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Abstract:	Copper poses a great risk to living organisms in the environment, calling for an alternative environmentally friendly and cost-effective technique such as microbial sorption (biosorption) for its remediation from aqueous solution. The potential of hyperbilized immobilized <i>Ragillus circulans</i> and <i>Sageharappage Caravisiae</i> to serve
	as biosorbents for biosorption of copper ions from aqueous solution in a batch process system were
	investigated. The biosorbents developed were characterised using Scanning electron microscope and Fourier
	transform infrared spectroscopy. The effects of environmental parameters on biosorption of copper ions onto
	immobilised biosorbents of Bacillus circulans and Saccharomyces Cerevisiae in a batch process system were
	also compared and investigated. The optimum parametric values such as pH, contact time, biosorbent dosage
	and initial concentration for: Copper ions onto Bacillus circulans were measured as 4, 60 min, 1.0 g and
	80 mg/l, respectively; Copper ions onto Saccharomyces Cerevisiae were recorded as 3, 80 min, 0.4 g and
	100 mg/l, respectively. The biosorption mechanism shows better fitness of Freundlich isotherm model and the
	kinetics follows Pseudo second order for both immobilised Bacillus circulans and Saccharomycesc cerevisiae
	for copper ions contacting. The results show that immobilised Bacillus circulans and Saccharomyces
	Cerevisiae are highly efficient in the removal of copper ions from aqueous solution. The biosorption capacity
	and percentage removal for copper ions onto Bacillus circulans gives better optimum performance to
	Saccharomyces cerevisiae contacting.
Keywords:	Biosorption, Saccharomyces cerevisiae, Bacillus circulans, Environmental parameters optimisation,

## Introduction

Biosorption generally known as Bioremediation involves the use of living organisms (biological systems) which include both plant and animals to detoxify and remove or reduce the environmental pollutants from air, soil and water (Guo, *et al.*, 2010; Gargouri *et al.*, 2011; Adams *et. al.*, 2015; Oyetibo *et al.*, 2017, Luka *et al.*, 2021; Luka *et al.*, 2024). Living and dead metabolically inactive microorganism or their derivatives can remove metal ions from a polluted environment (Abbas *et al.*, 2014; Farhan and Khadom, 2015; Luka *et al.*, 2022; Luka *et al.*, 2024). To be precise, Biosorption is a passive, rapid, reversible and independent metabolic energy process carried out by active or inactive microorganisms (Vendruscolo *et al.*, 2017; Luka *et al.*, 2024).

The constant wide spread of heavy metals (copper ions) into the aquatic environment can be attributed to anthropogenic activities and is one of the major problems of the ecosystems today ((Fu and Wang, 2011; Macek and Mackova, 2011; Wu *et al.*, 2012; Wang, *et al.*, 2014; Jones *et al.*, 2016; Ogunsile and Bamgboye, 2017; Luka *et al.*, 2024). Treatment of water before use is the only solution to related water-borne diseases easily identified with contaminated or polluted water (Jones *et al.*, 2016). For instance, Copper ions can cause diarrhea, dizziness and neurotoxicity (Abbas *et al.*, 2014).

Various techniques have been implemented for removal of heavy metals such as ion exchange, chemical precipitation, membrane filtration and reverse osmosis. These methods have high cost for energy and maintenance and the generation of toxic byproducts; and they are not environmentally friendly compared to biosorption (Colak *et al.*, 2011; Luka *et al.*, 2024). The use of freely suspended microbial biomass has its negative effects which includes small particle size, low mechanical strength and problems associated with separation of biomass from the effluent (White *et al.*, 1995). Due to these reasons, immobilised Bacillus Circulans and Saccharomyces Cerevisiae were used as biosorbents for the biosorption in a Batch System. The aim of this study is to compare the potential of immobilised Bacillus circulans and Saccharomyces Cerevisiae to remove copper ions from wastewater in a Batch System as well as investigating the environmental parameters affecting the biosorption.

# **Materials and Methods**

## Preparation of Stock Solution and Serial Dilution

A stock solution (100 mg/l) of copper ions was prepared from copper nitrate trihydrate  $(Cu(NO_3)_2, 3H_2O)$ . To determine the potential of immobilised *Bacillus circulans* and *Saccharomyces Cerevisiae* to remove copper ions from wastewater and for optimisation of some selected environmental factors in a batch process system. Synthetic wastewater was prepared by serial dilution of the stock solution of Copper ions solution each into distilled water with the desired metal ions solution ranged between 20 -100 mg/l and Nitrate salt was used as the counter ions for metal ions because of its low tendency to form complexes (Amirnia, 2015; Luka *et al.*, 2021; Luka *et al.*, 2023).

## Immobilisation of Biosorbent

Entrapment method that involved drop wise addition of biomass suspended in polymer matrix as adapted by Kumar *et al.*, (2011); Ilamathi *et al.*, (2014); Luka *et al.*, (2021) and Luka *et al.*, (2023) was used. The beads were produced when the alginate biomass slurry was introduced drop wise into 250 *ml* of dissolve dehydrate  $0.1 M CaCl_2$  using 20 *ml* syringe for polymerisation. The resultant beads after production were kept using Haier Thermocool Refrigerator HRD – 231SX at 4 °C (freezer) in  $CaCl_2$  solution for 12 - 14 h for hardening as depicted in Plate 1. It was then washed with 500 *ml* distilled water twice. The drained beads were spread on a dryer tray using GENLAB ENGLAND – N535 oven and kept at 80 – 100 °C for an hour and later adjusted

to 40 °C till the whole beads were dried for avoidance of burning sensation to the bead produced as shown in Plate 2. After drying, each strain of immobilised biomass bead was kept inside tight bottle for future use as biosorbent.



Plate 1: Fresh and Life Immobilised Biosorbent before Heat Killing



Plate 2: Washed and Drained Beads Spread on a Dryer Tray ready for Drying in an Oven

### Characterisation of Immobilised Bacillus circulans

The characterisation was done before and after biosorption of copper ions, respectively. PHENOM Prox Netherland Scanning electron microscope (SEM) gives information on surface morphology of each biosorbent of immobilised *Bacillus circulans* and *Saccharomyces Cerevisiae*, respectively; while AGILENT TECHNOLOGIST USA – Cary 630 Fourier transformed infrared spectroscopy (FTIR) gives information on functional groups presence and chemical composition of each biosorbent of immobilised *Bacillus circulans* and *Saccharomyces Cerevisiae*, respectively (Luka *et al.*, 2021; Luka et al., 2023).

### **Batch Process System**

Batch process system that was used in this study are Erlenmeyer conical flasks of  $100 \, ml$ . Five flasks for each metal ions were incubated in a rotary shaker type at 150 (*rpm*) and room temperature (32 °C) for 100 *min* to obtain equilibrium for each biosorbate and biosorbent used to monitor a particular parameter under investigation. The mixtures were decanted and the residual concentration of each Copper ions solution after biosorption were determined using VGP210 - AAS, respectively. The uptake value at equilibrium was calculated using equation (1) (Colak *et al.*, 2013; Dobaradaran *et al.*, 2016; Ahmad *et al.*, 2018; Luka *et al.*, 2021):

$$q_e = \frac{V(C_i - C_e)}{m}$$

(1)

Where,  $q_e$  is the uptake of metal (mg/g),  $C_i$  and  $C_e$  are the initial and equilibrium metal concentrations in the water (mg/l) respectively; V is the volume of solution (l) and m

is the mass of adsorbent (g). The removal percentage (R) of the metal ion by the biosorbent was achieved using equation (2) (Galedar and Younesi, 2013; Ahmad et al., 2018):

$$R(\%) = \frac{c_i - c_e}{c_i} \times 100$$

## 2.5 Optimisation of environmental Parameters

Optimum values of pH, biosorbent doses, contact time and initial metal concentration (Luka et al., 2024) were determined for copper ions biosorption in batch mode. The optimum value of pH for each metal ions at an initial concentration of 100 mg/l and 1.0 g biosorbent in 50 mlof metal solution at room temperature (32 °C) for 100 min at pH varying from 3.0 - 7.0using HANNA INSTRUMENTS -  $pHep^{(R)}$  pH meter were obtained by adding 0.1 M NaOH or HCl. The effect of contact time was investigated at room temperature, each metal ion initial concentration of 100 mg/l and 1.0g biosorbent in 50 ml of metal solution at 32 °C for 100 min and an optimised pH. The effect of contact time was studied by varying at 20, 40, 60, 80 and 100 min, room temperature, initial concentration of 100 mg/l and 1.0g biosorbent in 50 ml of metal solution. Furthermore, effect of biosorbent dosage were optimised by using biosorbents amounts of 0.2, 0.4, 0.6, 0.8 and 0.1 g in 50 ml of an initial metal ions concentration of 100 mg/l each, at room temperature and at optimised pH as well as contact time. For optimisation of initial metal concentration, biosorption investigation at optimised parametric values were achieved with initial metal ions concentrations each of value 20, 40, 60, 80 and 100 mg/l respectively (Samuel et al., 2013). In batch system, the mixture of biosorbent and wastewater were decanted and the clear supernatant from aqueous sample were analyzed using VGP210 - Atomic absorption spectroscopy for copper ions concentration. The final concentration of each metal ions was measured using Standard method for examination of water and wastewater as adapted by Luka et al., (2021).

### **Results and Discussion**

# Characterisation of Immobilised *Bacillus circulans* and *Saccharomyces cerevisiae*

### SEM analysis of immobilised Bacillus circulans

The SEM images results for biosorbents before and after biosorption of two different biosorbates are presented in Plates 3 - 6. Plates 3 - 4 show the change in morphology displayed by *Bacillus circulans* before and after biosorption with copper ions, respectively.



**Plate 3:** SEM Image of Immobilised *Bacillus Circulans* before Biosorption at 500x magnification



**Plate 4:** SEM Image of Immobilised *Bacillus Circulans* for Copper ions after biosorption at 500x magnification

The results shown by SEM images of immobilised *Bacillus circulans biosorbent* for copper ions for each analysis clearly show a change in the surface morphology of the biosorbents after exposure to biosorbates. The surface of biosorbents were observed to be made up of heterogeneous structure and micropores which were known for more biosorption and internally bioaccumulation of biosorbates. The results of this studies show similarity to that of Ririhena *et al.*, (2018) and Luka *et al.*, (2021) who reported that SEM of *Saccharomyces cerevisiae* after biosorption show changes and have more holes than before biosorption which look more hydrated and plumper. This morphological change occurs due to copper ions biosorption at the cell wall.

# 3.1.2 SEM analysis of immobilised Saccharomyces Cerevisiae

Plates 5 - 6 display the images of immobilised *Saccharomyces cerevisiae biosorbent*. Showing the change in morphology before and after biosorption with copper ions, respectively.



**Plate 5:** SEM Image of Immobilised *Saccharomyces Cerevisiae* before Biosorption at 500x magnification



Plate 6: SEM Image of Immobilised Saccharomyces Cerevisiae for Copper ions after biosorption at 500x magnification

The surface of Saccharomyces *cerevisiae* before biosorption (Plate 5) contains many traps but shorter than that of *Bacillus circulans* (Plate 3). The micrograph of *Bacillus circulans* before biosorption has tinier micropores on the surface to

that of *Saccharomyces cerevisiae*. After biosorption of copper ions onto *Bacillus circulans* and onto Saccharomyces *cerevisiae*, the surface structure of both biosorbents were transformed significantly as shown in Plates 4 and 6 for *Bacillus circulans* and *Saccharomyces cerevisiae* biosorbents, respectively. Concerning copper (Plates 4 and 6), the micropores structure become smoother after biosorption compared to before biosorption of the biosorbate onto the biosorbents (Abdul Raffar *et al.*, 2015; Nadeem *et al.*, 2016).

#### FTIR analysis of immobilised Bacillus circulans

The results of the FTIR Spectra for biosorbents before and after biosorption with biosorbate are presented in Figures 1 - 4.

Figures 1-2 show the trend obtained for *Bacillus* circulans



Figure 1: FTIR Spectra of Immobilised *Bacillus Circulans* before Biosorption

The FTIR spectra for *Bacillus circulans* (Figure 1) showed the functional groups that are present in the biosorbent before biosorption. The major peaks were noted at 3261.4, 2922.2, 2110.4, 1617.7, 1405.2, 1028.7 and 667.2  $cm^{-1}$ ; this implies the assigned functional group are -OH stretch for carboxylic acid, C - H stretch for aldehyde,  $C \equiv C$  stretch for alkyne, N - H bend for amine, -OH bend for phenol/ tertiary alcohol, C - N stretch for primary amine and -OH for alcohol out of phase bend, respectively (Hossian, 2013; Kariuki *et al.*, 2017).

Figures 3 – 4 give the outcome of using *Saccharomyces cerevisiae as* biosorbent.



Figure 2: FTIR Spectra of Immobilised *Bacillus Circulans* for Copper ions after biosorption

The peaks shifting assigned to copper ions after biosorption were 3268.9, 1599.0, 1416.4, 1088.4, 1032.5  $cm^{-1}$ . This verifies the interaction between hydroxyl, carboxyl, amine, phenol and alcohol functional groups with  $Cu^{2+}$ . *FTIR analysis of immobilised Saccharomyces Cerevisiae* 



Figure 3: FTIR Spectra of Immobilised Saccharomyces Cerevisiae before Biosorption

The FTIR spectra of *Saccharomyces cerevisiae* biosorbent before biosorption (Figure 3) were recorded at 3272.6, 2922.2, 2102.2, 1621.4, 1405.2, 1233.7, 1028.7 and 805.1  $cm^{-1}$ ; the corresponding functional groups are -OH stretch for carboxylic acid, C - H stretch for aldehyde,  $C \equiv C$  stretch for alkyne, N - H bend for amine, -OH bend for phenol/ tertiary alcohol, C - O stretch for alcohol/carboxylic acids/esters/ethers, C - N stretch for primary amine and C - H for aromatic out of plane bend, respectively.



Figure 4: FTIR Spectra of Immobilised Saccharomyces Cerevisiae for Copper ions after biosorption

A new FTIR peaks (Figure 4) were found at 3906.3, 3738.5, 3291.2, 3365.8, 3179.2, 2922.2, 2307.2, 2117.1, 1923.3, 1871.1, 1625.1, 1558.0, 1420.1, 1318.5, 1036.2, 931.8 and 797.7  $cm^{-1}$  after biosorption of copper ions which differ from that of *Saccharomyces cerevisiae* biosorbents before biosorption (Hasan *et al.*, 2016; Nadeem *et al.*, 2016; Benjreid *et al.*, 2018).

#### **Optimisation of environmental Parameters**

The results for optimisation of environmental parameters affecting biosorption of copper ions onto immobilised *bacillus circulans* and *saccharomyces cerevisiae* which include pH of biosorbate, contact time, biosorbent dosage and initial concentration of biosorbate are presented in Figures 5 - 12, respectively. Figures 5 - 6, present optimum values of pH of biosorbate.



Figure 5: Effect of pH on Biosorption Capacity of Copper onto Bacillus Circulans and Saccharomyces Cerevisiae



Figure 6: Effect of pH on Removal Percentage of Copper onto *Bacillus Circulans* and *Saccharomyces Cerevisiae* 

Regarding copper ions onto Saccharomyces Cerevisiae (Cu Y) as biosorbent, maximum biosorption capacity (Figure 5) and percentage removal (Figure 6) were obtained at pH of 3 with values of 3.16 mg/g and 63.25 %, respectively. Whereas copper ions onto Bacillus circulans (Cu B) as biosorbent measured maximum biosorption capacity (Figure 5) and percentage removal (Figure 6) at pHof 4 with values of 3.65 mg/g and 72.92 %, respectively. The results obtained are in line with Jones et al. (2016) which shows fluctuations in biosorption efficiency with a pН constant increase in of the biosorbates of  $Cd^{2+}$ ,  $Fe^{2+}$ ,  $Ni^{2+}$  and  $Zn^{2+}$  using Mucilage from Dicerocaryum Eriocarpum Plant as biosorbent.

The batch investigation on effect of contact time is presented in Figures 7 - 8 for copper and ions, chronologically.



Figure 7: Effect of Contact Time on Biosorption Capacity of Copper onto *Bacillus Circulans* and *Saccharomyces Cerevisiae* 



Figure 8: Effect of Contact Time on Removal Percentage of Copper onto *Bacillus Circulans* and *Saccharomyces Cerevisiae* 

Considering, copper ions, optimum biosorption capacity (Figure 7) and percentage removal (Figure 8) for Saccharomyces cerevisiae were observed at contact time of 80 min with values of 3.66 mg/g and 73.18 %, respectively. Decrease was recorded for Saccharomyces cerevisiae at 100 min. Whereas biosorption capacity (Figures 7) and percentage removal (Figure 8) for copper ions onto Bacillus circulans increases from 20 min to 60 min and a decrease at 80 min. Therefore, maximum biosorption capacity and percentage removal were recorded at contact time of 60 min with values 3.69 mg/ g and 73.85 %, respectively. The decrease in biosorption capacity of copper ions (Figure 7) and percentage removal of copper ions (Figure 8) after optimum values were observed might be due to gradual exhaustion of the number of vacant site available. The results of this research show similarity to those of Colak et al. (2011), where Biosorption of lead from aqueous solutions by Bacillus strains possessing heavy-metal resistance as biosorbent show that larger number of vacant surface sites are available for biosorption in the initial stage and virtually remain constant with increase in contact time.

From contact time optimisation, the following can be drawn: the optimum values for copper ions were obtained at 100 min and 60 min using *Bacillus circulans* as biosorbent with optimum percentage removal values recorded as 73.85 %. In addition, copper ions optimum percentage removal was found at 60 min. This is in agreements with the explanation given by Jones et al. (2016) for  $Cd^{2+}$ ,  $Fe^{2+}$ ,  $Ni^{2+}$  and  $Zn^{2+}$  ions using Mucilage from *Dicerocaryum Eriocarpum* Plant as biosorbent, where biosorption increased rapidly at the beginning and became very slow or steady with time.

The results for optimum biosorbent dosage are shown in Figures 9 - 10.



Figure 9: Effect of Biosorbent Dosage on Biosorption Capacity of Copper onto *Bacillus Circulans* and



Figure 10: Effect of Biosorbent Dosage on Removal Percentage of Copper onto *Bacillus Circulans* and *Saccharomyces Cerevisiae* 

Secondly, copper ions as recorded a decrease in the values of biosorbent capacity with increase in the values of biosorbent dosage from 0.2 g to 1.0 g. The maximum biosorption capacity of copper ions onto *Bacillus circulans* (Cu B) and copper ions onto *Saccharomyces cerevisiae* (Cu Y) were observed at 0.2 g as depicted in Figure 9 with value 15.87 mg/g and 15.26 mg/g, respectively. The highest percentage removal for copper ions onto *Bacillus circulans* (Cu B) and copper ions onto *Saccharomyces cerevisiae* (Cu Y) were measured at 1.0 g with value of 72.15 % and at 0.4 g with value of 61.83 % as displayed in Figure 10, respectively.

Based on the results recorded on optimisation of biosorbent dosage the following can be established: The values of biosorption capacity decreases for copper ions using either immobilised *Bacillus circulans* or *Saccharomyces cerevisiae* as biosorbent with increases in biosorbent dosage. The maximum percentage removal using *Saccharomyces cerevisiae* biosorbent was recorded as 61.83 % at 0.4 g for copper ions; while, maximum percentage removal using *Bacillus circulans* biosorbent was measured as 72.15 % at 1.0 g also, for copper ions.

The illustration for effect of initial concentration of biosorbate on biosorption capacity and percentage removal are presented in Figures 11 - 12.



Figure 11: Effect of Initial Concentration on Biosorption Capacity of Copper onto *Bacillus Circulans* and *Saccharomyces Cerevisiae* 





Regarding copper ions, optimum biosorption capacity for copper ions onto *Bacillus circulans* (Cu B) and copper ions onto *Saccharomyces cerevisiae* (Cu Y) were observed at 100 mg/l as observed in Figure 21 with value 3.56 mg/g and 7.87 mg/g, respectively. The maximum percentage removal for copper ions onto *Bacillus circulans* (Cu B) and copper ions onto *Saccharomyces cerevisiae* (Cu Y) were measured at 80 mg/l with value 88.58 % and at 100 mg/l with value 62.94 % as displayed in Figure 22, respectively.

Based on the results recorded for effect of initial concentration of biosorbate the following can be made: The values of biosorption capacity increases with increase in initial concentration values for copper biosorbates. The percentage removal increases with increase in initial concentration from 20 mg/l to 100 mg/l for copper ions. This may be attributed to increase in concentration gradient with increase in initial concentration of biosorbate used as a driving force to minimise or eliminate mass transfer resistance between the aqueous biosorbate and the solid biosorbent. This can be explained by Jones et al. (2016), who reported that biosorption of  $Cd^{2+}$ ,  $Fe^{2+}$ ,  $Ni^{2+}$  and Zn<sup>2+</sup> onto Mucilage from *Dicerocaryum Eriocarpum* Plant as biosorbent increase with increase in the initial concentration of biosorbates. The results also agree with Kariuki et al. (2017), where biosorption of lead and copper ions using rogers mushroom biomass as biosorbent increase with increase in the initial concentration of each biosorbate.

The results of biosorption isotherm testing for Langmuir isotherm model are presented in Figure 13 and the Langmuir model isotherm parameters are presented in Tables 1.

**Biosorption Isotherm and Kinetics Testing** 



**Biosorption Isotherm** 



Figure 13: Langmuir Model Fitness for Copper ions onto Bacillus Circulans and Saccharomyces Cerevisiae

Table 1: Langmuir Isotherm Model Parameters

Sample	$q_{max} \ ({\rm mg/g})$	K <sub>L</sub> (l/mg)	RL	$\mathbb{R}^2$
Cu Y	-0.2308	-0.0248	-0.6757	0.1089
Cu B	-1.5056	-0.0253	-0.6536	0.9144

The Langmuir isotherm model gives maximum value of coefficient of determination ( $\mathbb{R}^2$ ) for copper ions onto *Bacillus circulans* (*Cu B*) as 0.9144 and lowest value recorded as 0.1089 for copper ions onto *Saccharomyces cerevisiae* (*Cu Y*) as shown in Table 1. The monolayer saturated capacity ( $q_{max}$ ), Langmuir constant ( $K_L$ ) and separation factor ( $\mathbb{R}_L$ ) give negative values for the two contacting processes which implies that the biosorption data do not give better fit of Langmuir isotherm model (Hossain, 2013; Kariuki *et al.*, 2017; Luka *et al.*, 2021).

#### Freundlich model

Figure 14 illustrated Freundlich isotherm model fitness testing and Freundlich model isotherm parameters are presented in Tables 2.



Figure 14: Freundlich Model Fitness for Copper ions onto Bacillus Circulans and Saccharomyces Cerevisiae

Comparism of Biosorption of Copper ions by immobilised *Bacillus circulans* and *Saccharomyces Cerevisiae* from Wastewater **Table 2**: Freundlich Isotherm Model Parameters model's values for copper ions biosorption onto

Sample	1/n	n	K <sub>F</sub> (mg/g l)	R <sup>2</sup>
Cu B	5.7072	0.1752	1.84 X 10 <sup>-8</sup>	0.9219
Cu Y	42.399	0.0236	3.5 X 10 <sup>-67</sup>	0.2529

The Freundlich isotherm model parameters shown in Table 2 presented the coefficient of determination  $(R^2)$  for the biosorption of copper onto biosorbent with the maximum value recorded as 0.9219 for copper ions onto Bacillus circulans(Cu B) and lowest value found as 0.2529 for copper ions onto Saccharomyces cerevisiae (CuY). The values of intensity of heterogeneity  $(\frac{1}{n})$  were recorded to be more than one (1), which shows that the biosorption process is cooperative. Furthermore, values of intensity of biosorbate biosorption onto biosorbent (n) is less than one for all biosorption process which indicates biosorption of all biosorbate onto biosorbent are favourable. The equivalent maximum biosorption capacity for Freundlich constant  $(K_F)$ gives value measured in the ranged of  $3.5 \times 10^{-67}$  –  $1.84 \times 10^{-8} mg/g l$ . The higher the value of  $K_F$  implies higher maximum biosorption capacity of biosorbate onto biosorbent. Therefore, Freundlich isotherm parameters are indicative of favourable fitness of Freundlich isotherm model for all biosorbate biosorption onto biosorbent (Hossain, 2013; Kariuki et al., 2017; Verma et al., 2017; Ahmad et al., 2018; Luka et al., 2021;).

## Batch Biosorption kinetic models testing

Results of biosorption kinetic testing are presented in Figures 15 - 17.

#### Pseudo first order model

Pseudo first order model plots are given in Figures 15 and model parameters are shown in Table 3



Figure 15: Pseudo first order Model Fit of Copper ions onto *Bacillus Circulans* and *Saccharomyces Cerevisiae* 

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Sample	<i>K</i> <sub>1</sub> (1/min)	$q_e \; ({ m mg/g})$	$\mathbb{R}^2$
Cu Y	-0.0015	2.1888	0.0333
Cu B	0.0193	2.233	0.1393

Figure 15 and Table 3 show that Pseudo first order Kinetics Model plots and Parameters obtained recorded lower values of coefficient of determination ( $\mathbb{R}^2$ ) and the experimental  $q_e$  values are not in agreement with the model's values for copper ions biosorption onto the different biosorbents tested. This is an indication that Pseudo first order Kinetics Model does not fit the data for biosorption of copper ions onto *Bacillus circulans* and *Saccharomyces cerevisiae* (Verma *et al.*, 2017; Ahmad *et al.*, 2018; Luka *et al.*, 2021).

## Pseudo second order model

Pseudo second order model graphs are illustrated in Figure 16 and model parameters are presented in Table 4



Figure 16: Pseudo second order Model Fit of Copper ions onto Bacillus Circulans and Saccharomyces Cerevisiae

#### Table 4: Pseudo second order Model Parameters

Sample	$K_2$ (g/mg min)	$q_e \ ({ m mg/g})$	h (mg/g min)	$\mathbb{R}^2$
Cu Y	0.1297	3.35	1.4560	0.9717
Cu B	0.7655	3.50	9.3990	0.9685

The experimental values obtained were tested using Pseudo second order kinetics model and Parameters as depicted in Figure 16 and Table 4. Higher values of coefficient of determination  $(R^2)$  for all the biosorption carried out ranged from 0.9717 - 0.9685 were recorded and the experimental  $q_{\rho}$  values agree with the model's values for all analysis. The  $R^2$  shows that *Bacillus circulans*(Cu B) gives better fitness of Pseudo second order kinetics model to Saccharomyces cerevisiae (CuY). This also indicate that copper ions onto different biosorbents used in this research agree with Pseudo second order kinetics model and the rate-limiting step may be chemisorption that involves valance forces through exchange or sharing of electrons (Hossain, 2013; Verma et al., 2017; Ahmad et al., 2018). The values of  $(K_2)$  which suggest the biosorption process in two reactions. The first one is the fast reaction, and equilibrium is attaining quickly while the second is slower reaction that can proceed for long time. The  $K_2$  values for copper biosorption of biosorbent are less than one which indicate the mechanism is not chemisorption. The initial biosorption rate of copper (h)recorded the highest value as 9.3990 mg/gmin for Bacillus circulans (Cu B) with the lowest value found as 1.4560 mg/gmin for Saccharomyces cerevisiae (Cu Y). Intra particle diffusion model

Intra particle diffusion model results are demonstrated in Figure 17 and model parameters are presented in Table 5.



Figure 17: Intra particle Diffusion Model Fit of Copper ions onto Bacillus Circulans and Saccharomyces Cerevisiae

Table 5:	Intra Particle	Diffusion	Model	Parameters
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Sample	$K_{ip} \ (\mathrm{mg/g} \ \mathrm{min}^{0.5})$	C (mg/g)	$\mathbb{R}^2$
Cu Y	0.0539	2.7940	0.2052
Cu B	0.0134	3.3897	0.0137

Furthermore, experimental results obtained from batch contactors were also tested for Intra particle diffusion kinetic model and parameters as presented in Figure 17 as well as Table 5. Figure 17 clearly observed that for copper biosorption onto different biosorbents investigated an intercept (C) was established. The values of intercept ranged from 2.7940 - 4.5870 mg/g illustrating that the line does not pass through the origin and describing the boundary layer effects. Therefore, intra particle diffusion is not the rate controlling mechanism for biosorption (Amirnia, 2015; Nadeem *et al.*, 2016; Verma *et al.*, 2017; Luka *et al.*, 2021).

## Conclusions

The following conclusions can be made from the results of this research:

- 1. Bacillus circulans and Saccharomyces cerevisiae biosorbents were able to remediate copper ions from wastewater; indicating suitability of both biosorbents for biosorption as depicted by Scanning electron microscope and Fourier transform infrared spectroscopy.
- 2. Immobilised *Bacillus circulans* gives better performance to *Saccharomyces cerevisiae* as biosorbents for the biosorption of copper ions from wastewater in a Batch process system.
- Regarding batch contactor models, Freundlich isotherm model parameters are indicative of best fitness for copper biosorption onto the two different biosorbent examined. The kinetics obeyed Pseudo second order kinetics model with higher values of R<sup>2</sup> for all experimental data which ranged from 0.9717 – 0.9685.

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