
	0										
	-0.5				-0.5	1					
		0	0	0.5	-0.5	1					
	-0.5										
		0	-0.8	0.8	-0.8	0.8	0.8	1			
	-0.8										
		0	0.5	-0.5	0.5	0.5	0.5	0	1		
	0.5										
		0	0.97	0.97	-0.9	0.27	0.2	0.6	0.6	1	
	0.9										
		0	0.2	-0.27	0.27	-0.9	-0.9	-0.6	-0.6	-0.03	
1	0.28										

**Table.3.18 Pissurlem tank (Non Monsoon)**

	Total Hardness	Alkalinity	SO <sub>4</sub> <sup>-2</sup>	PO <sub>4</sub> <sup>-3</sup>	NO <sub>3</sub> <sup>-</sup>	DO	Temp	pH	EC µs	TDS	Turbidity (NTU)	Turbidity (NTU)
											1	Turbidity (NTU)
										0	0	TDS ppm
									1	0	-0.9	EC µs
								1	-0.6	0	0.7	pH
							1	0.7	0	0	0.2	Temp (°C)
						1	0.9	0.9	-0.2	0	0.4	DO
					-0.2	-0.2	1.0	-0.6	1	0	-0.9	NO <sub>3</sub> <sup>-</sup>
				1	-0.6	-0.6	0.80	-0.22	0	0	0.4	PO <sub>4</sub> <sup>-3</sup>
			1	-0.9	0.7	0.7	0.8	0.3	0.4	0	-0.3	SO <sub>4</sub> <sup>-2</sup>
	1	-0.9	-0.9	0.93	-0.8	-0.8	-0.9	-0.5	-0.2	0	0.1	Alkalinity
	1	-0.9	0.9	-0.9	0.6	0.6	0.8	0.3	0.5	0	-0.3	Total Hardness

\*Temp=Temperature ,°C All values are in mg/L except temperature, electrical conductivity,pH

**Table.3.19 Pissurlem tank (Monsoon)**

	Turbidity (NTU)	TDS ppm	EC $\mu$ s	pH	Temp (°C)	DO	NO <sub>3</sub> <sup>-</sup>	PO <sub>4</sub> <sup>-3</sup>	SO <sub>4</sub> <sup>-2</sup>	Alkalinity	Total Hardness
	1	0									
	0	0	1	1	1	1					
	1	0	1	1	1	1					
	-1	0	-1	1	-1	1					
	0	0	0.5	-0.5	0.5	1					
	-1	0	-1	1	-1	-0.5	1				
	0	0	0	0	0	0	0				
	-1	0	-1	1	-1	-1	1	0	1		
	-0.2	0	-0.2	0.2	-0.2	0.6	0.2	0	0.2	1	
1	0.27	0	0.27	-0.9	-0.2	0.6	0.27	0	1	-9	1

\*Temp=Temperature, °C All values are in mg/L except temperature, electrical conductivity, pH



### 3.6.1. Standardization of parameters for efficient biosedimentation of SMTTW

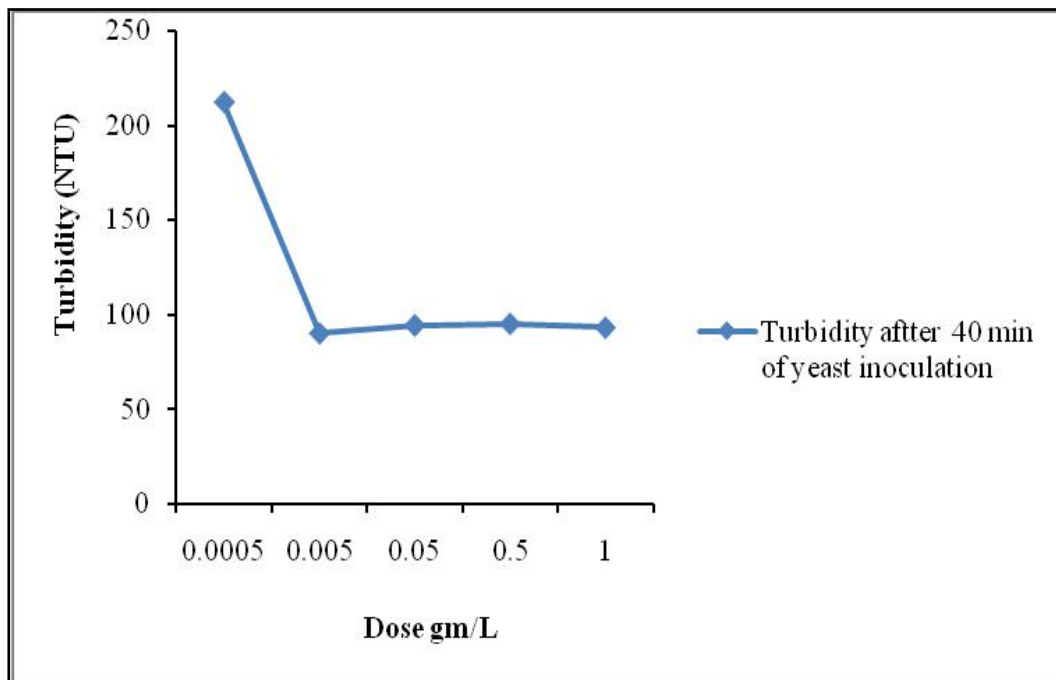


Figure 3.32: Turbidity of mine tailing water at yeast's different doses

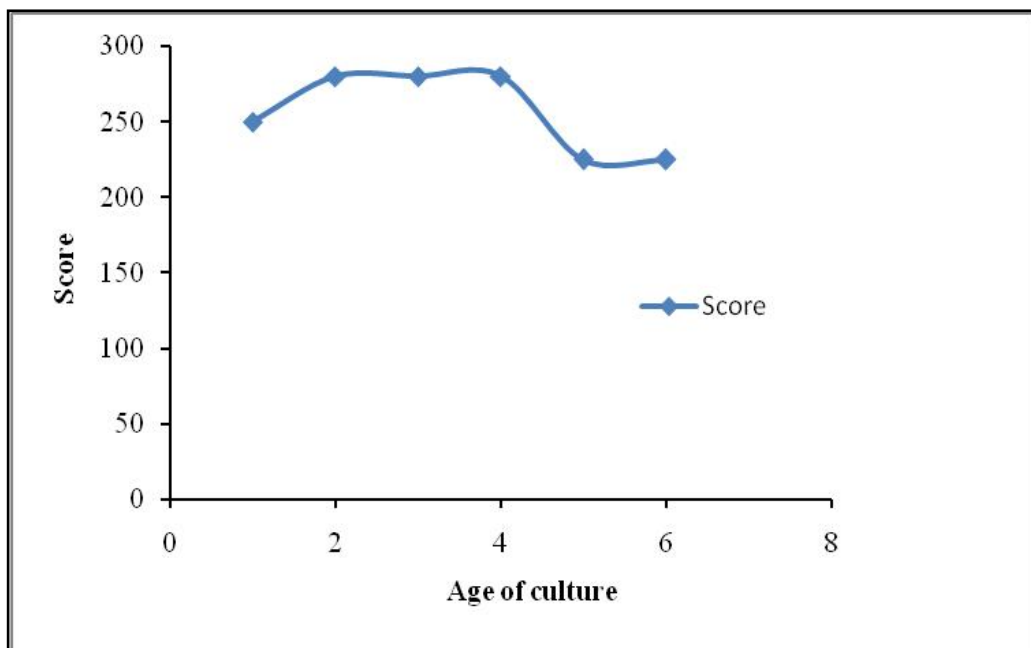


Figure.3.33: Biosedimentation efficiency score at different age of selected strain

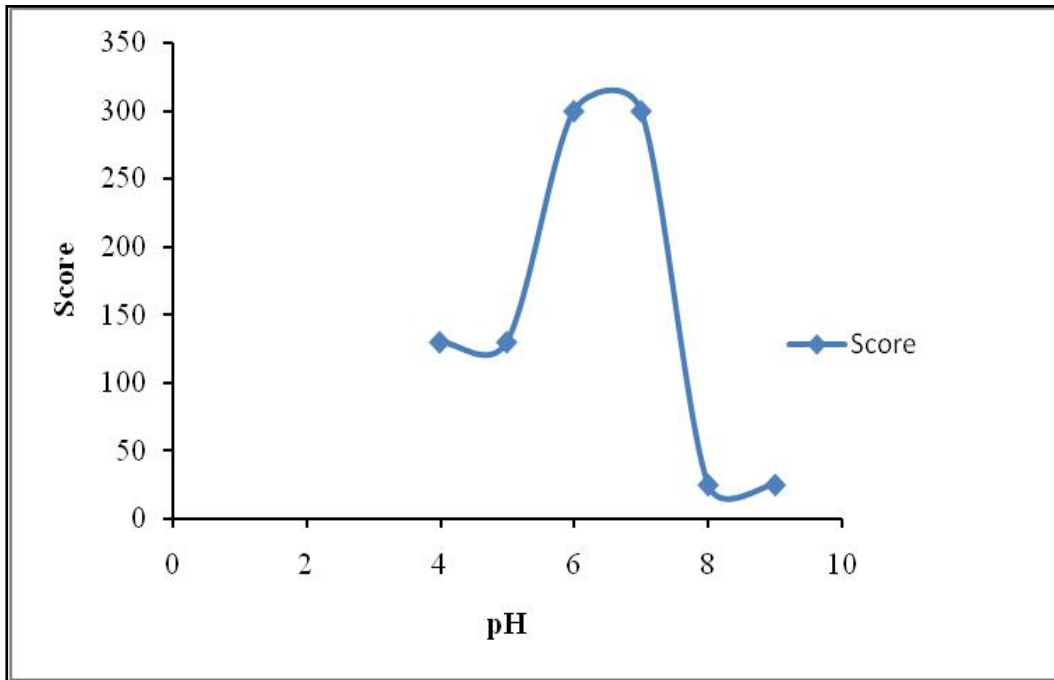


Figure.3.34: Efficiency score of biosedimentation at different pH of selected strain

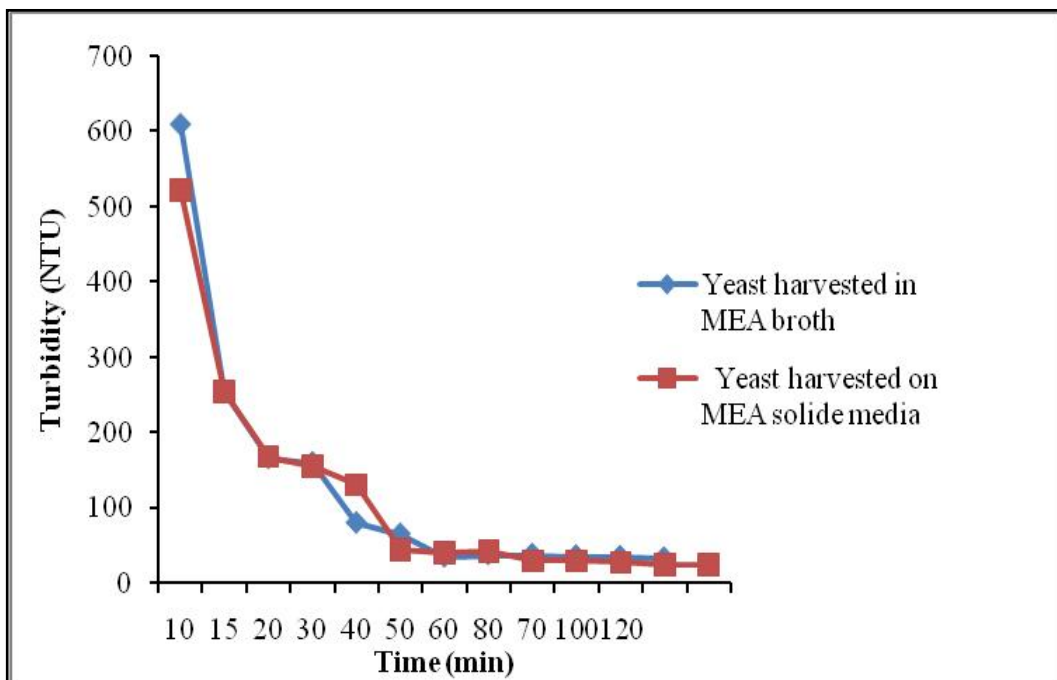


Figure.3.35: Kinetics of turbidity reduction

Biosedimentation efficiency score was highest for 4<sup>th</sup> and 5<sup>th</sup> day old culture, and biosedimentation score was highest at pH six and seven. (Fig.3.33). Moreover, appropriate dose for efficient biosedimentation is 0.005gm/L (Dry weight).

### 3.6.2. Biosedimentation efficiency score (BES) for floc assay after different heat treatment of strain

**Table.3.20 Growth of yeast plated on MEA(a) and BES after heat treatment(b)**

(a)

Treatment time (Sec)	Plate observation after 48 hr of inoculations
0	Mate growth
5	Mate growth
10	Circular white colony 200 in number
20	Colony 100 in number
30	10 colony in number
40	4 colony in number
50	No growth
60	No growth

(b)

Treatment time (Sec)	Biosedimentation efficiency score description			
	A	B	C	Total
0	80	100	100	280
5	50	80	80	210
10	25	80	80	185
15	25	80	80	185
30	80	100	100	180
40	50	80	100	130
50	25	80	100	180
60	25	25	100	150
Control	100	25	0	125

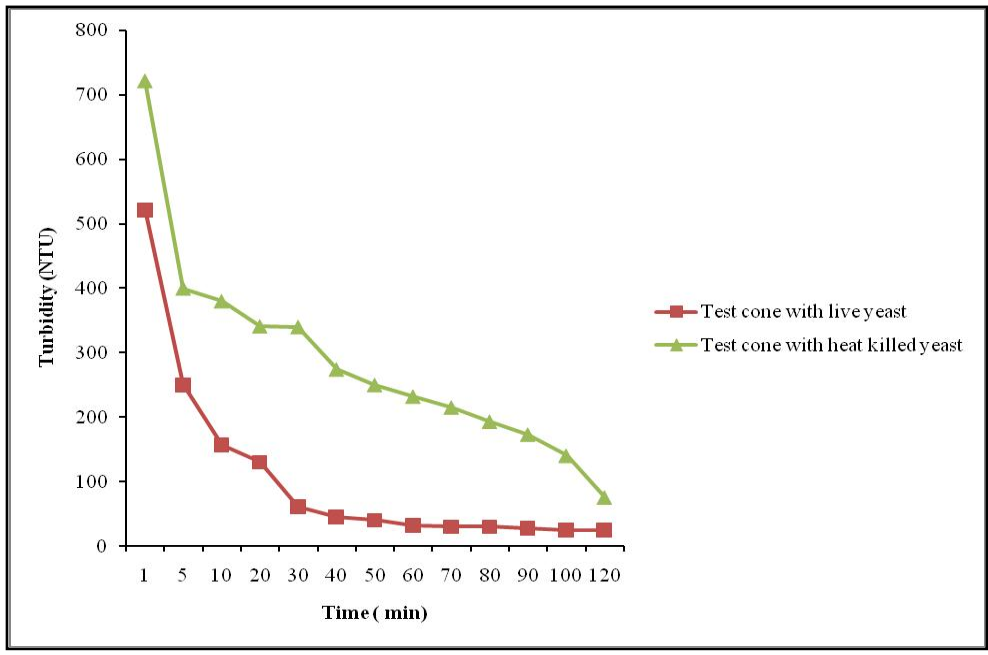


Figure.3.

**36: Biosedimentation kinetics with live yeast and heat killed yeast**

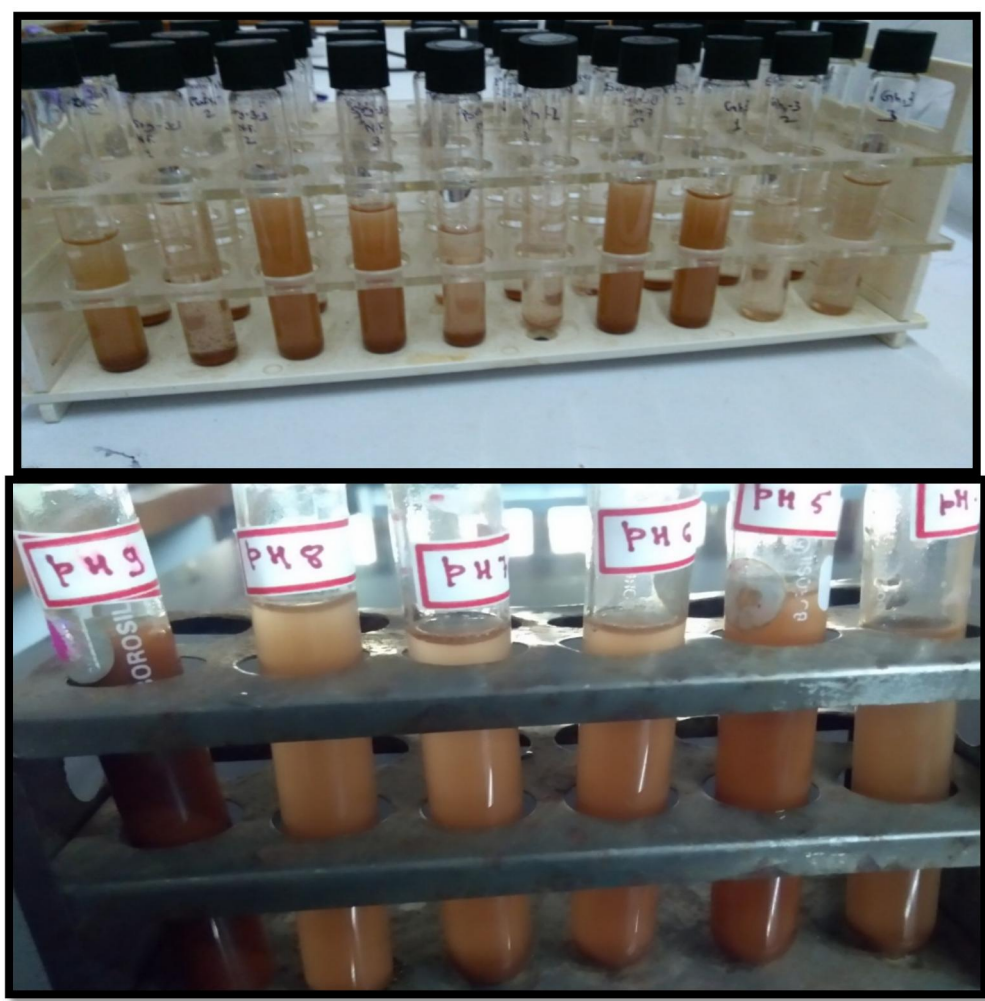
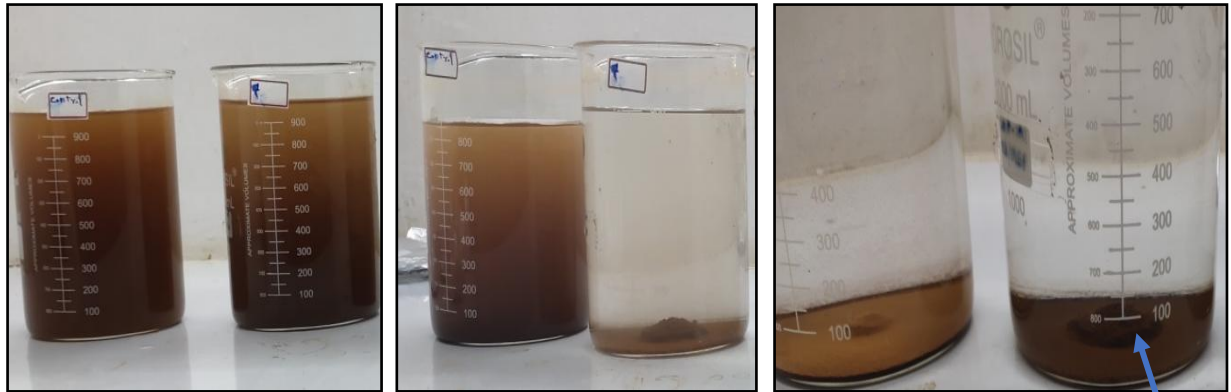


Figure. 3.37: Biosedimentation test assay



Dark colour sediment after yeast addition

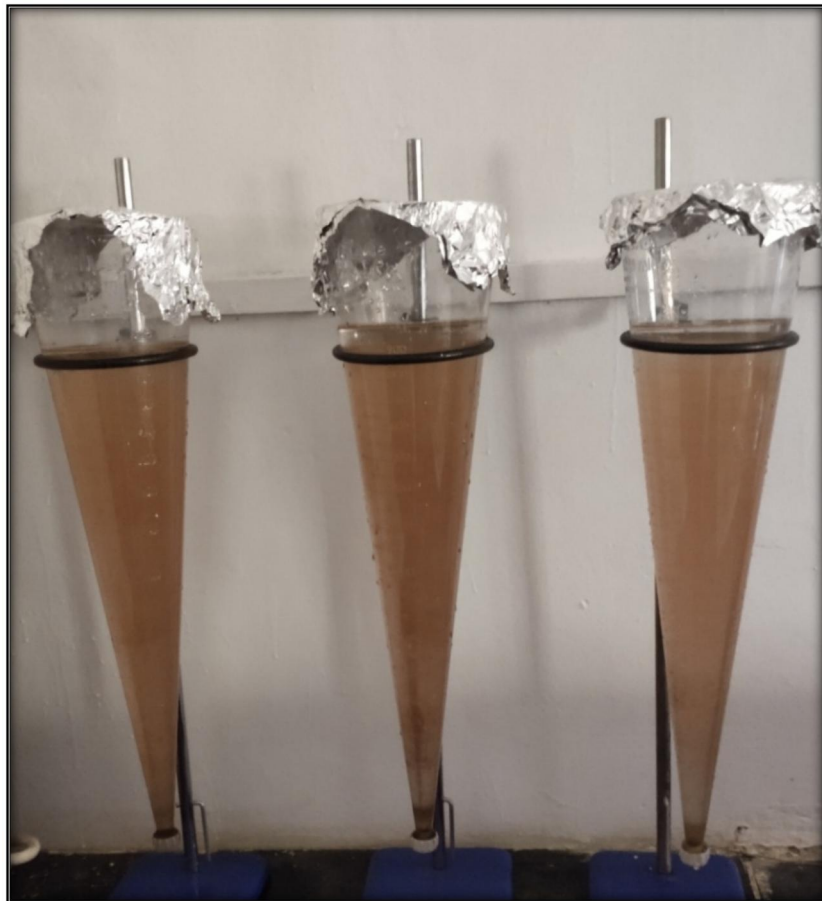


Figure. 3.38: Setup for biosedimentation test at 1 L scale

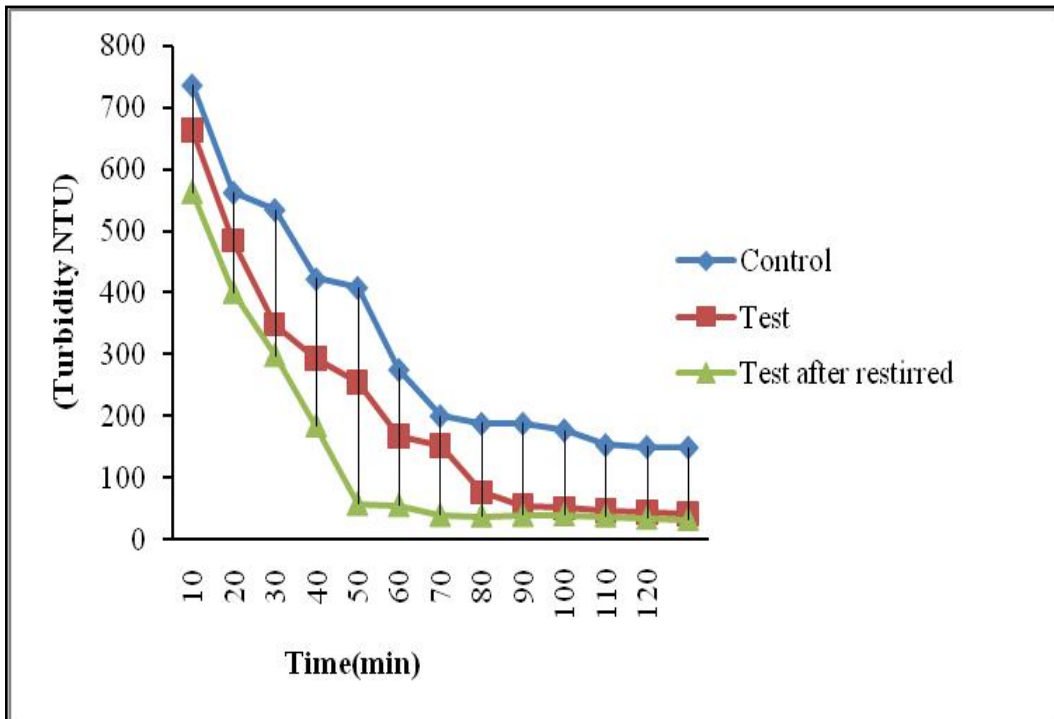


Figure 3.39: Kinetics of turbidity reduction in 1L Imhoff cone

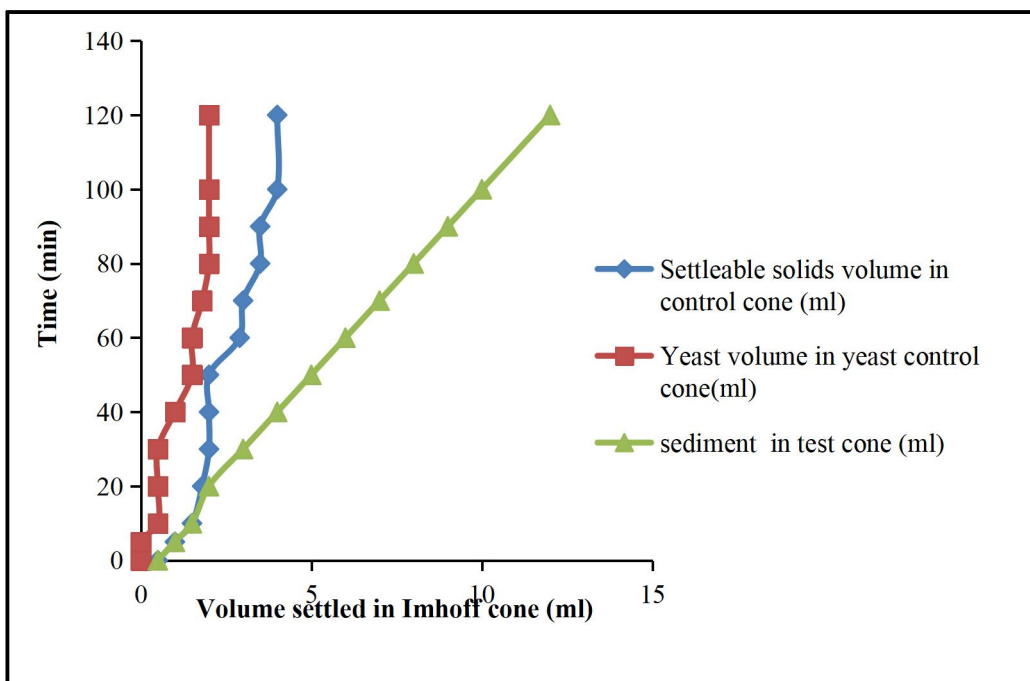


Figure.3.40: Sediment volume (ml) in Imhoff cone

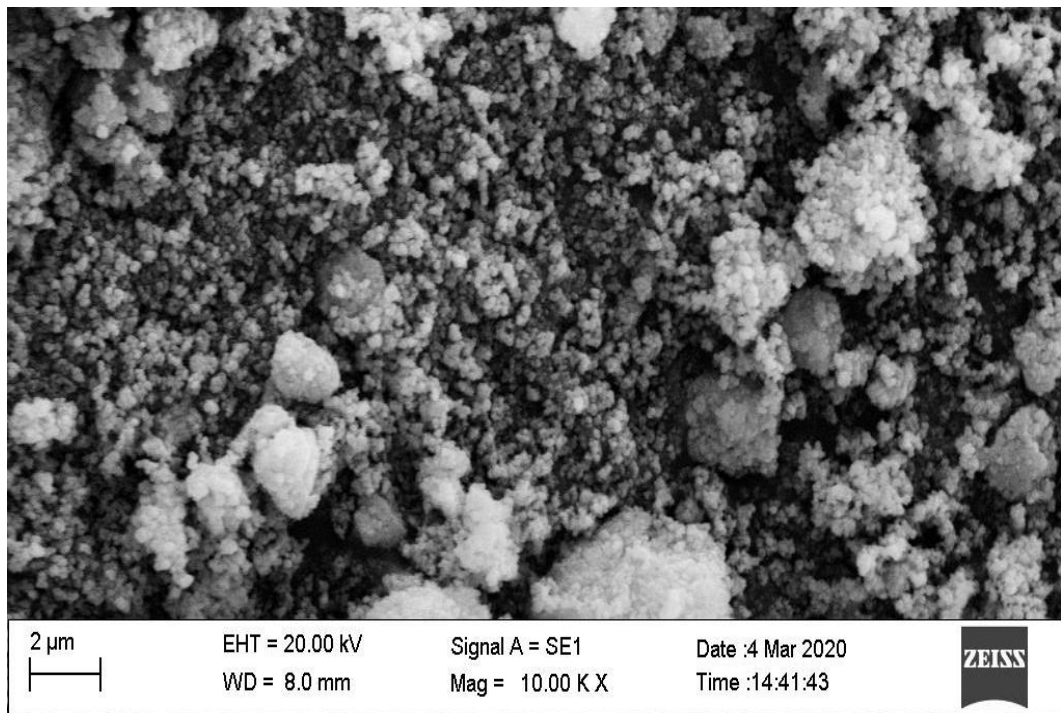
The reduction in turbidity of mine tailing water was higher in the test cone than in the control cone (Fig.3.36). Alive yeast reduced turbidity better than heat-killed (Fig.3.35). Further yeast grown on MEA plates and in MEA broth under shaking conditions demonstrated the similar trend of turbidity decrease (Fig.3.35). Moreover, soil sediment was higher in test cone with compared to control cone (Fig.3.40), 10 to 15 ml sediment was observed after 40-60 min of yeast addition. Sediment deposition was not compact. Sedimentation pattern revealed porous or filled with water or air. While in control cone sediment volume was less than 5ml within 120 min. Moreover sediment were tightly packed in control cone bottom.

Hence structural as well as quantity changes were observed among control and test cone sediment.

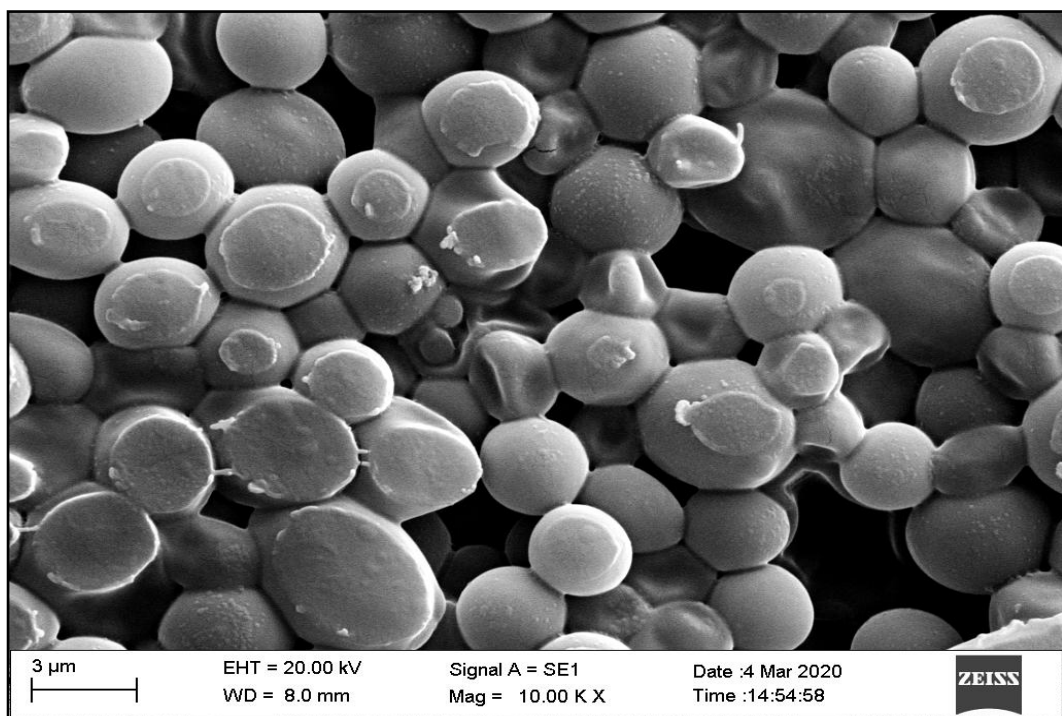
### **3.6.3. SEM analysis of the sediments**

The sediment of SMTTW showed rough morphology under SEM. After yeast was added, it became more flat. SEM Images of yeast revealed a spherical cell shape (fig.3.41(b)). Yeast cells are budding in the nature. The size of each cell varies. In figure(fig.3.42(c)), It can be seen yeast cells are not visible due to particle depositions, (with microscopic clay particles). SEM morphology of test yeast, test sediment, and control sediment revealed that clay particles were rough, and round in shape but sediment texture changed after interacting with yeast. Sediment is looking like flat sheets. Average size of control sediment and test sediment was found 0.232 and 3.567  $\mu\text{m}$  respectively using SEMJ software. Hence SEM images providing prove of interaction of clay colloids and yeast cells.

(a)



(b)



**Figure. 3.41: (a) SEM images of control sediment showing absence of microbial biofilm (b) Yeast cell**



(c)

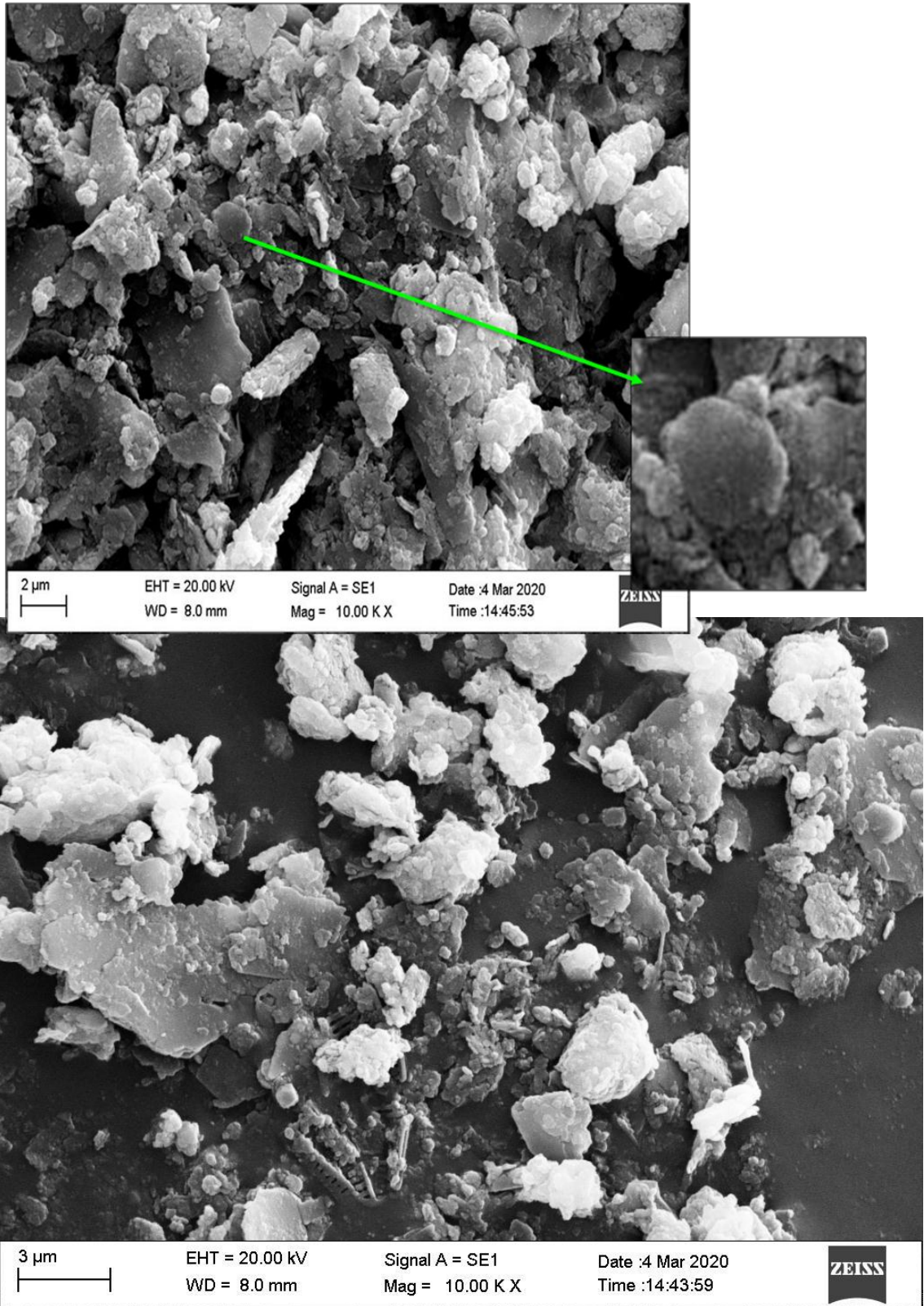


Figure. 3.42: (c) SEM images of test sediments

### 3.6.4. EDX study of sediments

The presence of Fe, Mn, Al, Pd, Au, and Si in mine tailing soil sediment (Control sample) was revealed by EDS (Fig.3.43 (a)). The presence of carbon element was detected by EDS in the mixture of yeast and mine tailing soil sediment (Test sediment) (Fig.3.43 (b)), in addition to all other elements observed in the control sediment. EDS spectra revealed, not much chemical composition changed in the soil sediment after yeast addition

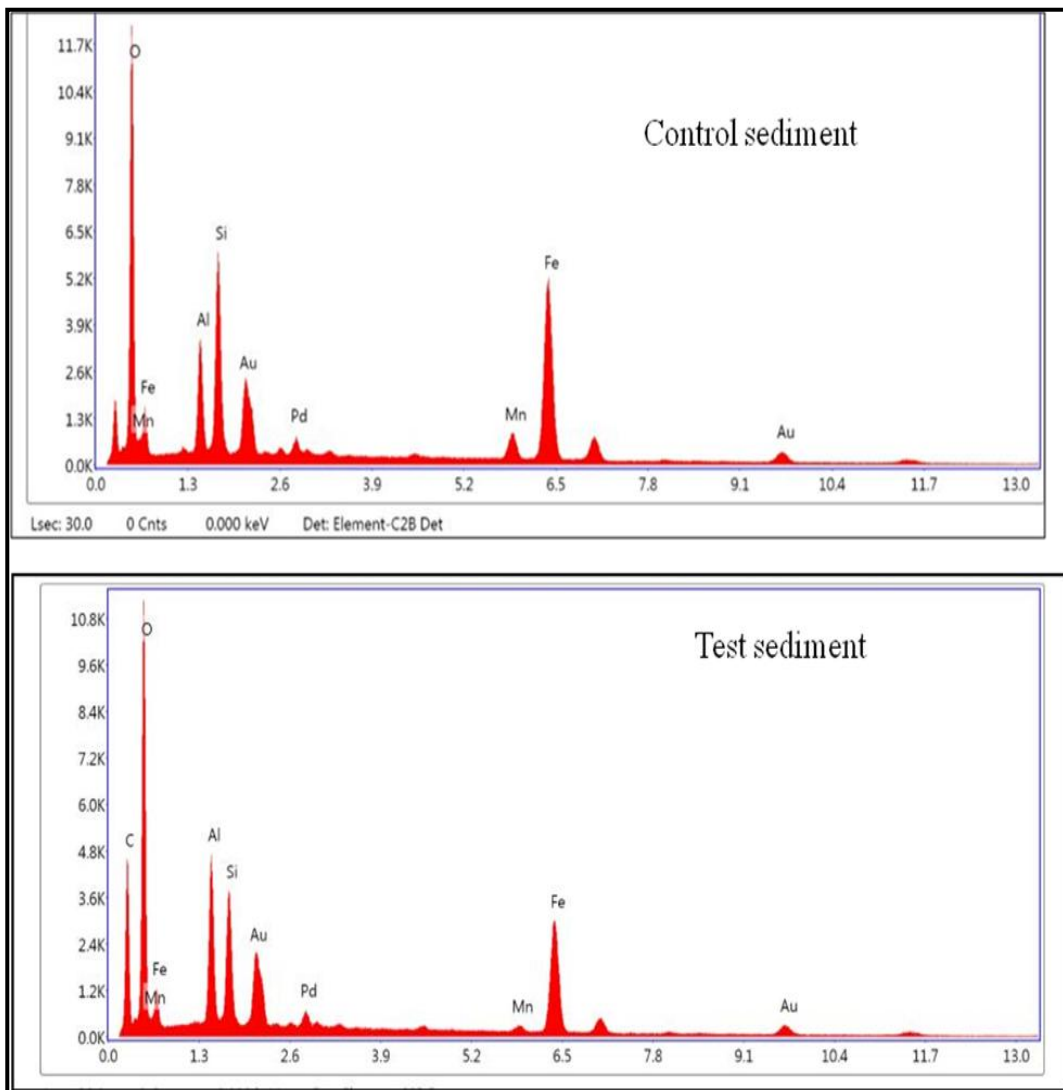


Figure. 3.43: EDS spectra of control sediment and test sediment

### 3.6.5. The FTIR spectra of control and test sediments

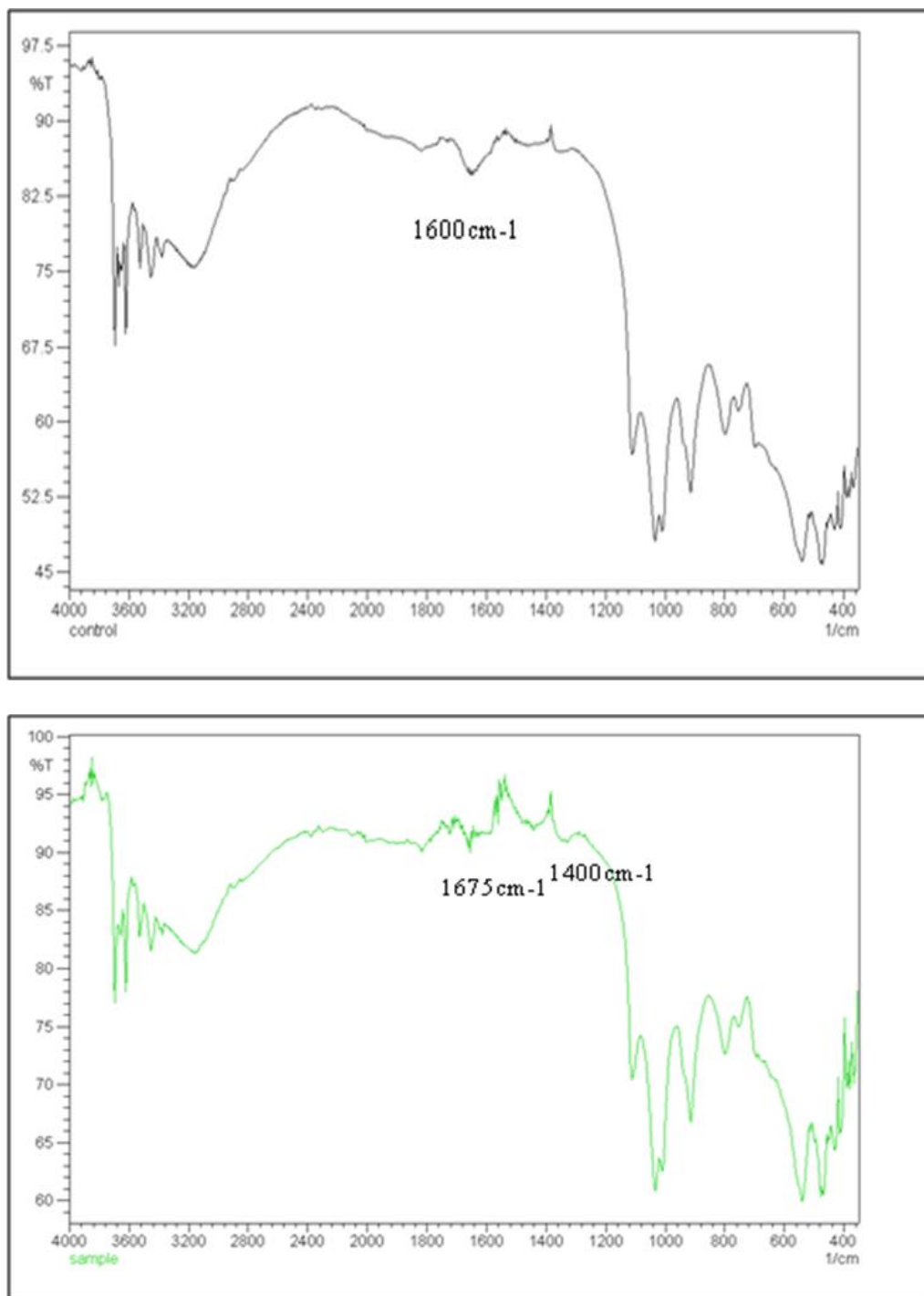


Figure.3.44: Spectra of control and test sediments

### 3.6.5.1 FTIR Spectra of sediments.

The IR spectra of soil sediment, soil sediment with yeast, were collected between the wave numbers of 4000 and 400 per cm and are shown in fig.3.44. sediment, and sediment with yeast spectra were studied.

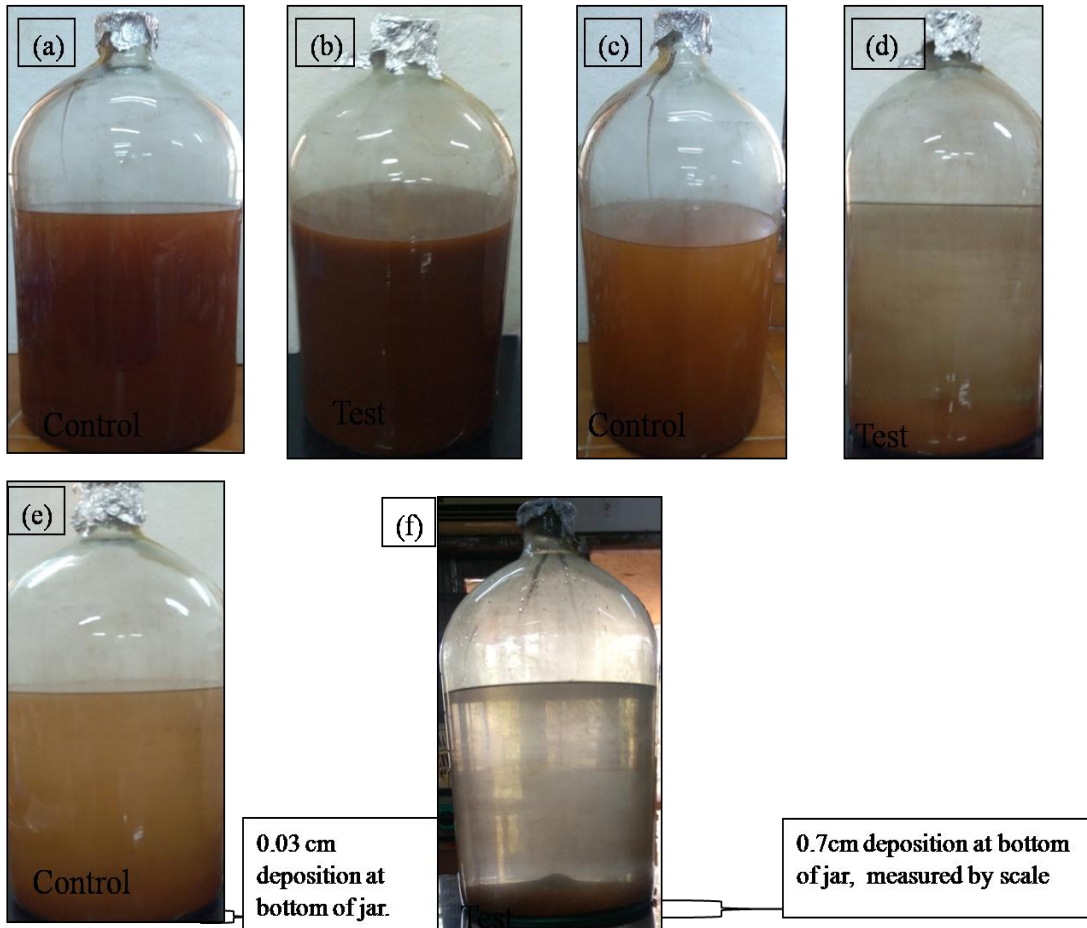
As seen in fig. A peak indicating P=O extending at 1,217 per cm occurs in the IR spectra of soil sediment. Peaks were identified in the soil and yeast mixed sediment IR graph ranging from 1,100 per cm to 500 per cm. In IR of yeast cells, a peak from 1,750 per cm appeared to be an aliphatic ketone.

The presence of amide is shown by the peak range 2800-3000  $\text{cm}^{-1}$ , which shows N-H stretching. The existence of an O-H bond in the peak range 2500-2000 suggests the presence of a carboxylic or hydroxyl group. Due to the presence of carbonyl chemical, the spectra exhibit a peak in the 1600-1750 range.  $1200\text{cm}^{-1}$  is the highest point. At  $1200\text{cm}^{-1}$ , the C-O group reaches its pinnacle. The presence of aromatic amines and nitro compounds is indicated by the presence of 525.12  $\text{cm}^{-1}$  peaks that show distinctive combinations of N=O and C-N stretching. Similar to the vibrating absorption of asymmetric and symmetric stretching in alkanes, alkenes, and alkynes, the peaks ranging between 1442.35 and 1056.50  $\text{cm}^{-1}$  were features of CC, C=C-H, and H-C-H, respectively. In general, the existence of organic molecules, compounds, and functional groups recorded for the sample demonstrates ion exchange capabilities that would confirm the attachment of clay colloids cations. Surface functional group dissociation aids dispersion by generating a negative or positive charge and/or supplying hydrophilic spots on a hydrophobic surface (Choudhary and Neogi 2017; Dupuy et al. 1997; Simate et al. 2012; Boehm 1994).

### 3.7. Biosedimentation study (20- 100 L)


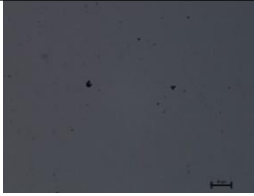
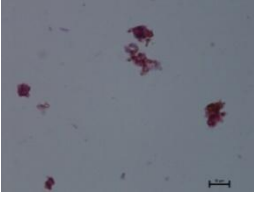
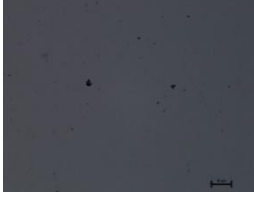
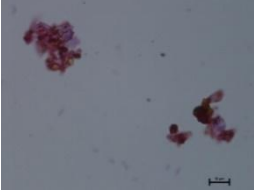
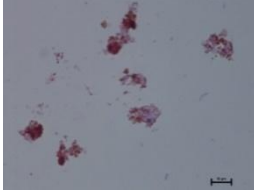
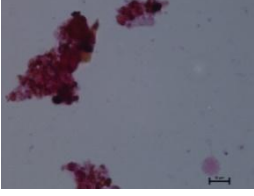
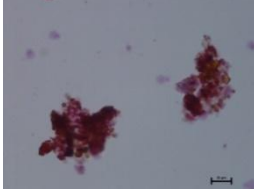
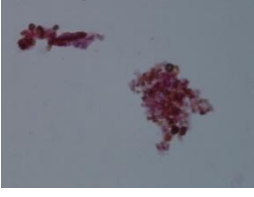

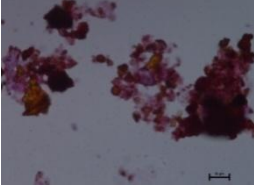
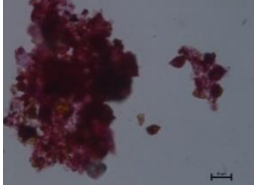
#### 3.7.1. Biosedimentation test in the 20 L bell jar

Biosedimentation efficiency from 1 L to 100 and results are follows



**Figure.3.45: Turbidity in Bell jar. Bell jar was kept on vibration free platform. (a) Control (b) Test at zero time. (c) Control (d) Test bell jar, after 10 minute of yeast addition in test jar) (e) Control (f) Test after 120 minute of yeast addition.**

**Table.3.21 Sediments morphology (at 100x magnification)**

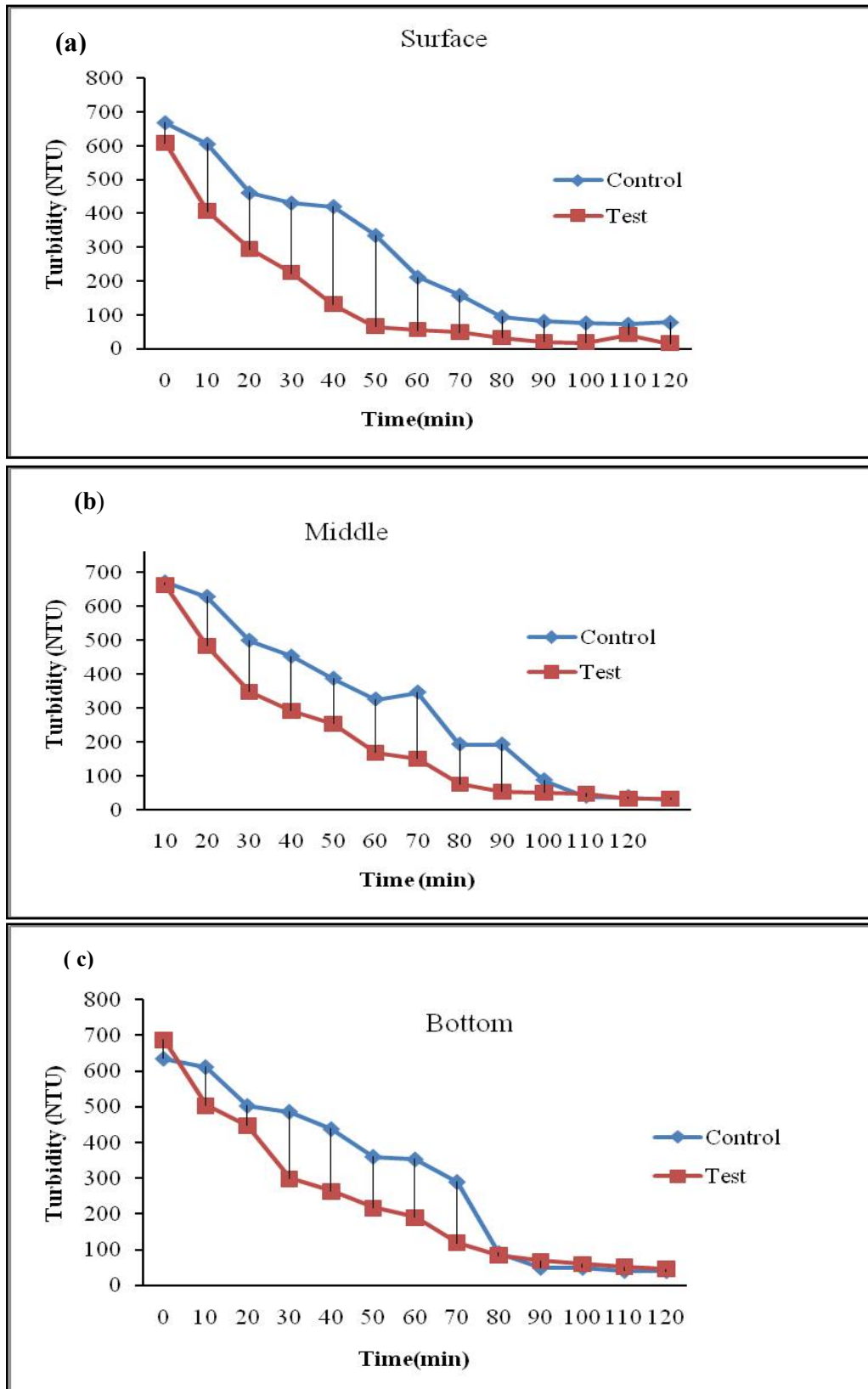
<b>Volumetric depth ml</b>	<b>Control</b>	<b>Test</b>
<b>14000</b>		
<b>12000</b>		
<b>10000</b>		
<b>8000</b>		
<b>6000</b>		
<b>4000</b>		

### **3.7.2.2. The biosedimentation kinetics of ( SMTTW) in 20 L Bell jar**

The biosedimentation kinetics of mine tailing water in 20 L at various depths revealed a decreasing tendency of turbidity at all depths. Turbidity in the test jar was lower than in the control jar after 60 min of yeast addition.

### **3.7.3. The biosedimentation kinetics of ( SMTTW) in 100 L tank**

In 100 L at various depths revealed a decreasing tendency of turbidity at all depths. Turbidity in the test jar was lower than in the control jar after 60 min. Moreover, sediment were compact as settled down rapidly after vigorous stirring, Fig3.47. Moreover, sediment compactness was high, as after restirring the water column turbidity reduced rapidly.



**Figure.3.46 Biosedimentation kinetics at different depths of bell jar (a) surface (b) middle (c) bottom of the Bell jar**



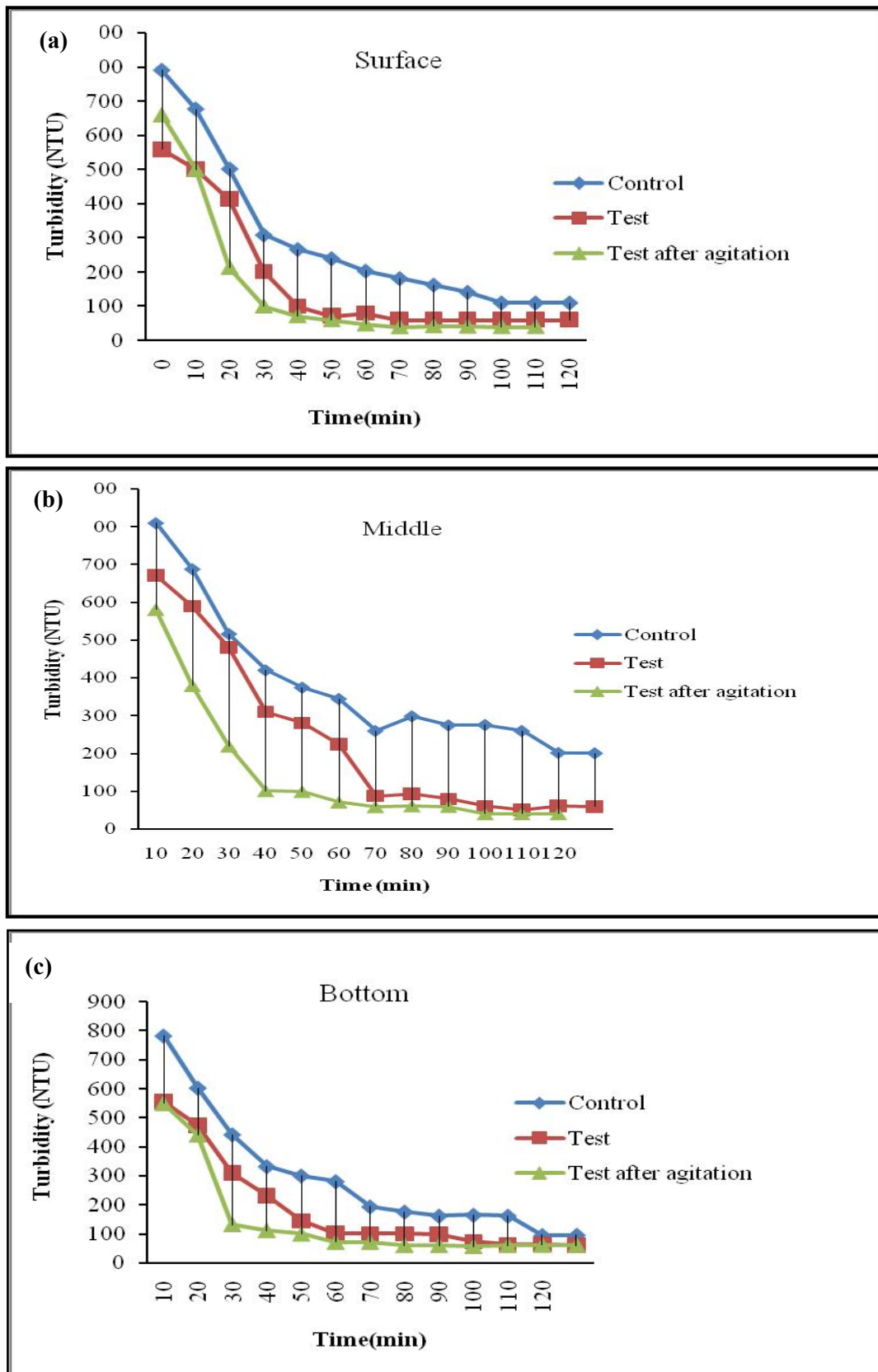
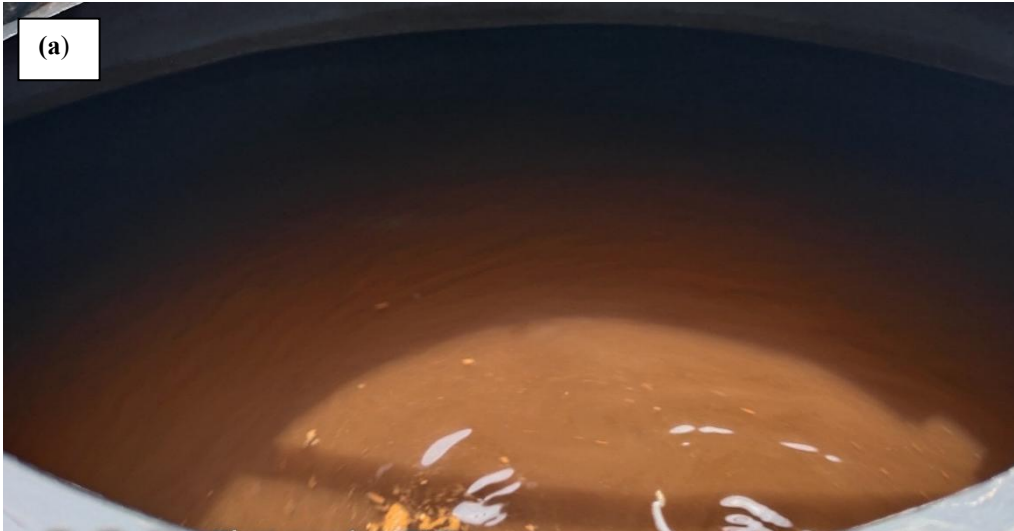


Figure.3.47: Biosedimentation kinetics at different depths of 100 L tank(a) surface(b) middle(c) bottom of the tank

### 3.8.1. Biosedimentation using selected cultures scale 500-1000 L

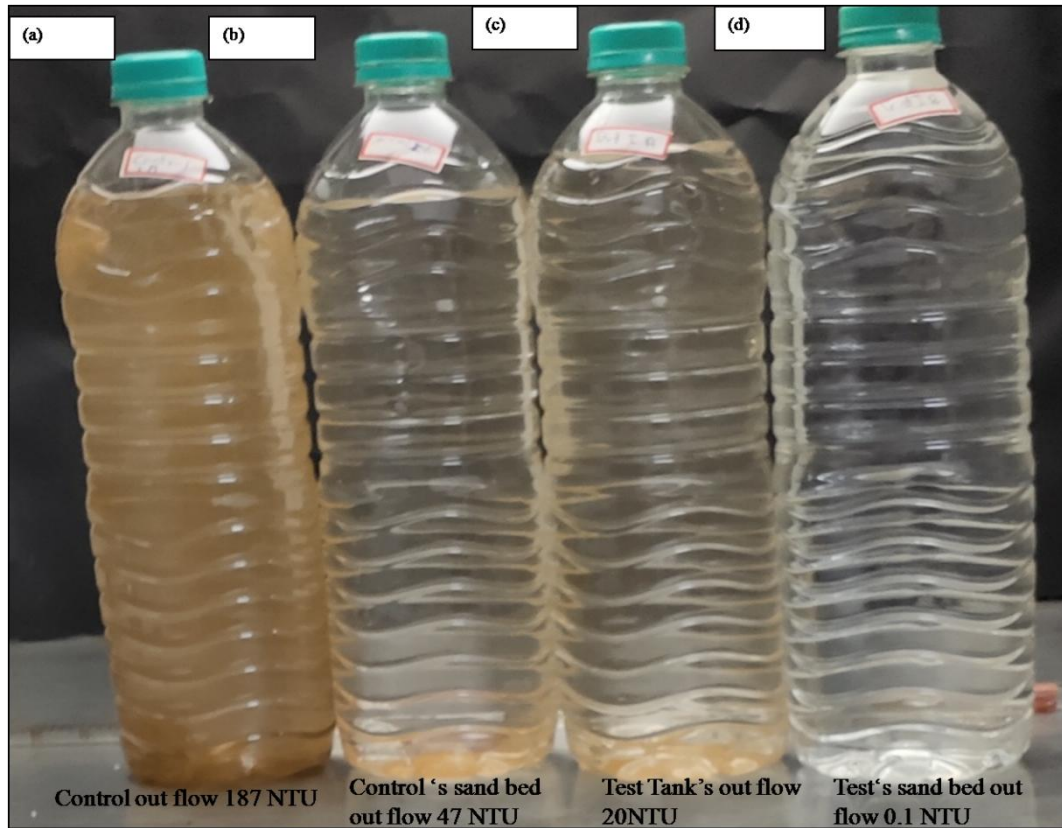


**Figure. 3.48: Visible turbidity in control tank, (a) after 0 min of stirring (b) after 1 hrs of stirring In the 1000L of tank.**

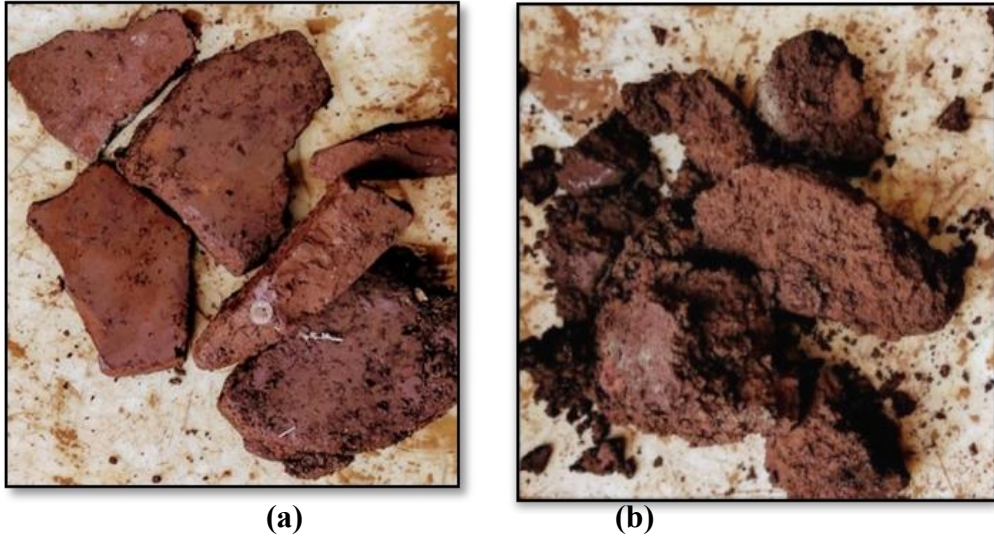


**Figure.3.49: Visible turbidity in reactor tank, (a) after 0 min of stirring (b) after 1 hrs of stirring in the 1000L of tank**

### 3.8.2. Out flow water collected from control tank, test tank and their respective sand bed



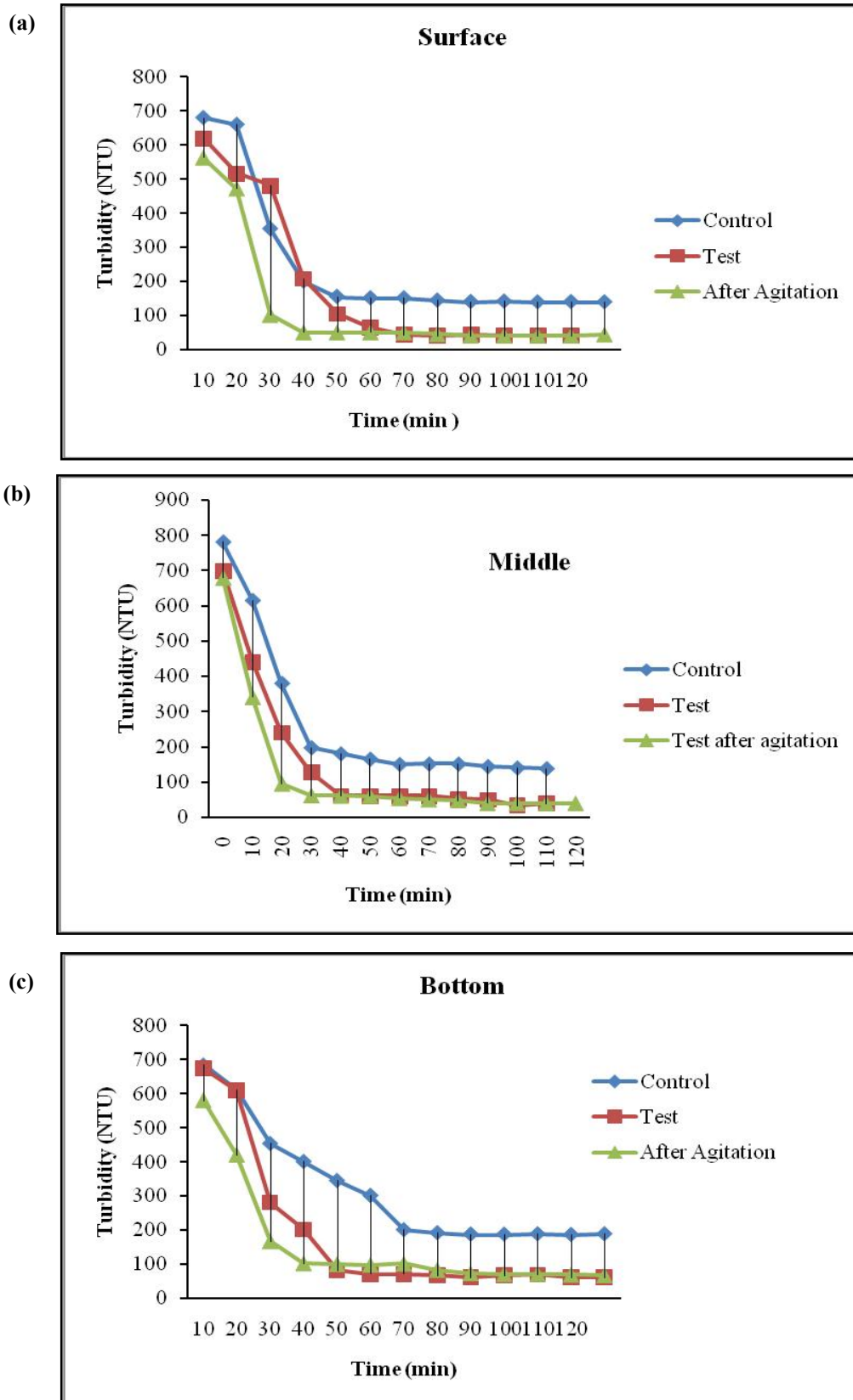
**Figure3.50: Turbid water, after 1 hr of stirring, collected from middle point of the tank height (a) Control tank outlet sample, (b) Control sand bed outlet sample (c) Reactor tank outlet sample (d) sand bed outlet sample connected to reactor tank.**



**Figure.3.51: (a) Settled Sediment from control tank showed clay like smooth texture. (b) Settled Sediment of reactor tank showed a coarsely aggregated outlook pattern of the texture.**

•  
**3.8.3. The biosedimentation kinetics of mine tailing water in 500L-1000L tank**

The results demonstrated that sedimentation occurring rapidly in the test reactor, or reactor in which yeast cells were inoculated. Moreover, sediment were compact as after restirring sediment did not broke and settled down rapidly. Results showed yeast mediated bioflocculation is scale independent.



**Figure.3.52: Biosedimentation kinetics in 500 Ltank(a)surface (b) Middle (Bottom)**

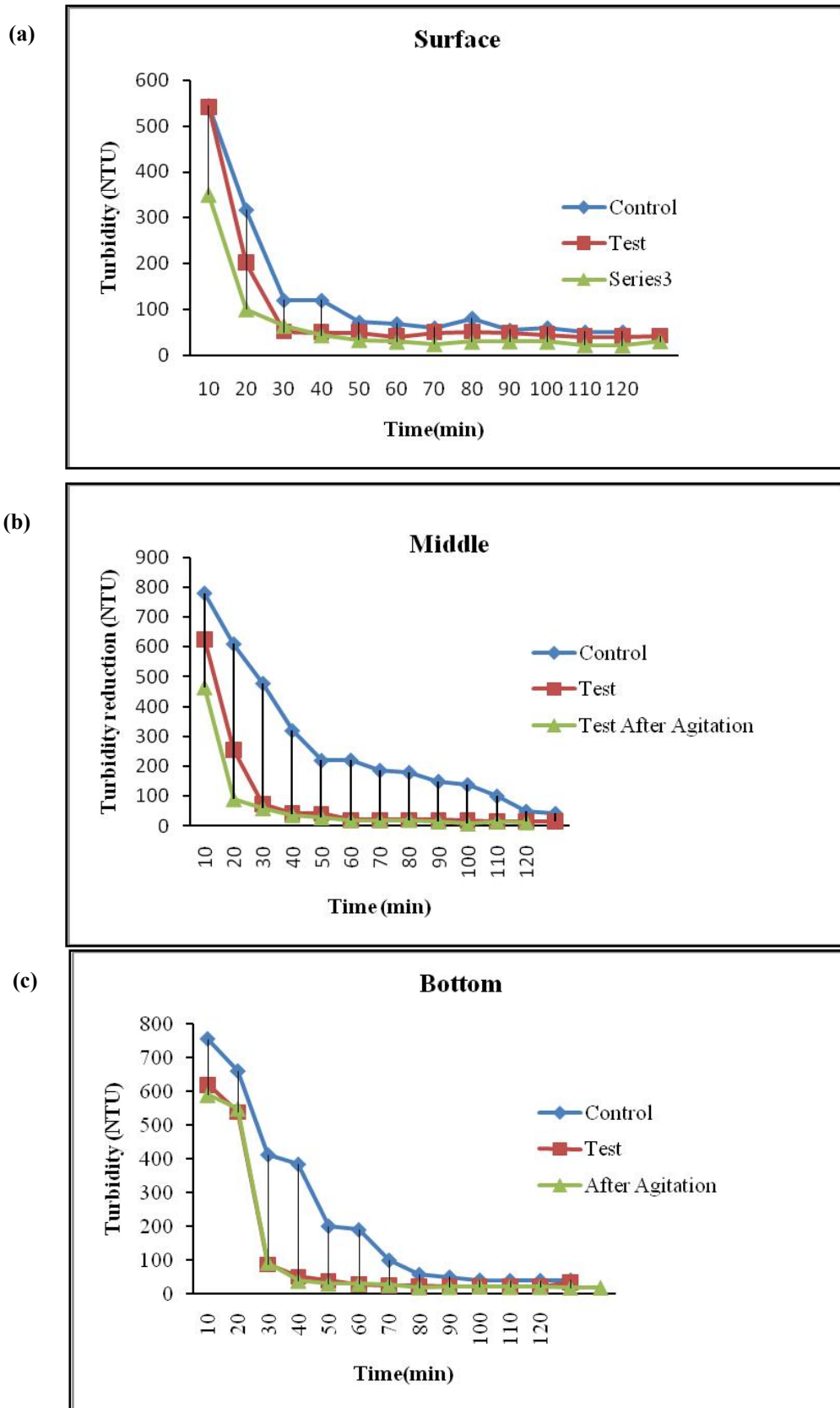
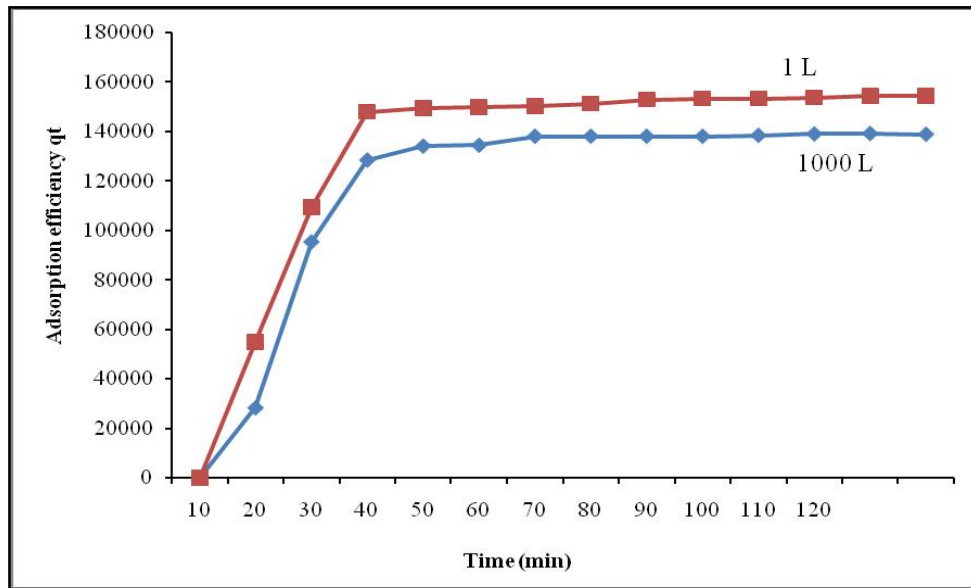


Figure.3.53: Biosedimentation kinetics in 1000L tank (a) surface (b) Middle (Bottom)

### 3.8.4. Adsorption efficiency of biosedimentation test



**Figure.3.54: Plot of adsorption efficiency of biosedimentation test using yeast in 1L and 1000 L mine tailing water**



### 3.8.5. Microbial study of effluent of tanks and its respective sand bed

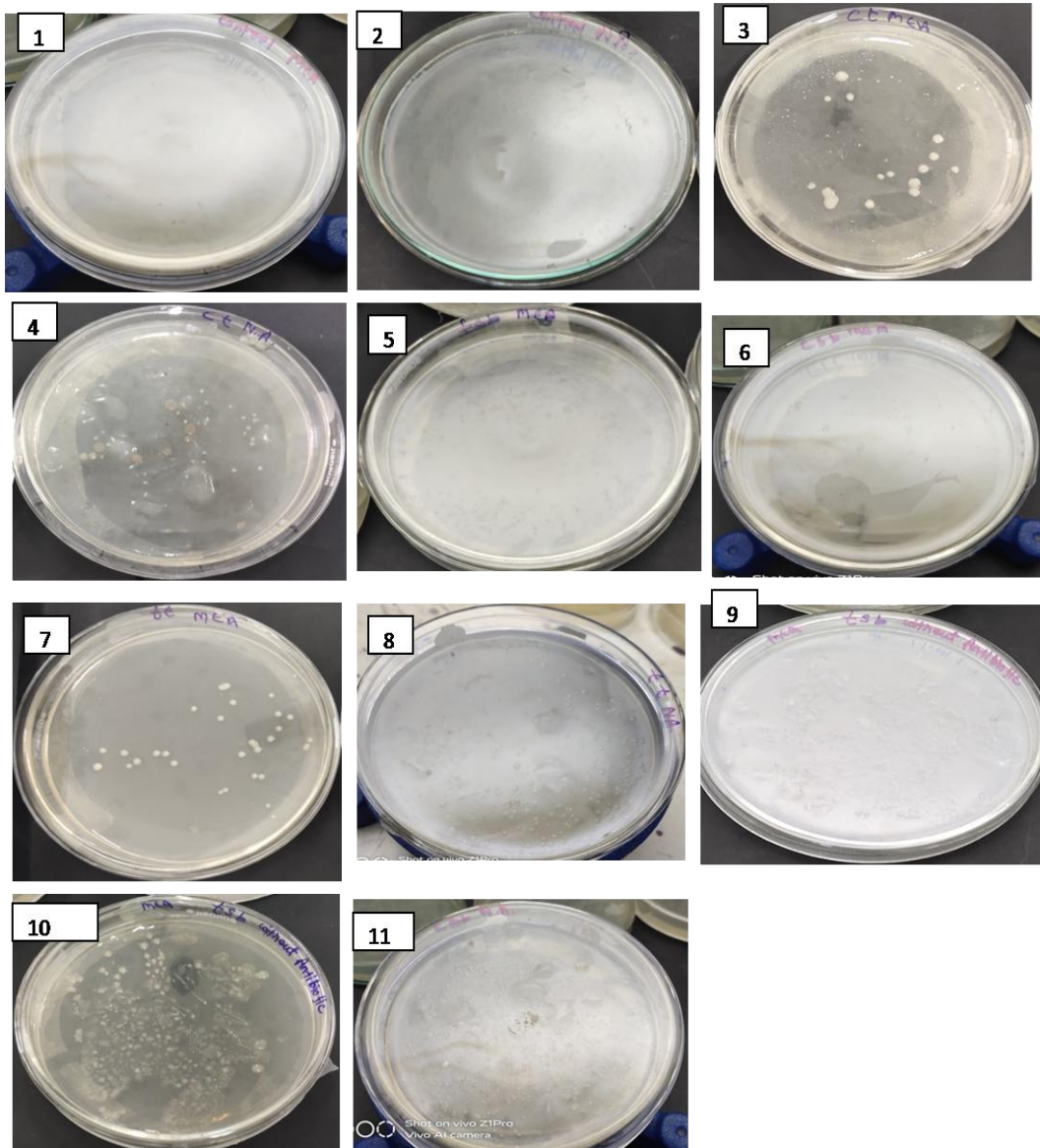


Figure.3. 55: Colonies from outflow of tanks

**Table.3.22 Colony characteristics**

<b>S. No.</b>	<b>Sample designation</b>	<b>Colony characterization</b>
1	Control MEA	No growth
2	Control NA	No growth
3	Control tank effluent Plated on mea (ct MEA)	Small white colony
4	Control tank effluent Plated on nutrient agar (ct NA)	Small colony
5	Test tank sand bed effluent Plated on MEA (ttsb MEA)	No growth
6	Control tank sand bed effluent Plated on MEA (ctsb MEA)	No growth
7	Control tank sand bed effluent Plated on nutrient agar (ctsb NA)	Small colony
8	Test tank effluent Plated on MEA (tt MEA)	Small white colony
9	Test tank effluent Plated on nutrient agar (tt NA)	Small colony
10	Test tank sand bed effluent Plated on MEA without antibiotics (ttsb MEA)	Small colony
11	Test tank sand bed effluent Plated on nutrient agar (ttsb NA)	Small colony

### 3.8.6. As a criterion for flocculant performance, the critical coagulant rate constant

The suggested criterion's validity is demonstrated by the results of a biosedimentation test in 1000L of turbid water in terms of residual turbidity of the flocculation process in order to meet the (WHO) turbidity standard for drinking water.

**c:** equals (5 NTU), which refers to the World Health Organization's (WHO) water turbidity standard (WHO 1996, 2006, 2008). The concentration of synthetic turbid water levels including: low (40 NTU), middle (70 NTU), and high (100 NTU) is the concentration of synthetic turbid water levels (780 NTU) **t:** equals the end of the chosen settling time (60 min) Substituting (c, c<sub>0</sub>, t) in Eq. (3) to obtain (k<sub>c</sub>)  $kC = 1/c - 1/c_0/t$  for second order reaction (Elimelech *et al* (1995)

#### Lower level (NTU<sup>-1</sup> min<sup>-1</sup>)

$$Kc = 1/0.1 - 1/40 / 60 \text{ (NTU}^{-1} \text{ min}^{-1}\text{)}$$

$$Kc = 0.1629 \text{ (NTU}^{-1} \text{ min}^{-1}\text{)}$$

#### Middle level

$$Kc = 1/0.1 - 1/70 / 60 \text{ NTU}^{-1} \text{ min}^{-1}$$

$$Kc = 0.1664 \text{ NTU}^{-1} \text{ min}^{-1}$$

#### High level

$$Kc = 1/0.1 - 1/780 / 60 \text{ NTU}^{-1} \text{ min}^{-1}$$

$$Kc = 0.1666 \text{ NTU}^{-1} \text{ min}^{-1}$$

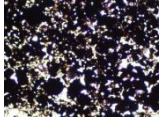
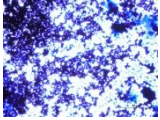
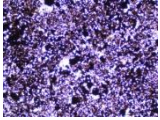
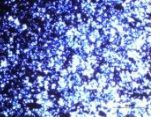
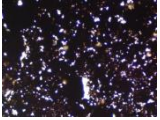
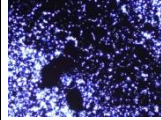
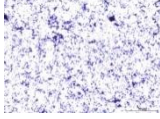
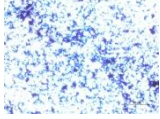
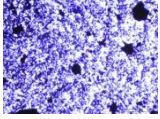
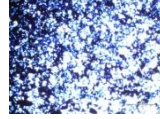
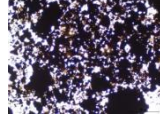
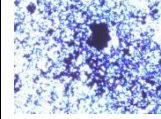
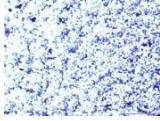

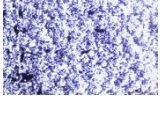
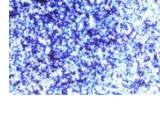
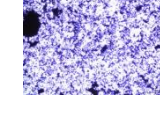
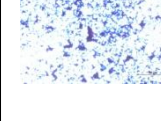
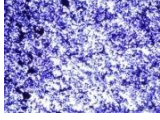

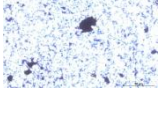

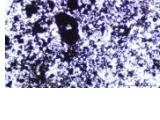

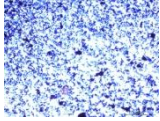
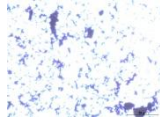
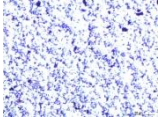
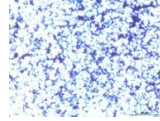
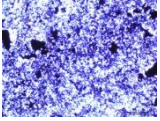
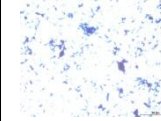
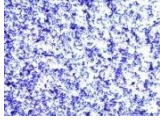



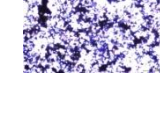

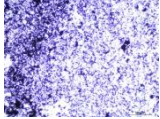
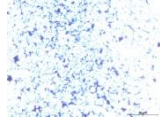
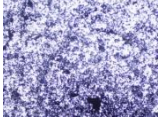
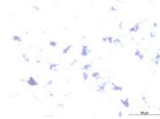
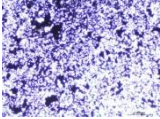

#### Rate constant for control tank and test tank at time 60 min

$$Crc = (1/c_t - 1/c_0)/t = (1/187 - 1/780)/60 = 0.000067 \text{ NTU}^{-1} \text{ min}^{-1}$$

$$Trc = (1/c_t - 1/c_0)/t = (1/40 - 1/780)/60 = 0.000396 \text{ NTU}^{-1} \text{ min}^{-1}$$

As a result, after employing yeast to remove turbidity in mine tailing water, the rate constant increased by more than 6 times. As a result, yeast may have a substantial role in reducing turbidity.

**Table.3.23 Morphology of sediment at different depth of the 1000L tank (at 100x magnification)**

Depth					
Surface (Control)	Surface Test	Mid (Control)	Mid Test	Bottom (Control)	Bottom Test
					
					
					
					
					
					
					

### 3.8.2.7. Sediment micromorphology

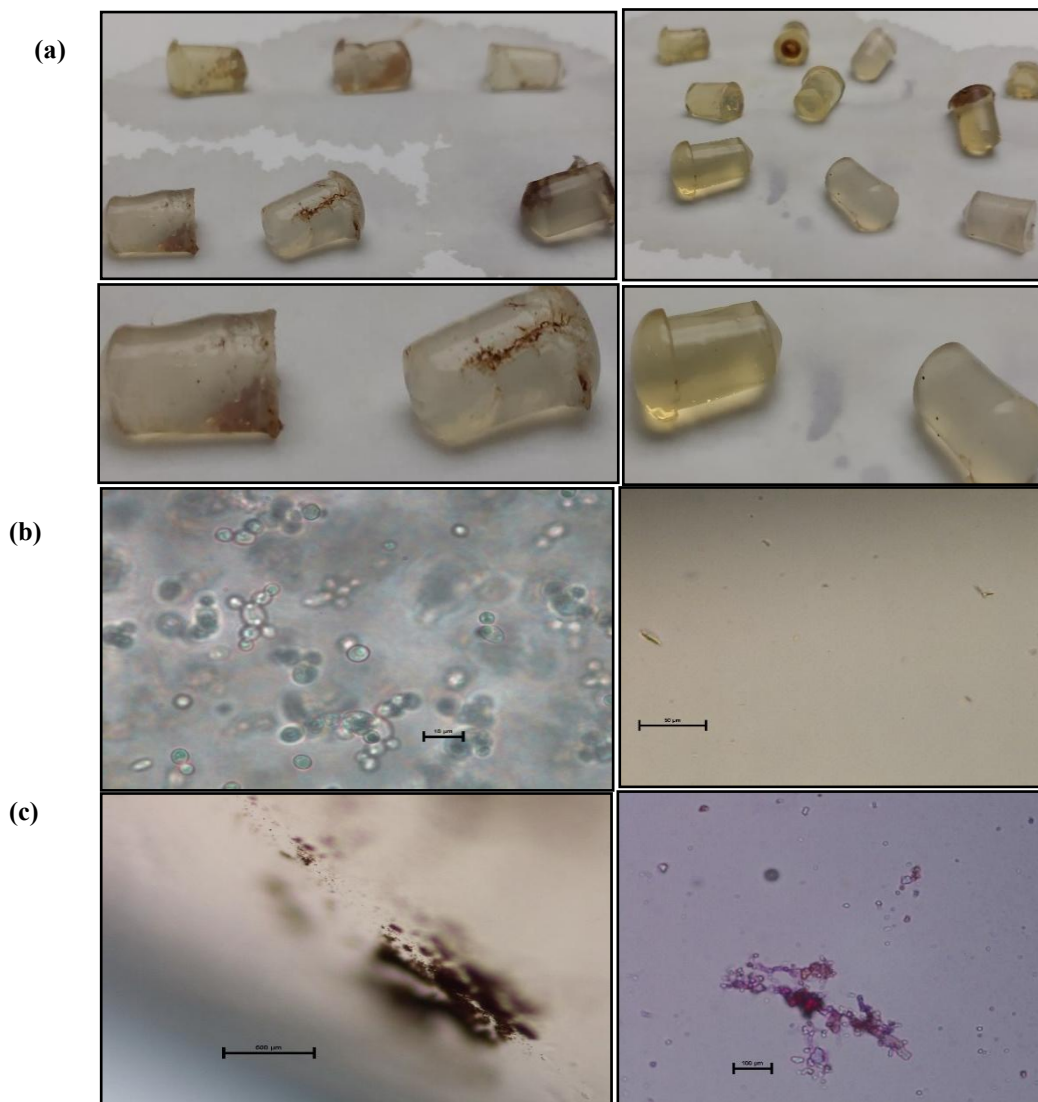
Micromorphology of the sediment revealed the concentration of suspended particles reducing with respect of time and it is reducing faster in the test reactor (Table3.27).

**Table.3.24 Water Parameters for 1000L biosedimentation test**

Parameter	Control tank	Control tank (sand bed)	Test tank	Test tank (sand bed)	Permissible Limit (WHO)
pH	6.4 ±0.05	6.40±.00441	6.40±.004	6.4±0.0044	6.5-8.5
Temp (°C)	29±1.06	29.1±.11	29±.88	29±0.99	-
TDS (ppt)	74.93±0	66.189±0	68.30±0	69.91±0	500
Conductivity $\mu$ s	136	127.6±	129	127.3	500
D.O.	6	6	6	6	No standard
Po <sub>4</sub> <sup>-3</sup> (mg/L)	0.75±0.306	0.75±0.614	0.25±0.16	-	0.1
So <sub>4</sub> <sup>-2</sup> (mg/L)	70.5±0.81	59.4285±.29	52±.25	-	400
No <sub>3</sub> <sup>-1</sup> (mg/L)	1.17±.0027	1.901±.003	0.6559±0.004	1.13±.0046	10
Alkalinity (mg/L)	4±0.44	4±0.333	3±0.5	2±0.5	200
Cl	6.025 0±.25	5.6 ±0.17	5.672±0.118	5.67 0±.27	≤200
Hardness (mg/L)	34±0.29	28±0.12	24±0.101	24±0.17	500
Na (mg/L)	6.484±0.5175	7.8480±.53	6	6.848	250
K (mg/L)	0.6432±.0.31	0.6432±.480	0.6432±.0.3	0.6432±.0.41	-
Fe (mg/L)	0.1928±0.01	0.0178±0.003	0.1427±0.008	0.0025±0.001	0.5
Mn (mg/L)	0.21145±0.01	0.00227±0.0002	0.12798±0.005	0.00224±0.00032	0.3

### 3.9.1. Observation of beads after and before immobilization and its turbidity removal efficiency

Beads from both the test and control beakers were thoroughly washed and examined under a stereomicroscope. Sediment were found in the test beads' surface interiors, but not in the control beads. In the test beaker, the drop in turbidity was not considerable. Furthermore, control beads turned yellowish in colour, whereas test beads stayed white and did not alter colour. Though sediment were visible in the test beads, they were not present in the control beads.



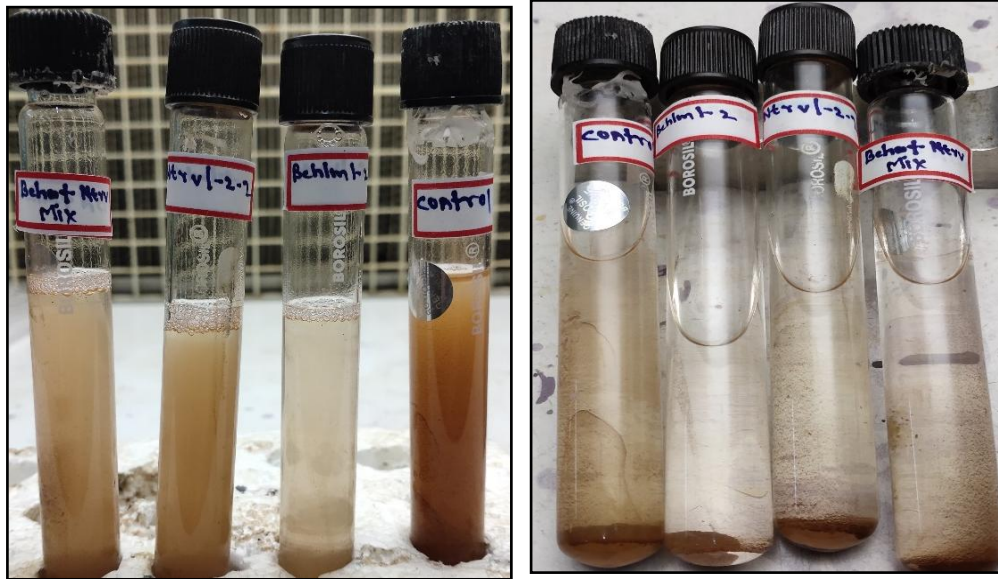
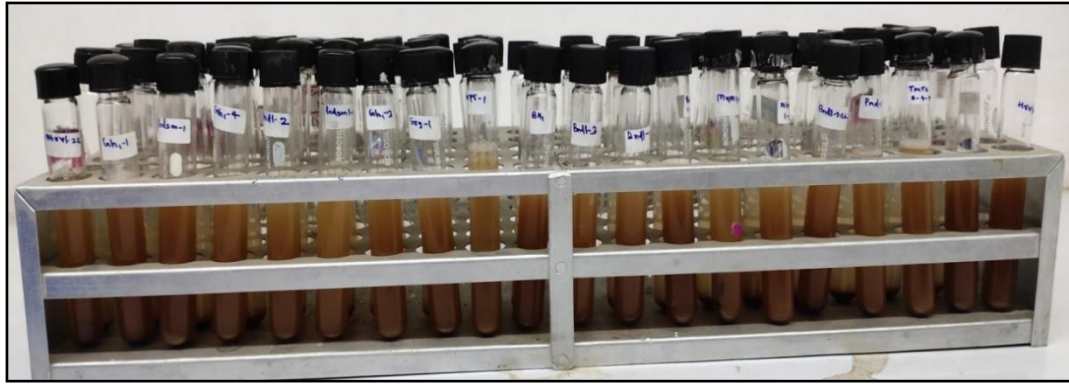
**Figure.3.56: (a) Beads with yeast (left) and without yeast (right), longitudinal section of beads with yeast revealed presence of yeast in the layer of agar (left). (b) Longitudinal section of beads without yeast (right). (c) Beads surfaces (left), longitudinal section with yeast, (right)**

### 3.10.1. Biosedimentation efficiency score for floc assay for yeast cultures combination study

The combination of the test strain and its own strain yielded the highest flocculation efficiency score, followed by Ntrv1-2-2.

**Table.3.25 Biosedimentation efficiency score of combinations of cultures**

<b>Strain designation</b>	<b>score</b>	<b>Strain designation</b>	<b>score</b>	<b>Strain designation</b>	<b>score</b>
Bchlm-2	100	Gh1-4	130	Srg-3-3	180
Pndsm1-2	180	Gh1-6	-	Krp-3	130
Srg-2-2	75	Krpr-2	-	S.C.-2	160
Bndl-1-1-1	50	Bndl-1-1-1	50	Srg-1	105
Pndsm-3	75	Bnd-2-2	50	Tmrs-2-2-3	105
Gh1-5	105	Crblm -1	-	Bndl-1	105
Ch1-1	105	Srg-3-3	-	Pnd-2	50
Krpr-1	75	Bnd-3	50	Bnd-2-2-1	130
Crblm-2	160	Tmrs-2-2-4	200	Ntrv1-2-2	50
Pndsm-1	130	Bchlm-1-2	280	Ntrv1-2	200
Pndsm2-2	130	Mym1-2	160	Hrvlm	130
Pndsm WA	130	Bchlm-1	160	Bh1	160
Gh1-1	130	Pnd-3	130	Control	25

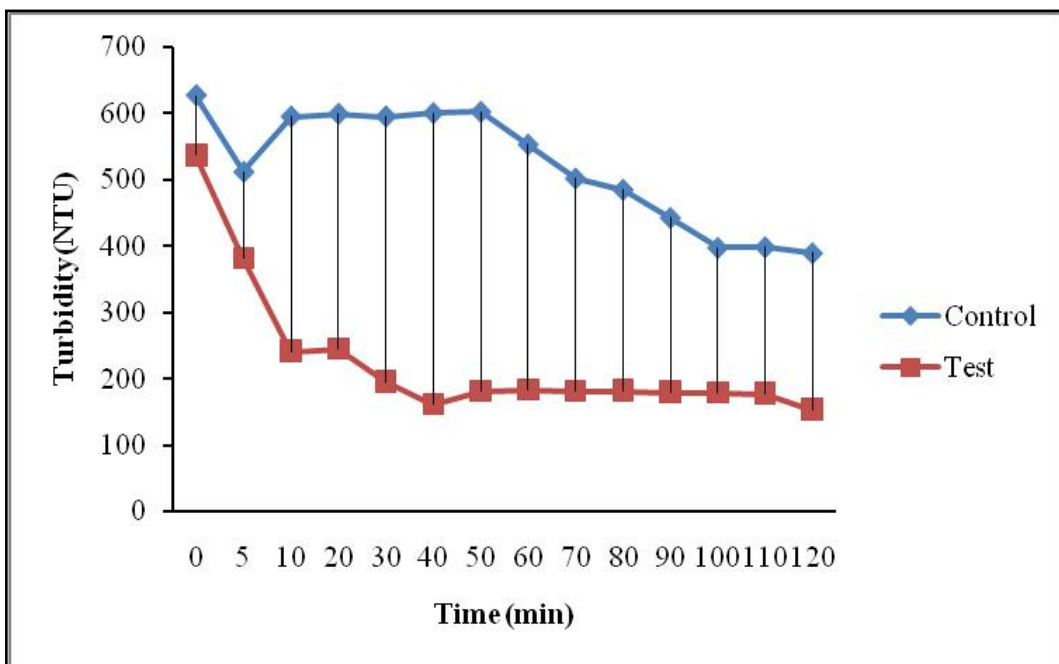
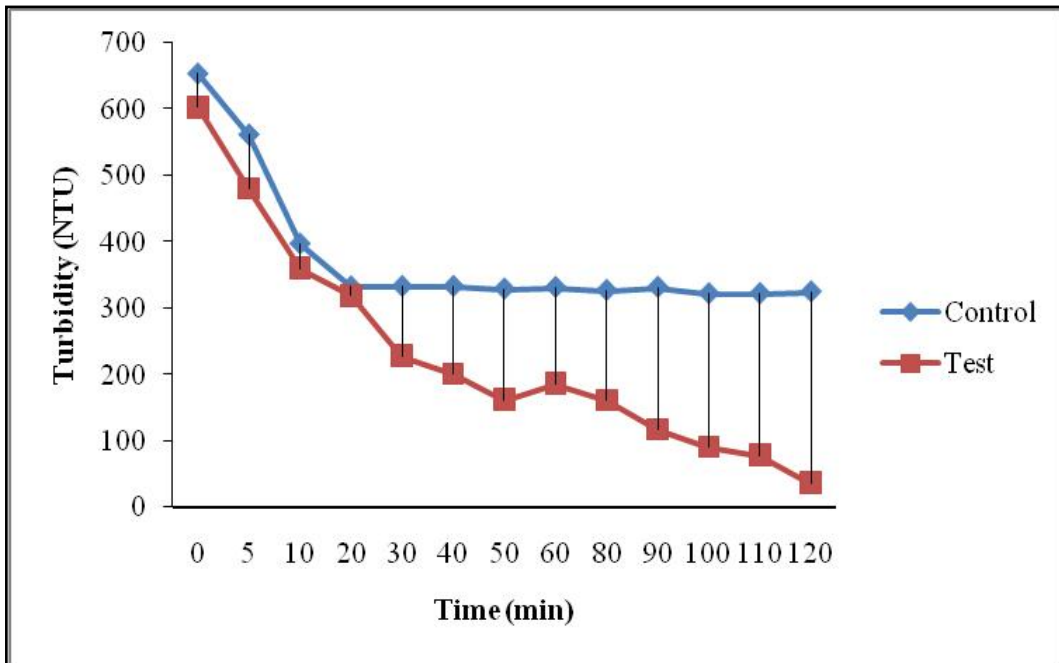


**Figure. 3.57: Biosedimentation assay for combinations study of yeasts cultures**

### **3.11. Biosedimentation study of natural mine tailing water**

The turbidity of mine tailing water remained high for a long time (Control cone), however yeast treatment reduced turbidity to less than 200 NTU after 60 minutes.(Fig3.58). In the control cone, the turbidity of natural mine tailing water stayed at 300 NTU till 60 minutes. Mine tailing turbidity remained same in the control cone for natural turbid water.

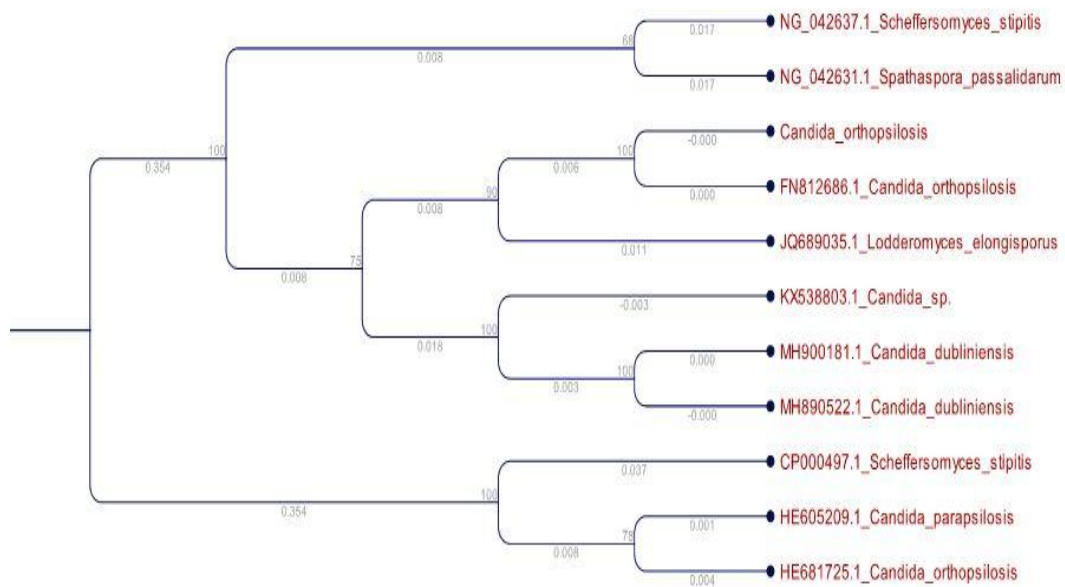




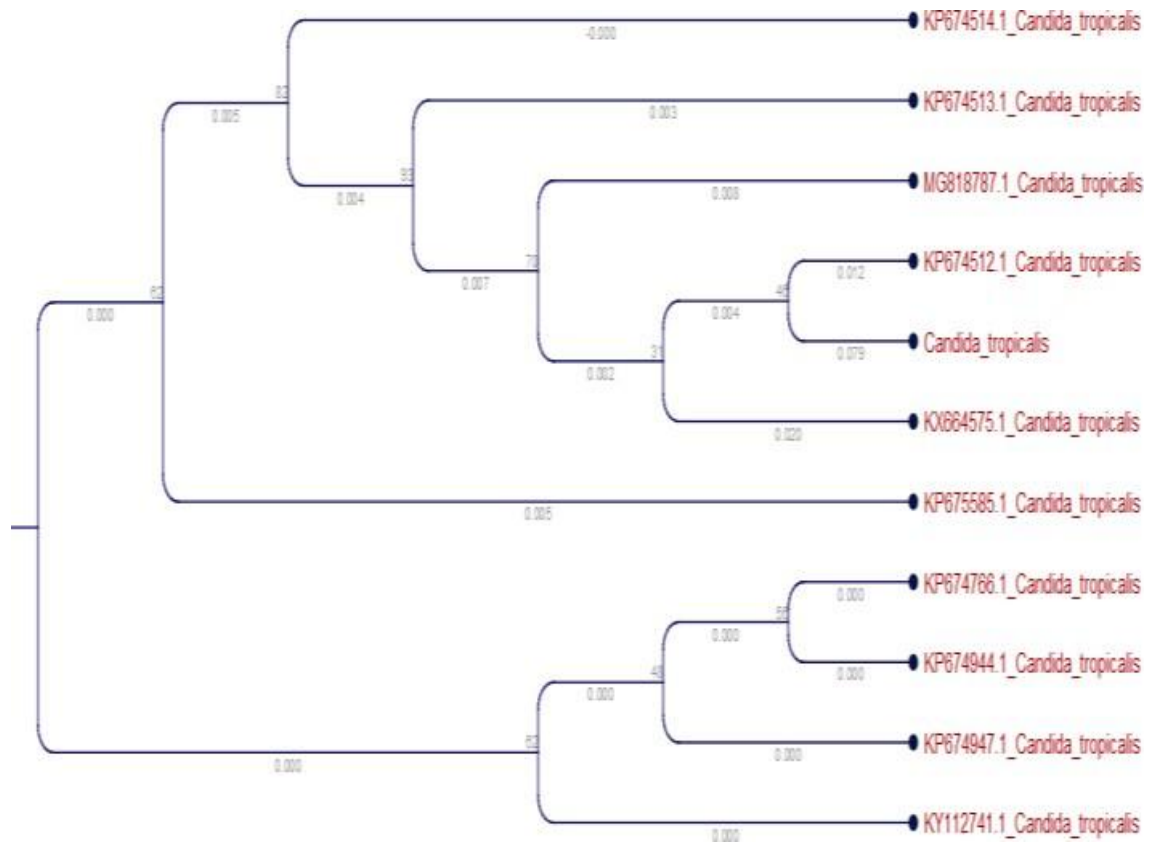
**Figure.3.58: Biosedimentation kinetics of natural mine tile water in Imhoff cone (a) at 900 ml of Imhoff cones depth (b) at 600 ml of Imhoff cones depth (All turbidity reduction values are the average of 9 turbidity reduction values**

### 3.12. Molecular Identification of promising culture

*Candida Orthopsilosis* was shown to be the most effective strain for biosedimentation after molecular identification. The cultures had a strong resemblance to *Candida Orthopsilosis* isolated from Ireland.



**Figure.3.59: Phylogenetic analysis of *Candida orthopsilosis*. Isolate showed 0.006 dissimilarity with type species *Candida orthopsilosis* ( FN812686.1)**



**Figure.3.60: Phylogenetic analysis of *Candida tropicalis*. Isolate showed 0.012 dissimilarity with type species *Candida tropicalis* (KP674512.1) obtained from gastric mucosa, China.**

# **CHAPTER IV**

## **DISCUSSION**

#### **4.1. Survey of freshwater habitats from mining and non-mining areas of Goa**

Goa showed a huge number of freshwater, bodies such as seasonal pond, during monsoon season. Most of the reservoir source was rainwater (e.g., cemented temple pond or tank and ground water (e.g., springs).

Since mining activities were not going on from the past twelve years, so the value of water parameters of these water bodies under study remained below the permissible level. Temperature varied from 24.5 to 30 °C and pH showed a variation from 6.3 to 7.8 variations in the studied seasonal water bodies.

Moreover, because of anthropogenic activity, slightly alkaline pH was observed at Netravali's freshwater bodies. The Electrical conductivity of water of these sampling sites under study varied from 40 to 140  $\mu\text{s}$  and remained almost in same range in monsoon and non monsoon season of every sites indicates the constant condition of the water body.

TDS were found to be within range of 15-130 ppm. Furthermore, Shirigao pond showed low TDS as no human activity was observed. Safa masjid pond showed highest TDS 140 ppm as the fauna population were high.

Renu et al., (2014) reported various physico-chemical parameters of the freshwater bodies of Goa. The pH in all locations was neutral to slightly acidic, varying within a small range of 6 to 7. And max pH limits of 7.9 was found during monsoon season.

The results validated Rump et al., (1988) assertion that any scenario when the water is neither excessively acidic nor very alkaline, it may be believed that pH is regulated by carbon dioxide, bicarbonate-carbonate system, which was first proposed by pioneer limnologist Hutchinson (1976).

#### **4.2. Isolation of natural aquatic yeast cultures**

This is first-time time that aquatic yeast were isolated successfully from mining areas of Goa by Kamat et al., (2010; 2013) using the membrane filter technique.

So after getting leads from the previous work freshwater yeast were isolated successfully. Among all the isolation plates, petriplates without antibiotics (MEA 2 %) showed growth of both yeast and bacteria. So it can concluded that antibiotics was effective for

the isolation of pure yeast cultures. This antibiotics faropanem was used first time by our group. Pure yeast culture growth was observed in the antibiotic-treated petri plates, and a similar pattern of growth was observed in all the triplicate plates. Further, the purified yeast's fourty cultures were maintained on MEA (2%) showed huge diversity of yeast.

Further, research can be done on freshwater yeast variety in Goa's water bodies, which revealed a great diversity of aquatic yeast and can be used for human welfare. All talukas showed huge yeast diversity. Bihcolim, and Ponda followed by Netravali waterfall, University seasonal pond, and then Sirigao showed yeast diversity.

Many different natural and artificial habitats, such as soil, freshwater and marine environments have yielded yeasts (Fell et al., 2001; Samarasinghe et al., 2021). Freshwater yeasts have not been much studied. However, no yeast was reported except some fungal species e.g., *Penicillium*, *Aspergillus*, *Rhizopus* in the lakes (Bandh et al., 2019). *Candida sp.* Have been isolated from resort salt water and the Shind river, Sonamar, and Kashmir (Bandh et al 2011; Wani et al., 2014). Vidya 2021 reported yeasts from the mangrove soil sediments of Kerala. In Kerala *Aspergillus* was reported as freshwater fungi isolated from Vembandu lake (Tomas et al., 2017).

Arora et al., 2021 isolated yeasts from coastal wetlands of Andaman Islands, India. *C. auris.* yeast isolated from mangroves soil sediments in Gujrat (Patel., 2020).

Most of the freshwater yeasts are underexplored in the world (Grossart et al., 2019). Previously pigmented yeasts was isolated from the river of mining area of Goa (Garg et al., 2011). So further studies need to be conducted to focus on isolation of freshwater yeasts from different freshwater bodies.

The highest number of aquatic yeasts are isolated and reported by Argentina Carlos et al., (2010)

#### **4.2.1. Characterization of yeasts cultures**

Yeast designated as Bchlm-1-2 had a high phosphate solubilization property, was robust to various drastic environmental conditions such as fluctuation in pH changes and high temperatures and grew on mine tailing media as black colonies. Colony colour changed from black to white on MEA media.

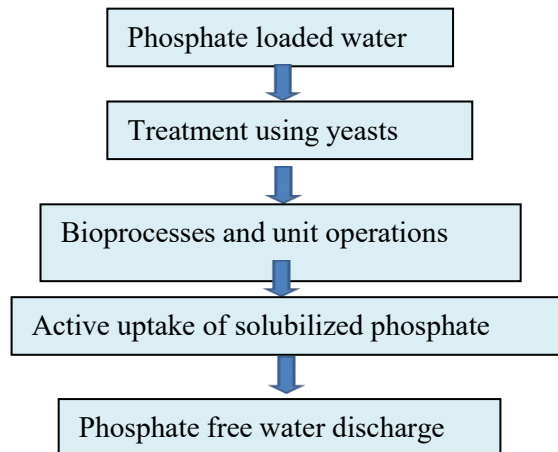
Besides from their biosedimentation activity, all strains can further investigated for their biotechnological applications.

Strains, such as Crmbm, Ntrvl-1-2, Ntrvl-2-2, and Srg-2-1, showed the presence of pseudo mycelium when cultivated on specified media, as demonstrated by various chemical tests.

Bchlm-1-2 tested positive for catalases. Bchlm-1-2 produced a positive test result with sucrose. All sugar tests were negative for Pnd-3, Pndsm-1, Hrvlm-1-2, Tmrs-2-3, and Tmrs-2-2-4, Ntrvl-2-3. With all sugars, Bh1 and Pndsm-2-2 were weak. Bndl-1-1-1, Pndsm-1-2, and srg-2-1 all tested positive for sucrose but negative for maltose. Ntrvl, Krpr-2, Pndsm-2-2 were positive in the presence of maltose sugar but negative in the presence of sucrose; moreover, Pnd-2 generated gas bubbles. Bndl-1-1, Gh1-5, showed positive with glucose and negative with all sugar.

Among all the studied cultures, seven freshwater cultures gave a positive test in phosphate solubilization and three cultures gave a negative results. Bchlm 1-2 was the best strain, displaying a very clear zone of phosphate solubilization around the colony on the media. The clean zone reached its maximum within 48 hours of strain inoculation and maintained a constant size until the seventh day of the inoculation.

Some soil fungal inoculants reported optimum phosphate solubilization effectiveness even after the 15<sup>th</sup> day of inoculation, while others did not (Elias F.,2016). Furthermore, microscopic analysis revealed that crystal deposition was present on test plates but not in control plates. Calcium oxlate, a reaction product, was predicted as the crystalline structure. Crystal concentrations were highest near the edges of colonies and least near the inner edges of petriplates.



**Figure.4.1: Schematic diagram for biotreatment of phosphate loaded water**

Among all yeasts strains, 12 strain were tentatively identified up to genus level using morphological and biochemical features. Almost 28% cultures belongs to *Candida sp.*, 2.5% belongs to *Rhodotorula sp.*

### **4.3. Screening of strain capable of biosedimentation of colloidal turbidity.**

**4.3.1.** Promising strain for biosedimentation of clay colloids was screened manually, using the biosedimentation efficiency score.

**4.3.2.** Cultures were classified and quantified for their biosedimentation property based on margin characteristic. Dimorphic behaviour was visible in micromorphology of stained unidentified yeast cultures. The fractal analysis of oligotrophic freshwater yeasts and its relationship to their bioflocculation capacity is little understood. For the first time, fractal analysis of colony edges of freshwater yeast isolated from Goa's several freshwater reservoirs has been performed.

Previously yeast colonies were characterized by its whole surface (Prado., et al 2014). Growth of yeast colony with respect to colony height studied earlier (Ravindranath., et al 1998). Papagianni et al., (2006) quantified the fractal nature of mycelial aggregation in *Aspergillus niger*. Morphological study of yeasts on the basis of the margins of colonies with image analysis was not much done so far.



**4.3.2.1.** The image analysis-based technology was used to quickly screen many yeast strains for application in wastewater treatment polluted by Goa's mining industry.

The fractal dimension of colony margins has been used to grade the yeast strains in terms of efficient biosedimentation.

Those yeast with a complex edge were examined for biosedimentation of colloidal turbidity in mine tailing water and were identified using ComiesJFrad.

Tables 3.21 and 3.23 in result section demonstrates that Bchlm-1-2 had the highest fractal dimension of its colony margins and created compact sediments with clay colloids, while Srg-3-3 did not have complex margins but generated compact sediments with clay colloids and showed the highest reduction in colloidal turbidity in NTU among the cultures with simple colony margins.

The Fractal analysis of yeast colony margins could be a beneficial tool for characterizing freshwater yeast strains and establishing a clear positive or negative link with their bioclarification capability using the fractal dimensions.

Because the same procedure and results were repeated with five best strains, it is possible to conclude that the sediments formed by different strains in simulated turbid mine water were first characterized by a mathematical tool to analyse their fractal dimension to correlate with its biosedimentation capacity, and it was discovered that those strains with complex margins were forming sediments with a high fractal dimension.

Apart from mathematical analysis, the most notable strains were visually examined for floc compactness. All cultures with complex margins developed compact sediments eg. Bchlm-1-2, formed compact sediments. The fractal dimension of colony margins and their compact floc formation feature were compared for the first-time using fractal analysis of isolate colony margins.

In Fig. 3.19, a snapshot of colony borders of freshwater yeast cultures is shown, and their fractality indices are compared in terms of growth functions and placement

New technique enables a simple study approach. Yeasts develop in predictable patterns, and even after streaking, they do not distribute evenly throughout the plates. It was

simple to watch, photograph, and generate a random data set from a predetermined nested pattern of established colonies on solid media. Furthermore, the square pattern has allowed repeated colonies to expand in a linear fashion.

**4.3.2.1.** Colony margin growth was monitored and defined using temporal and spatial changes. Because the colonies were inoculated on the same day and in the same conditions, growth and morphological behaviour for each colony could be easily compared.

Three nested colonies grew in the same pattern, increasing in complexity in their margins over time, but their fractality indices differed depending on their position.

The underlying cause could be colony heterogeneity or a genetic switch to live in nutrient-depleted media. With increasing margin complexity, we also see an increase in the fractality index. Because these traits may be ruled by chemo-taxis interaction or gene expressions, more research into colony margins and their interactions is needed to determine the reason of variance in growth patterns.

Ruusuvuori et al., (2014) used an image-based technique to conduct the qualitative analysis of yeast colony margins. Two-dimensional top-down binary pictures were used to examine the morphology of *Saccharomyces cerevisiae* colonies (Gontar., 2018).

Present study claims usefulness of image based technique for the screening of most promising strain designated as Bicholim1-2 shown to be the most promising strain for biosedimentation of clay colloids.

#### **4.4.1. Study of potential test yeast morphology due to change in medium:**

The test strain was tested for a change features when grown in nutrient rich medium (Commercially available MEA), which is usually rich in carbon source for microbial growth and simulated mine water to verify its wild behaviour.

Whereas test strain was cultivated on two separate medium, significant morphological changes were observed. The colour of yeast colonies changed after strain transfer to the changed media could be due to phenotypic changes in response to nutritional stress conditions. After 7 days of inoculation strain which were grown on MEA media remained white shiny smooth and gelatinous surface with complex margins while strain

which were grown on modified media turned black in colour with shiny smooth and gelatinous surface with complex margins. Strain grown on modified medium were spread plated on fresh MEA and the modified medium and observed that strain retained black colour on the modified medium while it regains its white color on MEA medium (Fig.3.18). Bicholim1-2 demonstrated remarkable resistance in the modified medium which was simulated to the mine environment.

#### **4.4.2. Characterization of soil sediment and yeast using SEM and FTIR**

The isolate was found to have a large surface area and surface charge. Catalases are reduced. Phosphate solubilization efficiency was high. Strain can also reduce Fe and Mn in the mine tailing water.

The results of the FTIR analysis used, to examine the functional groups present in the soil sediment and yeast mixed soil sediment. The functional Groups of the yeast mixed sediment, and control sediment did not show much variation. Little displacement was observed in the mixed sample.

Both sample had comparable spectra. The OH functional group and the O-H stretching of the polymeric molecules were attributed to the prominent peak seen at 3400-3294  $\text{cm}^{-1}$ . Additionally, the peak between 2929 and 2928  $\text{cm}^{-1}$  was identified as C-H groups. Around 1660–1657  $\text{cm}^{-1}$  was the typical stretching peak of the carboxylic COO-double bonds of deprotonated carboxylate functional groups. Most of the peaks of studied samples showed peak at 2900  $\text{cm}^{-1}$ .

**4.4.3.** The extent of the dye's adsorption (Acian blue) indicates the strength of the negative charge on the cell surface. The presence of hydrophilic groups at the surface was primarily indicated by yeast's relative hydrophobicity. According to Laurent et al. (2009), hydrophilic molecules are often polar or charged while hydrophobic molecules are non-polar. A better availability of polar/charged groups like carboxyls at the yeast cell surface can be correlated with a lower relative hydrophobicity and a larger negative surface charge. The greatest concentration of negative and/or polar sites on the yeast surface probably corresponds to the greatest concentration of particle fixation sites. This results is matching with work of Kordialik (2008) concluded Copper and Lead removal

from yeast cells is less effective when their relative hydrophobicity is higher and their surface charge is lower.

Many researcher have reported that a hydrophobicity percentage of between 30 and 40% could ability to interact with epithelial cells and at least perform transient adhesion (Abdulla et al. 2014; Sidira et al. 2015; Ilavenil et al. 2016). In this study, the selected strain showed less hydrophobicity compared to baker yeast so it can be predicted the test strain is less virulent.

#### **4.5. Assessment of the quality of drinking water in mining areas of Goa**

Physicochemical analysis of different water samples were confirmed to be within acceptable limits. Furthermore, results of all parameters were found to be nearly identical in monsoon and non-monsoon seasons. Because mining did not take place on a large scale, turbidity was not as high as it was previously reported. Even though turbidity was found to be above the permissible levels in all water samples sampled from various locations.

The temperature of all the freshwater bodies under study closely matched with the general trend of ambient temperature and stayed consistently. In general, similar range of temperature was also reported by Sawaikar (2016), Vikrant (2001), Renu (2014).

There were no seasonal differences in the values of water parameters of the samples. Renu et al., (2014) reported nitrate concentration in Carambolim lake, Pilar lake was 0.38 and 1.31m respectively. And Phosphate concentrations reported 0.70 and 0.570 m respectively. Present study observed concentrations of Phosphate and nitrate ranged from 0.01 to 0.02 m and 0.28 to 0.85 respectively in monsoon sesason. The observations were similar to those by Sawaikar et al., (2016) and Vikrant et al., (2001). Water samples from Arvalem in monsoon showed strong correlation of turbidity with Nitrate, Phosphate, Sulphate. While Morlem pond showed strong correlation between turbidity and nitrate. Moreover, Morlem tank showed strong co- relation coefficient between turbidity and Phosphate. Bicholim tank showed modrate co-relation coefficient of turbidity with Nitrate, Phosphate, Sulphate non monsoon.

So, it is concluded that the low concentration of Nitrate and Phosphate indicate in the freshwater bodies were oligotrophic in nature.

#### **4.6.1. Lab scale bioflocculation using natural yeast cultures at upto scale 1-100 liters.**

The rate of biosedimentation of suspended particles in the test jar was higher than the control jar. The suspended particles settled to the bottom of the bell jar, and there was some wall deposition on the wall of the test bell jar. The water composition was essentially identical in slides taken from different depths of the bell jar and tank after 2hr of yeast inoculation and agitation. At a depth of 2 L(11cm), near bottom of the bell jar yeast cell hyphae were discovered with bacterial cells. Single coccid and yeast cells were found at the top of the jar at a depth of 14L (volumetric depth ) from the bell jar's bottom.

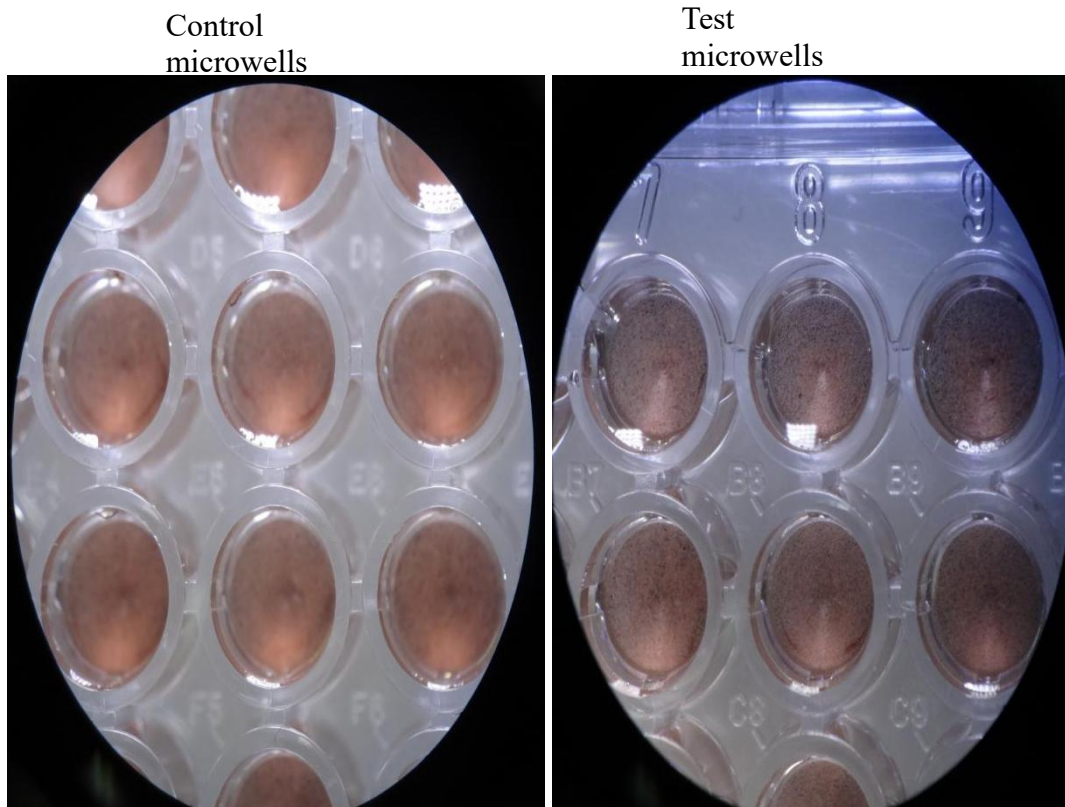
Furthermore, water samples collected at various depths in the tank and plated on MEA media revealed the same microbial makeup and colony size. Within the same time period, triplicates of withdrawn water samples exhibited identical turbidity reduction values.

The control tank, in which yeasts suspensions were not added, the turbidity was found to decrease slowly, and sedimentation also occurred slowly, whereas the turbidity in the test tank was found to decrease swiftly.

The treated water was decanted and then passed through a sand bed filter in order to get 100 percent turbidity-free water in a short period of time.

**4.6.2.** After 30 minutes of yeast addition to turbid water, the decanted of water can be run through a sand bed filter to obtain clean water with low turbidity, according to the results.

**4.6.3.** To enable clay colloids and microbial surface interaction, a mechanical stirrer is necessary.



**Figure.4.2: SMTTW in the micro well. (a) Control well showing turbid mine water, (b) Test well showing granular sediments, settled in the bottom due to binding of clay colloids with yeast.**

The benefit of yeast treatment is that it cuts down on the time, takes to settle the suspended particles and can be highly successful in removing the small particles that are difficult to settle out.

Furthermore, significant in turbidity reduction was seen in natural mine tailing water due to the addition of yeast.

While the turbidity remained high in natural mine tailing water compared with simulated mine turbid water in control reactor. The reason could be because natural mine tailing water contained clay colloids with proper water saturated.

Moreover, the sediment deposition was observed 1.5 and 4.5 ml respectively in control and test cone respectively during biosedimentation test of the natural mine tailing water.

So, it can be confirmed, and concluded that yeast mediated biosedimentation occurs in mine tailing turbid water as the sediment deposition in test cone is 3 times more than control cone in which the particles settled naturally.

The compactness of sediments formed were high, due to binding of yeast and clay particles. It can be observed from the biosedimentation kinetics graph that the sediments were compact since the turbidity quickly decreased when the test water column was restirred. And no breakage of biosediments were observed. Sediments settled fast after stirring the water. Moreover, Yu et al. 2011 studied the breakage of sediments formed by alum salt and concluded that sediments can break and regrow when the water column disturbed.

Leila Mosleh et al., (2014) reported a reduction in turbidity of water from 500 NTU to 85 NTU when using alum salt (0.005gm/L). Moreover, Aslani et al., 2012 reported a reduction of turbidity of water from 500 NTU to 10 NTU using alum salt (0.045gm/L), for a reduction in turbidity in an efficient time the dose of coagulant should be increased. While the present study can cause a reduction in turbidity of water using little amount of biomass (0.005 gm/L) from 780-800 NTU to 20-40 NTU within 1 hour, without breaking and remixing of the biosediments.

The findings could provide an insight into how yeast culture interacts with wastewater systems to eliminate colloidal turbidity. Yeasts can also be used as a coagulant because it is a good supply of fungal by products such amylase, chitin, and lactic acid (Oliveira et al., 2015). In comparison to bacteria, yeast can be also examined for its tolerance to inhibitory compounds.

#### **4.7. Standardization of bioflocculation process for field level application scale 500 litres-1000 litres.**

For the first time in the world yeasts isolated from freshwater habitats were used to study the biosedimentation of the clay colloids of the SMTTW. Mining activity contributes to water pollution, resulting in severely turbid water with SPM ranging from 4 to 22 gm per liter (AEQM, 1997) due to runoff of mine tailings soil during the southwest monsoon season, in Goa as we discussed.

Because freshwater yeasts have potent cell wall polysaccharides capable of attaching to inorganic colloids, the main goal of our research was to determine the practicality and bioefficacy of using them as active, live microbial cellular coagulants. After successfully screening, the most promising yeast strains (for biosedimentation of clay colloids), was investigated under various conditions.

For an efficient biosedimentation of clay colloids, all parameters were tuned, including SMTTW pH, system temperature, biomass of bioflocculant, age of yeast cultures, and coefficient of mine tailing water and yeast biomass. The stoichiometric connection between a flocculating yeast strain and a turbidogenic suspended material was investigated.

pH values were remained between 6-7 and temperature observed between 24-29 degree centigrade (ambient) for efficient sedimentation (throughout the experiment). Using an 0.005 gm/L (dry weight) of yeast suspension showed compact floc formation with clay and an efficient turbidity reduction.

Turbidity removal is maximum at an optimized dose of yeast suspension and when the dose is increased turbidity tends to reappear due increase number of yeast cells. 4<sup>th</sup> and 5<sup>th</sup> day old yeast proved more efficient as it coincided with the log phase of the selected strain. Bioflocculation kinetics were found to be greater after almost 40-60 min in the test reactors.

Hence pH and age of culture required to achieve a high reduction in turbidity ,as found 6-7 and 4<sup>th</sup>-5<sup>th</sup> day old respectively, using the scoring scheme (Table.2.3). When the pH was lower than 6-7, the water column was less clear, and when the pH was greater, the water became opaque (Fig. 3.34). Furthermore, wall deposition was higher in the column that was inoculated with 6<sup>th</sup> and 7<sup>th</sup> days old yeast biomass.

**4.7.1.** To keep the concentration of yeast biomass in SMTTW biosedimentation at a minimum, a 1:9 ratio of yeast suspension to mine tailing water was used.

**4.7.2.** Turbidity reductions in mine tailing water up to 1 L indicated a considerable reduction in turbidity within an hour after the yeast was added. The strain reduced the



turbidity from 780 to 73 NTU in one hour, demonstrating a 90% reduction in turbidity from initial. and 61% when compared to control under laboratory conditions.

**4.7.3.** Heat-treated yeast suspensions were not as effective at reducing turbidity than live yeast cells (Fig.3.36). This results gives an idea, that polysaccharide mediated biosedimentation is occurring .

**4.7.4.** As shown in Fig.3.48, volume of sediment was higher in the cone when yeast suspension was added, compared to the control cone (with only SMTTW containing 3.8 gm/L suspended solids but no yeast suspension). When just 100 ml of yeast suspension with 0.005gm/L was added to 900 ml of deionized water, the sediment volume was low (Positive control).

The turbidity in the decant of the bell jar and the 100L tank following the biosedimentation operation was reduced by about 76 and 88 NTU respectively, from 746 and 810 NTU. When the test and control cones decanted water were passed through the sand bed filter, the turbidity decreased by 0.1 NTU 40 NTU for the test and control cone respectively. The presence of Extracellular polysaccharide of yeast on the wall of the bell jar proves the mobilation of yeast in the container. Sarkaret al. (2018) conducted a comparative study of turbidity removal using biological and chemical agents and discovered that chemical coagulants were more effective for turbidity removal than natural coagulants, despite the fact that chemical coagulants increase TDS in treated water while natural coagulants decrease TDS.

**4.7.5.** Furthermore, agitating or stirring the water column resulted in a significant decrease in turbidity, thus provides a proper interaction of colloids and flocculants. The speed of agitation is suggested as 30-100 rpm for 10 min by Saritha et al., (2014), which is almost similar with that of the present study 50 rpm for higher scale biotreatment of turbidity removal.

Other parameters, such as phosphate and nitrate, were significantly reduced after yeast treatment in the SMTTW. Moreover, after passing through the sand bed filter, the mine water was colourless. Moreover, after plating of sand bed filter effluent on malt extract and nutrient agar media, no yeast colony was observed.

It can be concluded that yeast cells were caught by sand since nutrient agar plates revealed over the expansion of bacterial colony while yeast colony was lacking in plates. As a result, tertiary treatment is required to obtain turbid-free water in a short period of time. In addition, a disinfection system can be installed in the water treatment system.

For near-complete elimination of colloidal turbidity, the active cellular biocoagulant and the biotreatment process parameters can be tweaked. This could lead to the development of an environmentally friendly biotreatment technology to address the challenges created by turbidity in drinking water supplies in Goa's mining zones during monsoon.

#### **4.7.6. Yeast combination study for biosedimentation test**

A single yeast strain reduced turbidity better than a pair of yeast strains. Tubes and bottles with mixed cultures revealed massive inner wall deposition (Fig.3.59,3.56). And, in comparison to single culture's bottles, sedimentation rate was modest after forcefully agitation in bottles of mixed culture eg. As a result, it was discovered that a single culture was capable of significantly reducing turbidity in SMTTW

#### **4.8. Biosedimentation studies performed at lab scale upto the scale of 500 L to 1000L**

Aluminum sulphate, chitin, and sago were utilised in coagulation studies, and the results showed that the method efficiently eliminated turbidity from water using 0.1–0.4 g L<sup>-1</sup> of the coagulants (Saritha et al., 2017).

In present study the pH was observed similar to the other reported for efficient turbidity removal. The best pH range for removing turbidity was discovered to be 7 for alum and 6-8 for chitin, yielded the greatest turbidity removal (Saritha et al., 2017; Quasim et al., 2018). So this technique does not require pH maintenance.

**4.8.1.** Even after 1 hours, the turbidity of natural mine tailing water and SMTTW prepared from freshly collected (mine tailing soil) and processed remained between 400 and 300 NTU. It's possible that protracted storage has caused the soil to oxidize.

**4.8.2.** Significant turbidity reduction was observed in an efficient time in the 1000L test tank (in which yeast was added) when compared to the 1000L control tank (in which

yeast was not added), with sedimentation rate constants of 0.000067 and 0.000396 NTU<sup>-1</sup> min<sup>-1</sup> for control and test, respectively.

**4.8.3.** The 1000L turbidity test showed a bigger reduction than the 500L test, possibly because to the higher tank height, which provided more surface area for clay colloids and yeast cells to interact.

**4.8.4.** Plates (spread plated with water sample taken after experiment) did not reveal yeast growth on MEA plate, spread plated with test tank effluent, which was passed through the sand bed. Microbes grown in the same pattern in wastewater collected directly from the control and test tanks. A microscopic image of a microbial colony on a nutrient rich plate indicates that it was a bacterial colony. As a result, it can be assumed that the additional yeast was not present in the sand bed's outflowing contents.

**4.8.5.** The water tank was stirred 50 times in one minute for 10 minutes. The hydraulic retention time of water in the sedimentation tank may be 30-40 minutes because maximal turbidity reduction occurred within 30-40 minutes. The tank and sand bed had a flow rate of 41 minutes per litre. The turbidity of the sand bed effluent after 60 minutes was 50 NTU (for the control tank effluent) and 1.5 to 0.1 NTU (for the test sand bed effluent). There was no pressure applied during the hydraulic loading. As a result, by using pressure filters, multimedia filters with charcoal, and a further disinfection system, the best grade of treated water can be obtained from this process.

**4.8.6.** The flocculant cost associated with the proposed technique is comparatively very less than that of existing methods. In particular, the cost of coagulant for the treatment of 1000 m<sup>3</sup> of turbid water with alum, raw soybean, and defatted soybean is approximately 2441.48 INR, 939.03 INR, and 305.9 INR, respectively, as per the study shown by (Hussain.G., 2020). However, the flocculants cost associated with the proposed technique is only 63.38 INR for treatment of the same amount of turbid water. Yeast based flocculation is not only cost-effective but also has the potential of marketability. The daily costs of doses of coagulating the coagulation – flocculation process using the coagulants PACl + FeCl<sub>3</sub> and PACl were 55.87.3 INR and 6612.34 INR, respectively (Zafra, M et al., 2020). So the proposed technique is considered cost effective.

In this study the most challenging work was to prepare simulated turbid mine water of desired turbidity. Sediment concentrations cannot be determined easily or quickly in the field, and transportation to a laboratory for analysis is time-consuming and can be costly (Thackston and Palermo, 2000). As a result, these traditional methods are increasingly being replaced in favor of accurate, continuously-collected surrogate data for quantification of suspended solids that may be safer and less expensive to obtain, such as turbidity measurements.

Biosedimentation was almost reproducible at every scale in present work, it is suggested tank should be loner for more efficient biosedimentation.

#### **4.9. Yeast imobilization study in agar**

Beads were made in a cylinder shape. Beads that were not treated with yeast turned yellow, whereas those treated with yeast retained in their original colour after biosedimentation test, can be concluded that yeast retained the colour of beads.

Clays surface deposition was not seen in control beads, while test beads had both surface deposition and embedded clay. As a result, it can be inferred that immobilization and binding of clay with yeast surface occurred; however, because the turbidity of the water did not decrease much, this immobilization approach cannot be used to reduce turbid water turbidity.

#### **4.10. Molecular Study of only selected yeast Strains**

*Candida orthopsilopsis* was discovered to be the aquatic yeast utilising the PCR technique. This is the first time in the world that *C.orthopsilosis* has been isolated in a freshwater system. Earlier this species was isolated from other sources but not from freshwater. The characterization of this species also revealed that it is resistant to high pH, heat, and can thrive on modified media made from mine soil transported from the mining location. Moreover, among all studied cultures this strain displayed the best phosphorus solubilization property. Conclusion can be made that this wild type yeast can be investigated further for biotechnological uses.

After considering the phylogenetic tree of the yeast isolate, which was created by blasting with known species sequences, it revealed that it is genetically identical to

Ireland's isolate, whose gene bank number is (FN812686). Further, *C.orthopsilosis* species isolated from other environments, such as clinical, vegetation, soil, and air, beverage (Wong et al., 2020), isolated from naturally fermented Brazilian table olives (Simoes et al., 2021), can be compared to this isolate for its genetic variation.

Additionally, *C.orthopsilosis* isolated from olive oil was evaluated for safety, survival in gastrointestinal and digestion conditions, antimicrobial activity, cellular hydrophobicity, auto aggregation ability, adhesion to epithelial cells, co-aggregation, and inhibition of pathogenic bacteria adhesion and found to be good for probioactivity of the same strain (Simoes et al., 2021).

*Candida tropicalis* was identified from rain water utilizing the PCR technique. This is the first report ever to link yeast to rainwater from the southwest monsoon. This is the first report of *Candida sp.* and *Candida tropicalis* specifically in rainwater because *Candida tropicalis* has never been suspected of penetrating the southwest monsoon airborne cloud environment. The relationship between meteorological variables and yeast concentration can be studied further. At different sea levels, the variety of microbes in clouds can be examined, as well as their function in ambient settings and ecology. Since it has been proposed (Sarah et al., 2017) that cloud formation originates cloud condensation nuclei due to the presence of air particles and microbes, such as bacteria, fungi, and phytoplankton, their function in cloud nucleation could be examined.

#### 4.14 Future scope

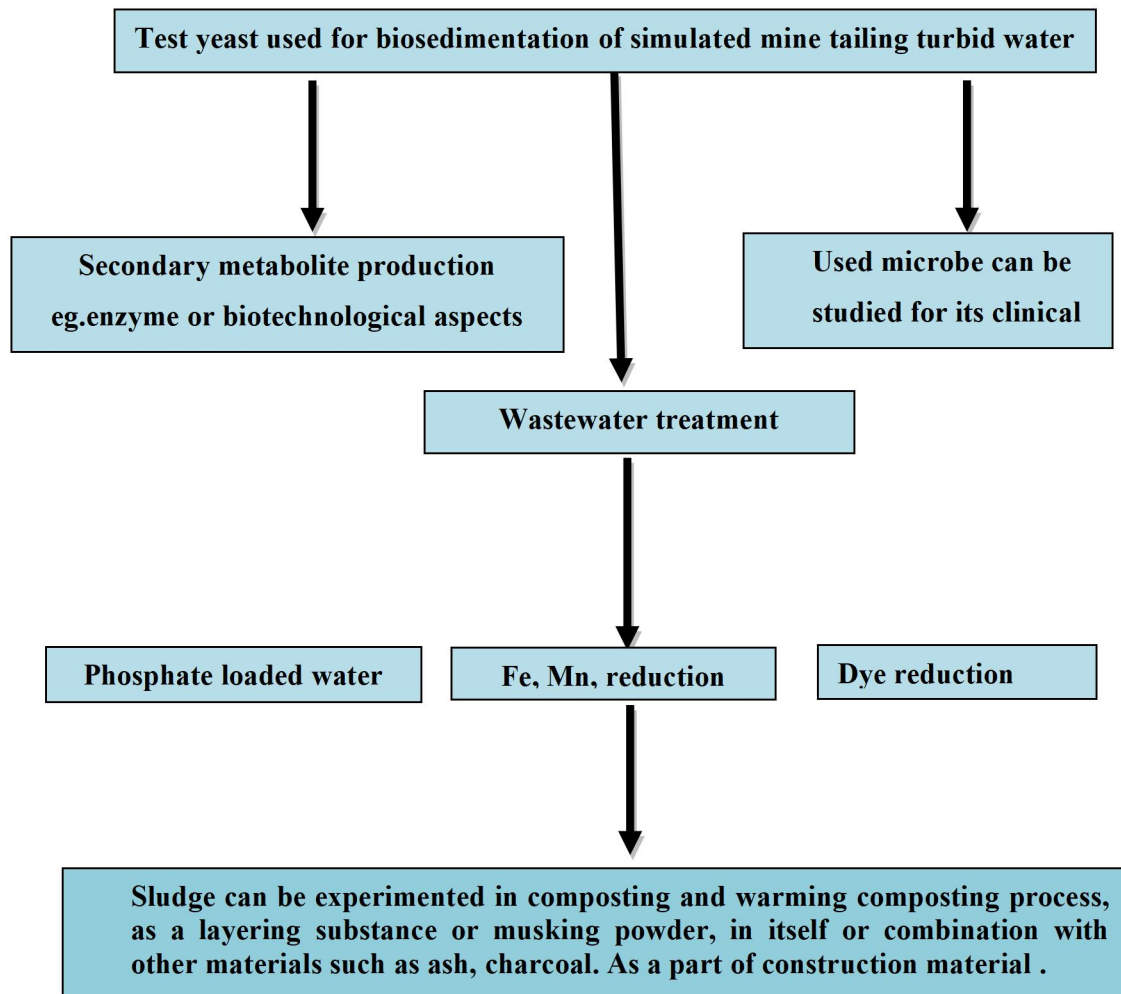


Figure.4.3: Diagram depicting the potential of the promising yeast in the future

# SUMMARY

## SUMMARY

The broad objective of this study was to evaluate promising yeast strain for the removal of colloidal turbidity from mine tailing water. It has been reported that colloidal turbidity was 4gm/l in the surface water reservoir of mining area in Goa. The key finding of this study is outlined as follows;

- First time in the world, freshwater yeast is used for the removal of turbidity of simulated mine tailing turbid water.
- Pioneer use of JFRAD in public domain for fractal analysis have performed. This image analysis based technique holds excellent potential for rapid screening of a large number of yeast strains required in different applications.
- Goa's freshwater bodies, revealed a great diversity of aquatic yeast. Freshwater ecosystem for yeast is not well explored .
- Culture was maintained on artificial media for long time without any change in morphological and turbidity removal character.
- Wild type yeast showed consistency in bioclarification property over long period of time.
- All studied water parameters were found to be similar in monsoon and non monsoon season and were below the permissible level.
- Yeast mediated biosedimentation is scale independent.
- Further, *C. orthopsilosis* species isolated from other environments, such as clinical, vegetation, soil, and air, can be compared to this isolate as it is isolated from nutrient poor freshwater body.
- Presence of yeast in rain water gives idea of circulation of microbes in the environment.

The study has several suggestions for further research work on the basis of research findings.



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# **APPENDICES**

## Appendix-I

### Media

#### **2% Malt Extract Agar (MEA)** (Commercial, obtained from HiMedia)

1 g refined bacteriological grade malt extract (HiMedia) and 3 g bacteriological grade agar (HiMedia). Dissolve and make the total volume to 600 ml with distilled water. Heat to boiling for dissolving / digest agar completely. Sterilize by autoclaving at 121 °C at 15 lbs pressure for 20 min. Allowed to cool at room temperature.

#### **Nutrient Agar** (Commercial, obtained from HiMedia)

Suspend 2.8.0 grams in 100 ml purified / distilled water. Heat to boiling to dissolve the medium completely. Sterilize by autoclaving at 15 lbs pressure (121°C) for 15 minutes. Cool to 45-50°C.

#### **Sodium Acetate Agar** (Commercial, obtained from HiMedia)

Suspend 6.19 grams (the equivalent weight of dehydrated medium per litre) of dehydrated medium in 100 ml purified / distilled water. Heat to boiling to dissolve the medium completely. Sterilize by autoclaving at 15 lbs pressure (121°C) for 15 minutes. Cool and mix well and pour into sterile Petri plates.

#### **Corn Meal Agar** (Commercial, obtained from HiMedia)

Suspend 1.7 grams in 100 ml purified/ distilled water. Heat to boiling to dissolve the medium completely.. Sterilize by autoclaving at 15 lbs pressure (121°C) for 15 minutes. Cool and mix well and pour into sterile Petri plates.

#### **Nitrate Agar** (Commercial, obtained from HiMedia)

Suspend 2.1 grams in 100 ml purified / distilled water. Heat to boiling to dissolve the medium completely. Dispense in tubes and sterilize by autoclaving at 15 lbs pressure (121°C) for 15 minutes. Allowed to cool and mix well and pour into sterile Petri plates.



**Pikovskya Agar** (Commercial, obtained from HiMedia)

Suspend 3.13 grams in 100 ml distilled water. Heat to boiling to dissolve the medium completely and sterilize by autoclaving at 15 lbs pressure (121°C) for 15 minutes. Incorporated with Bromophenol blue. Mixed well and pour into sterile Petri plates.

**Antibiotics solutions**

Stock solution of antibiotics solution was prepared by dissolving a tablet of Faropenem weighing 200mg in 50 ml sterile distilled water. Required concentration was obtained from stock solution.

**Stains**

**Lactophenol Cotton Blue** (Commercial stain obtained from HiMedia)

**Ammoniacal Congo Red (ACR)**

Dissolve 0.1 g of congo red powder (HiMedia) in 100 ml strong ammonia solution

**Reagents**

**0.1 N sodium hydroxide**

Dissolve 4 g NaOH and make the volume to 1000 ml in volumetric flask.

**2 M sodium hydroxide**

Dissolve 8 g NaOH and make the volume to 100 ml in volumetric flask.

**0.1 N hydrochloric acid**

0.1 N HCl by mixing 10.43 ml of conc. 35% HCl (Merck) carefully with distilled water and

0.2 make the volume to 1000 ml.

**0.1 M phosphate buffer pH (7.2)**

**0.2 Stock solutions preparations:**

A) 0.2 M di-sodium hydrogen phosphate, anhydrous (solution A) Take 1.42 g of Na HPO<sub>4</sub> (HiMedia) in 100 ml distilled water 2 4

B) 0.2 M sodium di-hydrogen orthophosphate, dehydrate (solution B) Take 1.56 g of NaH PO<sub>4</sub> . 2H O (HiMedia) in 100 ml distilled water 2 4 2 To prepare 0.1 M

phosphate buffer add 36 ml of solution A to 14 solution B (0.2 M phosphate buffer) + 50 ml distilled water. To this more 100 ml of distilled water was added.

### **Sterile glycerol**

Take 10 ml of glycerol (HiMedia) and make the total volume to 100 ml using 90 ml sterile distilled water, autoclaved.

### **Carbon dioxide free water**

Prepared all stock and standard solutions and dilution water for the standardisation procedure with distilled water that has been freshly boiled for 15 minutes and cooled to room temperature.

The final pH of the water was  $\geq 6.0$  and its conductivity was  $< 2 \mu\text{S/cm}$ .

### **Potassium biphthalate solution (approx. 0.05N)**

Crush 15 to 20 g primary standard potassium biphthalate ( $\text{KHC}_8\text{H}_4\text{O}_4$ ) to about 100 mesh and dry at  $120^\circ\text{C}$  for 2 h. Cool in desiccator. Weigh  $10.0 \pm 0.5$  g (to the nearest mg), transfer to a 1000 mL volumetric flask and make up the volume with  $\text{CO}_2$  free water.

### **Standard sodium hydroxide (0.02N) titrant**

Dissolve 0.8 g NaOH and dilute to 1000 mL using  $\text{CO}_2$  free distilled water. Store in air-tight, rubber stopper Pyrex/corning glass bottles to protect from atmospheric  $\text{CO}_2$ , Standardized against 0.05 N potassium biphthalate.

### **Phenolphthalein indicator**

Dissolved 0.5 g of phenolphthalein indicator in 500 mL 95% ethyl alcohol. Add 500 mL distilled water. Add drop wise 0.02 N NaOH till pink colour appears.

### **Methyl orange indicator**

Dissolve 0.5 g of methyl orange indicator and dilute to 1000 mL with  $\text{CO}_2$  free distilled water.

### **Standard sodium thiosulphate, 0.1 N**

Dissolved 25g  $\text{Na}_2\text{SO}_3 \cdot 5\text{H}_2\text{O}$  in 1000 mL of freshly boiled and cooled distilled water and standardized against potassium bi-iodate .

This initial storage is necessary to allow oxidation of bisulphite ion present. Added about 5 ml of chloroform as a preservative to minimize bacterial decomposition.

### **Starch indicator**

Prepared slurry by adding small quantity of water to 1.0 g starch powder. Add 100 ml boiling water to it and continue boiling for a few minutes until solution becomes clear. The solution is cooled and preserved by adding 1.25 g of salicylic acid or a few drops of chloroform.

#### **1. Buffer solution (pH 10)**

Dissolved 16.90 g  $\text{NH}_4\text{Cl}$  in 143 ml of conc  $\text{NH}_4\text{OH}$ . Added 1.25 g EDTA Mg salt to obtain sharp change in indicator and diluted to 25ml.

This was titrated with standard calcium solution to avoid interference produced by EDTA to the buffer. Stored in a plastic or borosilicate glass container for no longer than 1 month. Stopper tightly to prevent loss of ammonia ( $\text{NH}_3$ ) or pick up of carbon dioxide ( $\text{CO}_2$ ) from air.

#### **2. Eriochrome black T indicator**

Mix 0.5 g dye with 100 g NaCl to prepare dry powder dissolve 0.5 g of indicator in 100 ml of ethyl alcohol.

**Standard calcium solution (0.01N) (1 g/L)** Weigh accurately 1.0 g grade  $\text{CaCO}_3$  (dried at 180 °C for 1 h before weighing) and transfer to 250 ml conical flask. Placed a funnel in the neck of a flask and add 1+1 HCl till  $\text{CaCO}_3$  dissolved completely. Added 200 ml distilled water and boil for 30 min to remove  $\text{CO}_2$  Cool and added methyl red indicator.

Added  $\text{NH}_4\text{OH}$  (3N) till intermediate orange colour develops. Diluted to 1000 ml to obtain 1 ml = 1 mg  $\text{CaCO}_3$ , (1mL = 0.40 mg Ca or 0.243 mg Mg).

**Standard EDTA solution (0.01M)**

Dissolved 3.723 g EDTA-disodium salt and dilute to 1L. Standardised against standard calcium solution. 1mL = 1 mg CaCO<sub>3</sub>

**Inhibitor**

Dissolved 1.5 g hydroxylamine hydrochloride in 100mL 95% ethyl alcohol or isopropyl alcohol.

**Stock nitrate solution (1 ml= 100 1mg NO<sub>3</sub>-N)**

Dry potassium nitrate (KNO<sub>3</sub>) in an oven at 105°C for 24 h. Dissolve 721.8 mg in water and dilute to 1000 ml. Preserve with 2 ml chloroform per L. This solution is stable for at least 6 months.

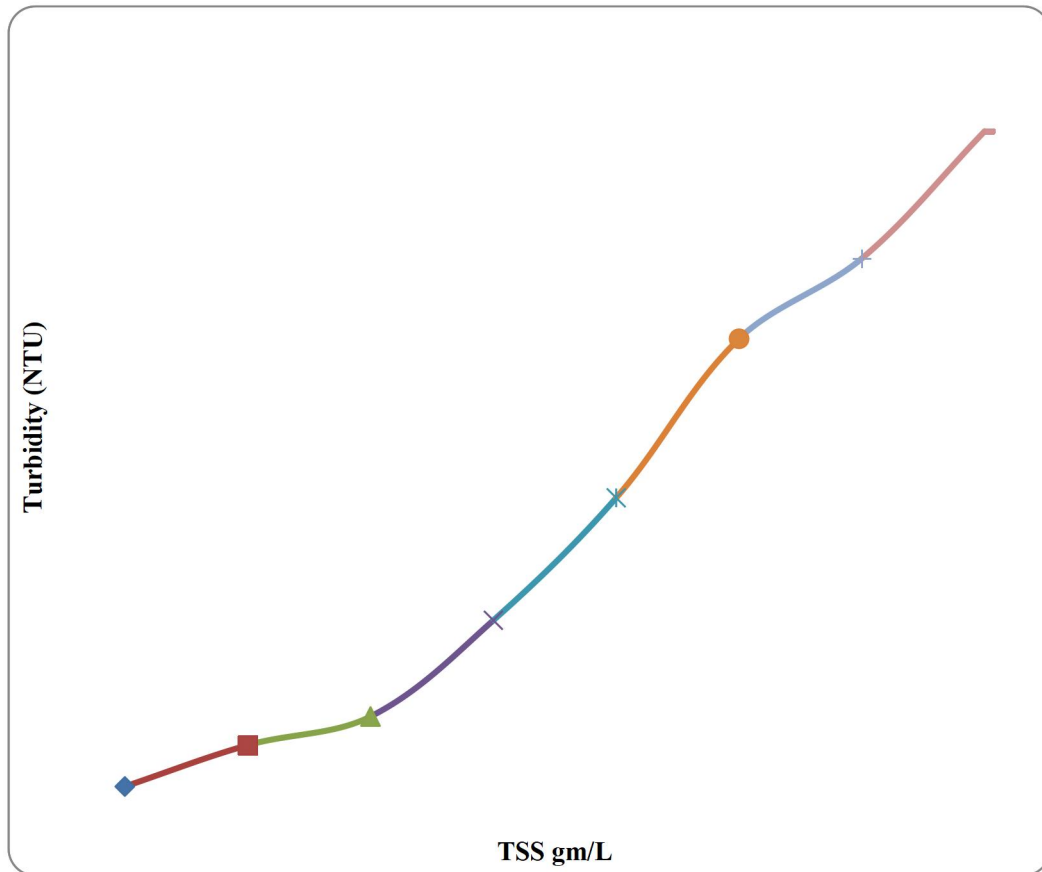
**Stock potassium solution (1.907g/L)** Dissolved 1.907 g of dry (110°C) KCl in 1000 ml of distilled water; 1 ml = 1 m

**Permissible level of water parameters.**

Parameters	Permissible limit		
	Drinking water IS 10500: 2012	WHO	EPA
pH	6.5 to 8.5	6.5-8.5	6.5-8.5
TDS (mg/l)	500	-	-
DO (mg/l)	-	-	-
Hardness (as CaCO <sub>3</sub> ) (mg/l)	200	500	-
Alkalinity (as CaCO <sub>3</sub> ) (mg/l)	200	-	-
Nitrate (mg/l)	45	50	
Sulfate (mg/l)	200	400	-
Phosphate (mg/l)	-	-	-
Fluoride (mg/l)	1		
Chloride (mg/l)	250	200	250
Turbidity (NTU)	5	0.1	-
Arsenic (mg/l)	0.01	-	-
Copper (mg/l)	0.05	1	1.5
Cadmium (mg/l)	0.003	0.003	0.005
Chromium (mg/l)	0.05	0.05	0.1
Lead (mg/l)	0.01	0.01	0.05
Iron (mg/l)	0.3	0.1	-
Zinc (mg/l)	5	5	-

### Preparation of alcohol series for SEM studies

30%, 50%, 70%, 90% Ethanol Take 30 ml of absolute ethanol and make the volume to 100 ml with respective volume of distilled water to get different concentration of ethanol.



**Figure 1. Turbidity vs SS graph of simulated turbid mine tailing water**  
**Relationship between dry weight of yeast biomass and wet weight of biomass. It is not exact but approximate.**

S.No.	Dry weight gm/L	Wet weight(gm/L)	Mean of wet weight
1.	0.005	0.045	0.047gm/l
2.	0.005	0.05	~0.05gm/L
3.	0.005	0.047	

### Research published

**Pal, S.** and Kamat, N. (2019). Fractal analysis of colony margins as an aid for screening freshwater yeast cultures for bioclarification of turbid polluted water resources. Journal of Applied and Natural Science, 11(2): 250- 256  
<https://doi.org/10.31018/jans.v11i2.2028><https://journals.ansfoundation.org/index.php/jans/article/download/2028/1763>

### (Annexure I)

**Pal, S.** and Kamat, N. (2020). Yeast in Southwest monsoon rainwater. (Fungal territory

Journal, Sci cell Publisher). [office.scicell.org/index.php/FT/article/view/95/61](http://office.scicell.org/index.php/FT/article/view/95/61).

## **(Annexure-II)**

### **Research under review**

A Quantified Approach to Test Bioefficacy of Phosphate Solubilizing Yeasts Useful in Phosphate Removal in Turbid water”. (Studies in Fungi)

### **Research communicated**

Potential of Novel Biotreatment Process Using Freshwater Yeast Cultures as Active Cellular Biofloculants for Reduction of Turbidity of due to entry of Mine Tailing Water.

### **Research Presentations**

#### **Oral**

Oral presentation on the topic, Effective use of yeast for bioclarification of turbid drinking water sources, at International Conference on Recent Innovations in Biological Research. Krishnammal college for Women. Department of Botany (DST –FIST Sponsored). PSGR . 17-18 Marc2022 (**Annexure III**)

Oral presentation on the topic, Present knowledge of sustainable bioremediation of highly turbid freshwater resources with prospects of application in mining areas of Goa, India, at National Conference on Change Environment Challenges, Solution and Strategies. Dempe college Goa. 8<sup>th</sup>-9<sup>th</sup> March 2018. (**Annexure IV**)

Oral presentation on the topic, Successful eco-friendly demonstration of yeast mediated bioclarification of inorganic colloidal turbidity in simulated mining wastewater at thousand litres scale, at National webinar on Technical Advances in Applied Microbiology. Department of Microbiology Goa University. 10<sup>th</sup>-12<sup>th</sup> November 2021. (**Annexure V**)

Oral presentation on the topic, Fractal analysis of fresh water yeast colony margins facilitates strain selection in biotreatment of turbid water, at International Symposium on Fungal Biology: Advances, Application and conservation and 45<sup>th</sup>



annual meeting of mycological Society of India. At ARI Pune Maharashtra. 19-21 November 2018. **(Annexure VI)**

### **Poster**

Poster presentation on the topic, Phosphate Solubilizing Freshwater Yeast: A Novel resource for biotreatment of turbid Phosphorous loaded polluted water. New Vistas in Botany (NSNVB, 2020) Dept. Botany Goa University 13<sup>th</sup>-14<sup>th</sup> February 2020.

### **(Annexure VII)**

Poster presentation on the topic, Yeast in Southwest monsoon rainwater: Some ecological and biometeorological implications, National symposium on Current Trend and Future Prospects in Plant Science Research at Banaras Hindu University Varanasi. 1-3<sup>rd</sup> February 2019. **(Annexure VIII)**

Poster presentation on the topic, Potential of novel biotreatment process using freshwater yeast culture as active cellular bioflocculants for reduction of turbidity of mine tailing water, National level seminar on microbial bioremediation: Novel approaches and trend. St. Xavier college Goa ,27<sup>th</sup>-28<sup>th</sup> February 2019. **(Annexure VIII)**

Poster presentation on the topic, Fractal analysis of colony margins as an aid for screening oligotrophic freshwater yeast cultures for bioclarification of turbid polluted water resources in the Iron ore mining region of Goa, International Conference on Trend in Biochemical and Biomedical Research Advances and challenges. Organized by Banaras Hindu University, Varanasi, India. 13<sup>th</sup>-15<sup>th</sup> February 2018. **(Annexure IX)**

### **Workshop, Seminar, Conference attended**

Novel Vistas in Plant Science, Department of Botany Goa University.

Attended Conference on water Management, Organized by Confederation of Indian Industry on 26 July 2019.

Contributed in Successful organization of Nobel Exhibition, Nobel media, Govt. of Goa at Kala Academy, Panaji, Goa.

Attended National webinar on “Advances in Plant Sciences” Department of Botany , Goa University.

Webinar on Book writing and related topics, Webinar on Springer Nature  
The Complete Step-by-step Guide to a Successful Career in Microbiology,  
Department of Microbiology, P.E.S's R.S.N College of Arts and Science, Farmagudi,  
Ponda-Goa.