

# Toluene Degradation By Free *Staphylococcus Gallinarum* And Immobilized On Multi-Walled Carbon Nanotubes

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## ABSTRACT

Hydrocarbons pollution is a most important environmental and health anxiety. Using free and immobilized bacteria could be a suitable attitude to find a proper bioaugmentation agent. A toluene degrading bacterium was isolated from oil-contaminated environs (located in Bandar-Anzali, Guilan, Iran). The strain was molecularly identified as *Staphylococcus gallinarum* ATHH41 (Accession number: KX344723) by partial sequencing of 16SrDNA gene. The response surface methodology (RSM) was expanded for biodegradation of the toluene by ATHH41. The central composite design (CCD) was utilized to optimize pH, temperature, and toluene concentration by ATHH41. In accordance with the optimization purpose of the Design-Expert software, the optimum circumstances of toluene degradation were obtained when pH, temperature and toluene concentration were adjusted to 7.68, 31.73°C and 630.04 mg.l<sup>-1</sup>, respectively. Multi-walled carbon nanotubes (MWCNTs) were used to immobilize the strain. Infrared spectroscopy and scanning electron microscopy showed that the cells adhered to the MWCNT surface and developed a biofilm. Results reveal that free cells were able to degrade 68.01% of the toluene as the sole carbon and energy source within 24 h under optimized conditions. The immobilized cells reached 95.68%.

**Keywords:** Carbon nanotube; Response surface methodology; *Staphylococcus gallinarum* ATHH41; Toluene.

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Widespread occurrence of accidental leakage and spillage of petroleum hydrocarbons from underground pipelines and storage tanks often contaminates groundwater and may pose health hazards to the nearby populace (Hilpert et al. 2015). Volatile monoaromatic hydrocarbons of crude oil and petroleum by-products, which are generally found together, are benzene, toluene, ethyl benzene, and xylene (BTEX). These contaminants are a source of health hazards and pollute surface and ground waters (Lee et al. 2018). The remediation of the contamination of BTEX compounds is difficult due to their relative water solubility (Yakout & Daifullah 2014). Hydrocarbon bioremediation depends on the biodegradation activity of soil bacteria. Bacterial biodegradation of these compounds is considered as the most active process in petroleum degradation, and bacteria are known as the primary degraders of spilled oil (Brzecz & Kaszycki 2018). The preparation of bacteria in a ready-to-use form that is appropriate to the contaminated site is one of the major issues in bioremediation. However, the use of free-living cells of oil degrading bacteria shows that they have limited efficiency and are not reusable in a continuous treatment system. Therefore, immobilization of bacterial cells on a solid support material is an approach to overcoming such problem (Nopcharoenkul et al. 2013). The technology of immobilized microorganisms can be applied in biological treatments to enhance the efficiency and effectiveness of biodegradation given the higher specific surface areas for microbial growth and better resistance against chemical toxicities and environmental stresses (e.g. pH, temperature, and toxic substances) compared to suspended cells (Wang et al. 2015; Yan et al. 2013). Bina et al. (2012) reported the efficiency of toluene adsorption to be 99.5% by multi-walled carbon nanotube in terms of 10 mg.l<sup>-1</sup> of toluene, 1 g.l<sup>-1</sup> of carbon nanotube, 10 min exposure time and neutral pH. Multi-walled carbon nanotubes (MWCNTs) are a capable candidate caused by the structure of their pores, the wide spectrum existence of surface functional groups, and their unique properties (Rahman et al. 2017). Anjum et al. (2019) showed that the surface modified MWCNTs presented a fast and efficient removal of BTX with the highest adsorption capacity. MWCNTs have got applications in various takes, such as the adsorption of pollutants due to their chemical, mechanical, electrical and thermal properties (Pourfayaz et al. 2013). MWCNTs have high surface area, hydrophobic property, and chemical and thermal stability. Thus, they can be suitable adsorbents for volatile organic compounds (Rahman et al. 2017).

This study aimed to isolate bacterial strain with toluene degradation ability, to molecularly identify toluene-degrading bacteria from oil-polluted soils, and to optimize the medium culture conditions to investigate the toluene biodegradation by free-living and MWCNTs-immobilized cells of

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30 isolate. In addition, the effects of MWCNTs concentration, environmental conditions such as pH,  
31 temperature, and toluene concentration were evaluated on the biodegradation efficiency of toluene.

## 32 **MATERIALS AND METHODS**

### 33 MEDIUM AND CULTURE CONDITIONS

34 Different polluted soil samples were gathered from the Caspian Sea (Bandar-Anzali, Guilan,  
35 Iran, (is located in the north of Guilan Province with the coordinates of 37 28' 16 North, 49 27' 44  
36 East)). Samples were stored at 4°C preceding to utilization. Toluene (purity of 99.5%) was filtration-  
37 sterilized and used as the sole carbon and energy sources to enrich culture media for the isolation of  
38 degrading bacteria. An amount of 5 g of soil sample was combined to 50 ml of mineral salt medium  
39 (MSM) complemented with 1% (V/V) toluene. The liquid mineral salt medium (MSM) comprised of  
40 (g.l<sup>-1</sup>) 4 g NaNO<sub>3</sub>, 1.5 g KH<sub>2</sub>PO<sub>4</sub>, 0.5 g Na<sub>2</sub>HPO<sub>4</sub>, 0.2 g MgSO<sub>4</sub>.7H<sub>2</sub>O, 0.0011 g FeSO<sub>4</sub>.H<sub>2</sub>O, 0.01 g  
41 CaCl<sub>2</sub>, and pH was regulated to 7 before autoclaving. The samples were incubated at 30°C shaken at  
42 150 rpm for 7 days. After an enrichment period, 1 ml of the culture was transferred into the fresh MSM  
43 medium and incubated at 30°C shaken at 150 rpm (Zhang et al. 2013). After three subcultures, 0.1 ml  
44 of the culture was spread on MSM and nutrient agar plates and incubated at 37°C for 24-48h.

### 45 IDENTIFICATION OF STRAIN ATHH41 BY 16S RDNA SEQUENCE

46 The bacterial chromosomal DNA was extracted, using the method of CTAB, and identified by  
47 1% agarose electrophoresis (Raieta et al. 2015). The forward primer was 27R-  
48 AGAGTTTGATCMTGGCTCAG and the reverse primer was 1502F-  
49 GGTTACCTTGTTACGACTT. For the PCR outcome system, states were as follows: 2.5µl DNA  
50 templates (70 ng/µl); 0.5 µL dNTP mixture (10 mM); 0.4 µL 27 F (10 µmol/L); 0.4 µL 1502 F (10  
51 µmol/L); 1 µL 10X PCR Buffer (2.5) with MgCl<sub>2</sub> (50 mM); 0.3 µL Taq DNA polymerase (5 U/µl); 17.4  
52 µL bringing up ddH<sub>2</sub>O. The PCR amplification states were as follows: force-degeneration at 95°C for 5  
53 minutes, degeneration at 95°C for 1 minute, annealing at 60°C for 30 seconds and at 72°C for 35  
54 seconds, 30 cycles, with another extension at 72°C for 5 minutes (Madueno et al. 2011). After  
55 purification, the PCR products were sent for sequencing by Iranian Biological Resource Center.

### 56 DESIGN OF EXPERIMENTS AND MODELLING

57 Twenty runs and six replications of the central points were chosen to verify the initial pH,  
58 temperature and toluene concentration for the highest degradation of toluene. RSM with a three-factor,  
59 three-level CCD design was managed to optimize the response, Y (toluene degradation) of three  
60 variables:

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$$Y = b_0 + b_1X_1 + b_2X_2 + b_3X_3 + b_{11}X_1^2 + b_{22}X_2^2 + b_{33}X_3^2 + b_{12}X_1X_2 + b_{13}X_1X_3 + b_{23}X_2X_3 \quad (1)$$

where Y is the forecast response factor;  $X_1$  is pH;  $X_2$  is temperature ( $^{\circ}\text{C}$ );  $X_3$  is toluene concentration ( $\text{mg}\cdot\text{l}^{-1}$ ),  $b_0$ ,  $b_1$ ,  $b_2$ ,  $b_3$ ,  $b_{11}$ ,  $b_{22}$ ,  $b_{33}$ ,  $b_{12}$ ,  $b_{13}$  and  $b_{23}$  are constant regression coefficients of the model, in which  $b_0$  is the intercept term,  $b_1$ ,  $b_2$ , and  $b_3$  are linear coefficients, and  $b_{11}$ ,  $b_{22}$ , and  $b_{33}$  are squared coefficients. On the other hand,  $X_1$ ,  $X_2$  and  $X_3$  are independent factors. Combinations of factors (such as  $X_1X_2$ ) represented the interaction between the individuals (Azaman et al. 2010). The genuine factor level relating to the coded factor levels are shown in Table 1. The ranges of factor levels for the experimental design were selected based on the original medium. The optimal culture conditions for maximum toluene degradation and the coefficients in the second-order polynomial (Eq. 1) were calculated by statistical analysis using the Design Expert Software (version 7.1).

**Table 01.** Levels and codes of variables for central composite design and related strains

Variables	Level code				
	-1.68	-1	0	1	+1.68
$X_1$	5.32	6	7	8	8.68
$X_2$	21.59	25	30	35	38.41
$X_3$	195.46	400	700	1000	1204.54

$X_1$ : pH;  $X_2$ : Temperature ( $^{\circ}\text{C}$ );  $X_3$ : Toluene concentration ( $\text{mg}\cdot\text{l}^{-1}$ )

Source: The Author

## 71 TOLUENE BIODEGRADATION ASSAY

72 The isolated Bacteria were grown at  $30^{\circ}\text{C}$ , 150 rpm, in MSM medium containing 1% (v/v) of  
73 toluene for 24 h. The cells were harvested by centrifugation at  $10,000\times g$  for 10 min and washed twice  
74 in sterile MSM and re-suspended with one-tenth volume of medium. This cell suspension was operated  
75 as inoculum for consequent experiments. The toluene degradation was done by dissolving the residual  
76 toluene of the medium in 3 ml n-hexane and reading the optical density of the toluene against a blank  
77 at 200-400 nm in a UV-visible spectrophotometer (UV-vis-3600, Mapada) (Berlendis et al. 2010).

## 78 PREPARATION AND CHARACTERIZATION OF MWCNT

79 1 g of MWCNTs (5-10 nm inner diameter, 20-30 nm outer diameter, surface area  $>110 \text{ m}^2\cdot\text{g}^{-1}$ ,  
80 purity above 98%, US Research Nanomaterials, Houston, TX, USA) was soaked in 60 ml of  $\text{HNO}_3$  and  
81  $\text{H}_2\text{SO}_4$  (3:1) and dispersed using a probe sonicator for 3 h (Zhang et al. 2011). The suspension was  
82 filtered through a  $0.45 \mu\text{m}$  membrane filter, and the MWCNTs were washed with deionized water until  
83 neutral pH was reached; then, it was dried for 12 h at  $60^{\circ}\text{C}$  and stored for further use (Pan et al. 2007).  
84 The MWCNTs were characterized by the scanning electron microscopy (SEM) (AIS-2100, Seron  
85 Technologies, Gyeonggi-do, Korea) and the Fourier transform infrared spectroscopy (FT-IR) (Perkin  
86 Elmer, Waltham, MA, USA).

87 **BACTERIAL IMMOBILIZATION BY MWCNTS FOR TOLUENE REMOVAL**

88 MWCNTs were dispersed in sterile distilled water to yield concentrations of 0.005, 0.025, 0.05,  
89 0.25 g.l<sup>-1</sup> under ultrasonication for 30 min. Then, 10 µL of bacterial suspension with a density of 0.5  
90 McFarland were re-suspended in MSM and 100 µL of MWCNTs suspensions were added. After  
91 incubation for 24 h with shaking at 150 rpm, toluene degradation was determined by dissolving the  
92 residual toluene of the medium in 3 ml n-hexane and reading the optical density of the toluene against a  
93 blank at 200-400 nm in a UV-visible spectrophotometer.

94 **SEM OBSERVATIONS AND FT-IR ANALYSIS**

95 The carbon nanotubes with adhered cells were analyzed by SEM after being rinsed three times  
96 with sterile distilled water to remove unattached cells. The surface organic structures were studied by  
97 FT-IR. The spectra were recorded at 4 cm<sup>-1</sup> and 0.01 cm<sup>-1</sup> of resolution between 4000 and 500 cm<sup>-1</sup>  
98 using a Perkin Elmer Spectrum one series model instrumental analysis with the KBr disc method.

99 **RESULTS**

100 **TOLUENE DEGRADING ISOLATE CHARACTERIZATION**

101 After sampling from oil-contaminated soils and enrichment procedures in MSM toluene-  
102 containing medium, toluene-degrading bacterial strain was isolated. The bacterium in the strain  
103 surviving presence of toluene isolated in this study was designated as *Staphylococcus gallinarum* ATHH41.  
104 *Staphylococcus gallinarum* ATHH41 cells were cocci-shaped, gram-positive, catalase, nitrate positive, and  
105 oxidase negative. Almost complete sequences of the 16S-rDNA of the strain *Staphylococcus gallinarum*  
106 ATHH41 (1380 bases) were determined. The BLAST algorithm downloaded from the Genebank  
107 database (<<http://www.ncbi.nlm.nih/BLAST>>) exhibited 99% identified with the closest match for  
108 *Staphylococcus gallinarum* ATCC35539. The strain reported in this paper has been deposited in the  
109 GeneBank database under the accession number of KX344723. Fig. 1 shows a phylogenetic tree of  
110 *Staphylococcus gallinarum* ATHH41 that was constructed using the MEGA (version 5.2) (Tamura et al.  
111 2011).

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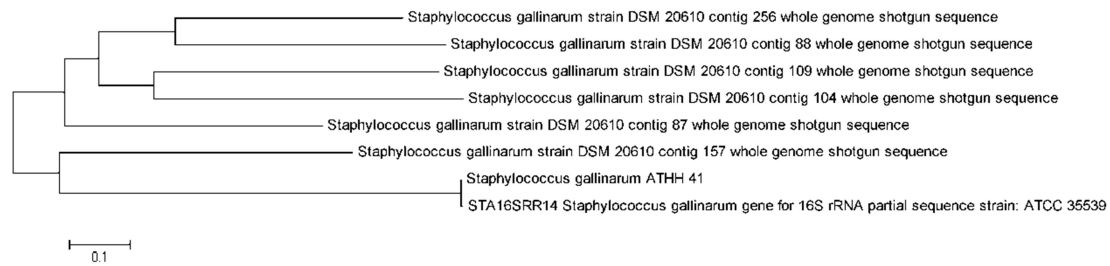
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**Figure 01.** Phylogenetic tree of the 16S rDNA sequence of *Staphylococcus gallinarum* ATHH41 and related strains



Source: The Author

117

### 118 RSM MODEL DEVELOPMENT

119 Instead of optimizing medium composition by one factor at a time approach, the statistical  
 120 RSM design provides the opportunity to determine the optimal conditions in any given parameter by  
 121 establishing the relationship between factors and predicted responses (Myers et al. 2016). The RSM  
 122 design was applied to obtain the precise factor values, which results in the higher toluene degradation.  
 123 The results are summarized in Table 2.

**Table 02.** RSM design for the three factors and their experimental results

Run order	Factors			Toluene biodegradation (%)	
	X <sub>1</sub> <sup>a</sup>	X <sub>2</sub>	X <sub>3</sub>	Experimental <sup>b</sup>	Predicted
1	7	30	195.46	60.703	60.40
2	6	25	400	48.016	47.90
3	8	25	400	57.155	57.14
4	6	35	400	59.168	60.36
5	8	35	400	67.958	67.52
6	7	21.59	700	54.941	55.38
7	5.32	30	700	51.229	50.22
8	7	30	700	69.636	69.22
9	7	30	700	69.722	69.22
10	7	30	700	69.729	69.22
11	7	30	700	69.501	69.22
12	7	30	700	68.301	69.22
13	7	30	700	68.394	69.22
14	8.68	30	700	65.393	66.21
15	7	38.41	700	64.537	63.91
16	6	25	1000	55.313	55.89
17	8	25	1000	68.8	67.74
18	6	35	1000	55.505	55.66
19	8	35	1000	65.179	65.43
20	7	30	1204.54	65.251	65.36

<sup>a</sup> X<sub>1</sub>: pH; X<sub>2</sub>: Temperature (°C); X<sub>3</sub>: Toluene concentration (mg.l<sup>-1</sup>).

<sup>b</sup> The results are presented as the means of duplicates.

Source: The Author

124

125 TOLUENE BIODEGRADATION

126 By applying multiple regression analysis to the experimentally determined data in Eq. (1), the  
 127 regression coefficients were estimated and the following second-order polynomial equation was  
 128 obtained for toluene biodegradation:

129 
$$Y = 69.22 + 4.75X_1 + 2.45 X_2 + 1.48X_3 - 3.89X_1^2 - 3.39X_2^2 - 2.24X_3^2 - 3.17X_2X_3 \quad (2)$$

130 The predicted optimum levels of  $X_1$ ,  $X_2$ ,  $X_3$  were obtained by applying regression analysis (Eq.  
 131 2), and they were 7.68 of pH, 31.73°C of temperature, and 636.04 mg.l<sup>-1</sup> of toluene concentration,  
 132 respectively. The prediction of toluene biodegradation was 70.73%. The coefficient of determination  
 133 ( $R^2$ ) of the regression for the response related to significant effects on the model was 0.96, which  
 134 means that the sample variation of 96% for toluene degradation was attributable to the factors. The  
 135 adequacy of the full quadratic model of toluene degradation was also evaluated with ANOVA. Model  
 136 summary statistics in Table 3 indicated the adequacy of the models including linear, two-factor  
 137 interactions, and quadratic terms. Linear and interaction models for toluene degradation were  
 138 significant.

**Table 03.** Analysis of variance for response surface quadratic model obtained from experimental design

Source	Sum of Squares	DF	Mean Square	F Value	Prob > F
Model	0.023	9	2.595*10 <sup>-3</sup>	26.88	<0.0001***
X <sub>1</sub>	3.162*10 <sup>-3</sup>	1	3.162*10 <sup>-3</sup>	32.76	0.0002***
X <sub>2</sub>	1.042*10 <sup>-3</sup>	1	1.042*10 <sup>-3</sup>	10.79	0.0082**
X <sub>3</sub>	6.851*10 <sup>-4</sup>	1	6.851*10 <sup>-4</sup>	7.10	0.0237*
X <sub>1</sub> <sup>2</sup>	0.013	1	0.013	134.65	<0.0001***
X <sub>2</sub> <sup>2</sup>	1.213*10 <sup>-3</sup>	1	1.213*10 <sup>-3</sup>	12.56	0.0053**
X <sub>3</sub> <sup>2</sup>	3.322*10 <sup>-3</sup>	1	3.322*10 <sup>-3</sup>	34.42	0.0002***
X <sub>1</sub> X <sub>2</sub>	1.431*10 <sup>-3</sup>	1	1.431*10 <sup>-3</sup>	14.83	0.0032**
X <sub>1</sub> X <sub>3</sub>	4.061*10 <sup>-4</sup>	1	4.061*10 <sup>-4</sup>	4.21	0.0674 <sup>ns</sup>
X <sub>2</sub> X <sub>3</sub>	1.081*10 <sup>-3</sup>	1	1.081*10 <sup>-3</sup>	11.20	0.0074**
Residual	9.653*10 <sup>-4</sup>	10	9.653*10 <sup>-5</sup>		
Lack of Fit	7.165*10 <sup>-4</sup>	5	1.433*10 <sup>-4</sup>	2.88	0.1353 <sup>ns</sup>
Pure Error	2.488*10 <sup>-4</sup>	5	4.977*10 <sup>-5</sup>		
Cor Total	0.024	19			
Std. Dev.= 9.825*10 <sup>-3</sup>		R-Squared=0.9603			
Mean=0.59		Adj R-Squared=0.9246			
C.V.= 1.67		Pred R-Squared=0.7229			
PRESS=6.737*10 <sup>-3</sup>		Adeq Precision=15.910			

X<sub>1</sub>: pH; X<sub>2</sub>: Temperature (°C); X<sub>3</sub>: Toluene concentration (mg.l<sup>-1</sup>)

\*Values of "Probability>F value" less than 0.05 indicate model terms are significant

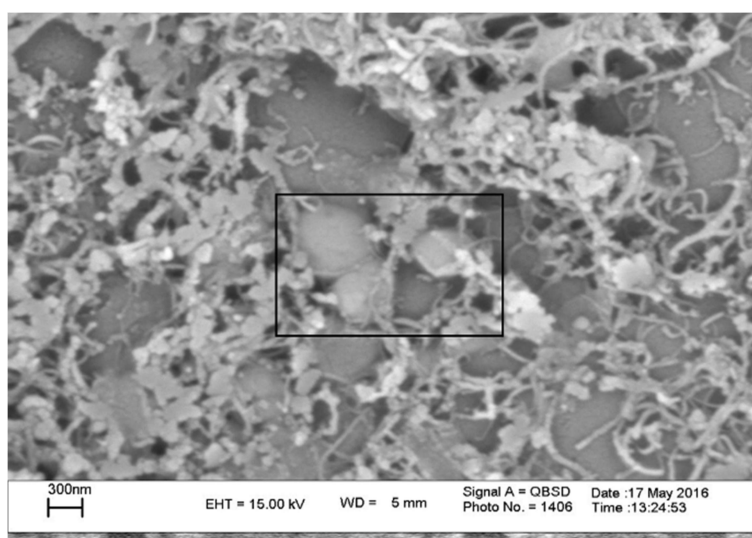
Source: The Author

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140 SEM OBSERVATIONS AND FT-IR ANALYSIS

141 The bacterial adhesion on the surface of MWCNTs in the presence of  $100 \text{ mg.l}^{-1}$  toluene was  
142 observed using SEM. Fig. 2 demonstrates that bacteria cells are trapped among the bundles of  
143 MWCNTs arrays. It can be due to the interactions of bacteria cells with the external surfaces of  
144 MWCNTs arrays. In addition, Fig. 2 indicates no major changes in the morphology of the bacteria cells  
145 after incubating with MWCNTs arrays. These SEM images reveal that MWCNTs clusters only capture  
146 the bacteria cells due to sieving mechanisms without any damage to the cell wall.

**Figure 02.** Scanning electron microscopy imagery of immobilized cells of *Staphylococcus gallinarum* ATHH41 with MWCNTs



Source: The Author

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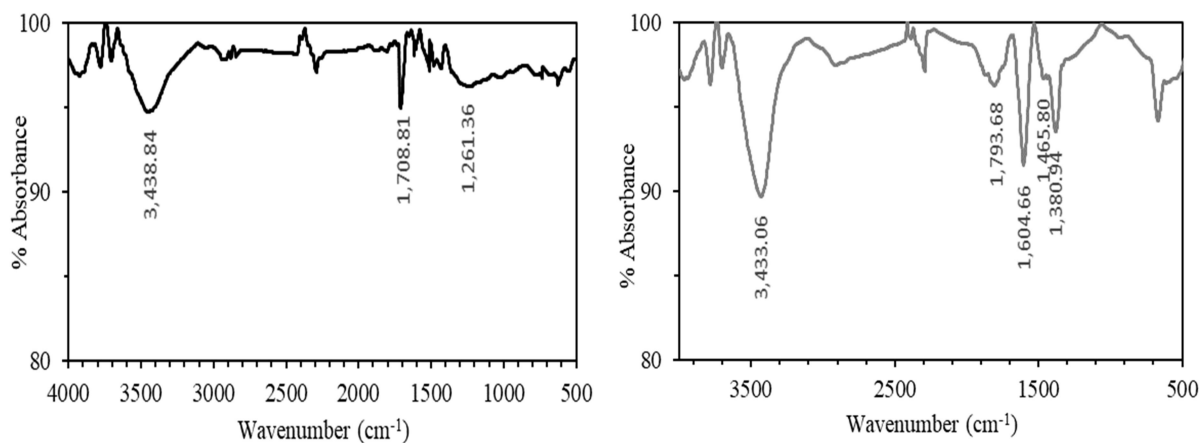
148 The whole spectrum of MWCNTs and MWCNTs/*Staphylococcus gallinarum* are compared in  
149 Fig. 3. The peak appearance in the areas about  $1708.81 \text{ cm}^{-1}$  can be ascribed to functional groups  
150 containing C=O stretching bond and the peak observed near  $3438.84 \text{ cm}^{-1}$  is attributed to the band  
151 vibration of O-H (Fig. 3A).

152 The peak appearance in the areas about  $3433.06 \text{ cm}^{-1}$  is attributed to the band vibration of O-  
153 H and the peak observed near  $1604.66 \text{ cm}^{-1}$  can be ascribed to functional groups containing C=O  
154 stretching bond. The peak appearance in the areas about  $1380.94$  and  $1465.80 \text{ cm}^{-1}$  is attributed to the  
155 band vibration of C-O and C-N stretching mode. In addition, the peak appearance near  $1793.68 \text{ cm}^{-1}$   
156 can be ascribed to functional groups containing C=O stretching bond (Fig. 3B). This peak is revealing  
157 of the presence of the functional groups and bacterial strain on the MWCNTs surface that have  
158 designed duration the formation MWCNTs/*Staphylococcus gallinarum* and purification processes. The  
159 peak observed near  $1798.68 \text{ cm}^{-1}$  can be ascribed to functional groups containing C=O single bond.



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**Figure 03.** FT-IR analysis of the A) carboxylate multi-walled carbon nanotubes, B) MWCNTS/*Staphylococcus gallinarum*

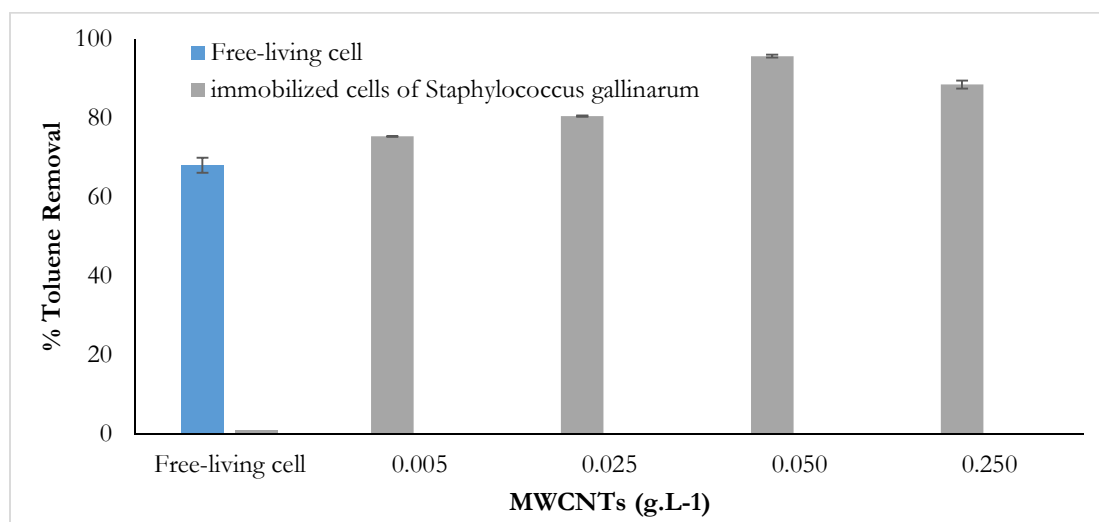


Source: The Author

160 **ADSORPTION PERFORMANCE OF FREE-LIVING AND IMMOBILIZED CELLS**

161 The removal of 636.04 mg.l<sup>-1</sup> toluene by the free and immobilized cells of *Staphylococcus*  
 162 *gallinarum* ATHH41 with different concentrations of MWCNTs under an initial pH of 7.68 and the  
 163 temperature of 31.73°C during 24 h, and shaking at 150 rpm were studied (Fig. 4). In addition, a higher  
 164 toluene removal percentage was achieved by immobilized cells by 0.05 g.l<sup>-1</sup> MWCNTs. Another thing  
 165 about the effect of carbon nanotubes was that carbon nanotubes at low concentrations had reverse  
 166 effect on a high concentration. In this study, toluene, at pH of 7.68, temperature of 31.73°C, and an  
 167 initial concentration of 636.04 mg.l<sup>-1</sup> was considerably degraded by 68.01% by the free-living cells and  
 168 up to 95.68% by immobilized cells of *Staphylococcus gallinarum* ATHH41 (Fig. 4).

**Figure 04.** The comparison of toluene removal percentage by free-living cells and immobilized cells of *Staphylococcus gallinarum* ATHH41 with MWCNTs



Source: The Author

169

## 170 **DISCUSSION**

171 This study investigated the biodegradation of toluene by free and immobilized *Staphylococcus*  
172 *gallinarum* strain ATHH41 and the following conclusions were drawn: Toluene degrading bacterium  
173 with high biodegradation activity and high tolerance of toluene, *Staphylococcus gallinarum* strain ATHH41,  
174 was isolated from the oil-contaminated soils. The genus *Staphylococcus* is gram positive with a thick  
175 peptidoglycan layer in the cell wall, and it shows tolerance to organic solvents, such as toluene,  
176 benzene, and xylene (Torres et al. 2011). This strain was capable of removing toluene from liquid  
177 mineral salt medium by 68.011% in 24 h. Multi-walled carbon nanotubes (MWCNTs) were used to  
178 immobilize the strain ATHH41. No changes have been observed in other studies in the structures of  
179 the carbon nanotubes after bacterial immobilizing, which is the benefit of the method. Using non-array  
180 CNTs have shown that CNTs rupture cell wall-membrane due to toxicity mechanisms, such as  
181 oxidative stress and physical damage (Kolangikhah et al. 2012) while this observation has not been  
182 observed here. The immobilized cells possess better storage stability and could remove toluene by  
183 95.68% in 0.05 g.l<sup>-1</sup> MWCNTs during 24 h. Based on the results, it is evident that the toluene  
184 degradation by immobilized bacteria is higher than by bacteria alone. The interesting point was that in  
185 spite of the increase in nanotubes concentration and the degradation effect, it was not linear and  
186 regular. The adsorption mechanism of toluene on MWCNTs is essentially ascribed to the  $\pi$ - $\pi$  electron  
187 donor-acceptor interaction among the aromatic ring of toluene and the surface carboxylic groups of  
188 MWCNTs. Positively charged toluene molecules attract the negatively charged molecules such as  
189 carbon nanotubes. Carbon nanotubes are effective adsorbent of BTX compounds and have a good  
190 potential for the removal of BTX compounds from the wastewater (Bina et al. 2012). Pang et al. (2011)  
191 showed that immobilized *Pseudomonas aeruginosa* with multi-walled carbon nanotubes (MWCNTs) were  
192 able to increase the absorption of Cr(VI) and the repeated operation of them. The MWCNTs show  
193 better toluene adsorption efficiency in 0.05 g.l<sup>-1</sup> MWCNT. When the MWCNTs contents were more  
194 than 0.05 g.l<sup>-1</sup>, toluene degradation would be decreased because of the toxicity of MWCNTs. Also, high  
195 MWCNTs contents cause a certain degree of inhibition to the microbial cells. The antimicrobial nature  
196 of CNTs depends on multiple variables related to their physical structure and composition. The  
197 exposure of microbes to CNTs induces severe oxidative stress in microbes pursued by cell membrane  
198 hurt and the release of internal cell contents (Kolangikhah et al. 2012). Thus, the establishment of  
199 proficient contact between the CNTs and bacterial cell surface determines the biocidal action of CNTs.  
200 However, this effort depends on a variety of factors, such as: (i) physical and structural properties of

201 CNTs (size and length); (ii) physical condition of CNTs (aggregated or dispersed); (iii) type and  
202 concentration of infections associated with CNTs and their availability to bacteria (heavy metal  
203 impurities); and (iv) number of layers (single or multi-walled) of CNTs. Normally, loosely-packed,  
204 debund-led, highly-dispersed, and shorter length tubes can easily penetrate through the cell membrane  
205 and display higher cell cytotoxicity (Al-Jumaili et al. 2017).

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268 **Degradação De Tolueno Por Staphylococcus Gallinarum Livre E**  
269 **Imobilizado Em Nanotubos De Carbono Multi-Carregados**

270 **Resumo**

271 A poluição por hidrocarbonetos é uma preocupação ambiental e de saúde mais importante. Usar  
272 bactérias livres e imobilizadas pode ser uma atitude adequada para encontrar um agente de  
273 bioaumentação adequado. Uma bactéria degradadora de tolueno foi isolada de ambientes contaminados  
274 com óleo (localizado em Bandar-Anzali, Guilan, Irã). A cepa foi identificada molecularmente como  
275 Staphylococcus gallinarum ATHH41 (número de acesso: KX344723) por sequenciamento parcial do  
276 gene 16SrDNA. A metodologia de superfície de resposta (RSM) foi empregada para biodegradação do  
277 tolueno por ATHH41. O projeto composto central (CCD) foi utilizado para otimizar o pH, a  
278 temperatura e a concentração de tolueno por ATHH41. De acordo com o propósito de otimização do  
279 software Design-Expert, as condições ótimas de degradação do tolueno foram obtidas quando o pH, a  
280 temperatura e a concentração de tolueno foram ajustados para 7.68, 31.73 ° C e 630,04 mg.l-1 ,  
281 respectivamente. Nanotubos de carbono de paredes múltiplas (MWCNTs) foram usados para  
282 imobilizar a cepa. A espectroscopia de infravermelho e a microscopia eletrônica de varredura  
283 mostraram que as células aderiram à superfície MWCNT e desenvolveram um biofilme. Os resultados  
284 revelaram que as células livres foram capazes de degradar 68.01% do tolueno como única fonte de  
285 carbono e energia em 24 horas sob condições otimizadas. As células imobilizadas atingiram 95.68%.

286 **Palavras-chave:** Nanotubo de carbono; Metodologia de superfície de resposta; Staphylococcus  
287 gallinarum ATHH41; Tolueno

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