

## THE DEGRADATION OF DIESEL OIL BY CONSORTIUM OF BACTERIA IN SHAKEFLASK CULTURES

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### **Abstract**

*Diesel fuel is widely used in manufacturing and transportation industries. Its production, transportation, use and disposal have the risk to contaminate the environment. In the other hand contaminated site usually can produce some microorganisms, through natural selection, that can utilize the hydrocarbons as source of energy and food thus clean the environment itself. For bioremediation purposes, 14 isolates of diesel oil degrading bacteria from Bandung, Jakarta, and Bali has been employed for diesel oil degradation. These 14 isolates, together with other variables, e.g. pH, Pb, sucrose and urea, were plotted in 23x24 matrixes of Plackett and Burman statistical method. Fermentation was carried out in batch cultures for 14 days. The results showed that 9 isolates and the availability of sucrose and higher initial pH have a positive/stimulatory effect on diesel oil biodegradation, 5 of which were very significant (99.99%), whereas 1 isolate has inhibitory effect (-98%). The results also revealed that 8 isolates, higher initial pH, and availability of urea have stimulatory effect on culture growth (more than 90% significant). The present of Pb in medium inhibit both growth and the degradation of diesel oil, although the effect was not significant (less than 60%).*

*Keywords: biodegradation, diesel oil, bacteria isolates, shakeflask cultures*

### **1. Introduction**

The pollution by diesel oil occurred in many places (petrol stations, workshops, factories, oil refinery plant, harbour), not counted the huge pollution by oil spill from tanker (Bragg et al, 1994). In Indonesia, the pollution by diesel oil in Tanjung Periok Harbour was 900 L per ship with oil content of 12,95 mg/L, this figure will count to 0.1 to 363 ton in year 2000 (Irawati, 1994). Similar condition is probably occurred in every

harbour in Indonesia (Feliatra, 1996, Udiharto, 1992), whereas in fact diesel oil contamination is regarded as toxic and hazardous pollutant (RI Governmental Regulation no.18, year 1999, code D1003d). To overcome oil pollution, bioremediation is the most environmental friendly way (Kadarwati et al, 1996) and many organisms can be employed for this purpose (Vieira et al, 2007, Al-Ghazawi et al, 2005, Edvotek, 2002). Many factors influence the efficiency of the

bioremediation (Barathi and Vasudevan, 2003, Barenholz et al., 2003, Boopathy, 2003, Bieszkiewicz et al., 2002, Oh et al., 2001). The aim of this research is to investigate the factors affecting the level of biodegradation of diesel oil by consortium of bacteria using statistical method. Consortium of bacteria have been proven effective in many petroleum hydrocarbon biodegradation work (Rahman et al., 2003, Al-Awadhi et al., 2003, Ciawi and Santi, 2000). We have isolated diesel oil degrading bacteria from oil contaminated sites in Bandung (Safitri et al, 2002 and Ciawi et al, 1999), Jakarta and Bali (Ciawi et al, 2003). The experiments employed the statistical method Plackett Burman, which has been used successfully in determining the factors affecting the level of metabolite production by microbes (Castro et al, 1992). This method is more economical and time saving compared with factorial method (Stose and Mayer, 1966).

## 2. Materials and Methods

All chemicals used in this research are pro analyst grade, except for urea, sucrose and diesel oil. The medium used for fermentations was a modification of Bussnell-Haas mineral salts medium which consist of 0.2 gr.l<sup>-1</sup> MgSO<sub>4</sub>, 0.02 gr.l<sup>-1</sup> CaCl<sub>2</sub>, 1.0 gr.l<sup>-1</sup> KH<sub>2</sub>PO<sub>4</sub>, 1.0 gr.l<sup>-1</sup> K<sub>2</sub>HPO<sub>4</sub>, 1.0 gr.l<sup>-1</sup> NH<sub>4</sub>NO<sub>3</sub>, 2 drops of 60% FeCl<sub>3</sub> solution, 1000 ppm sucrose, 1000 urea, 1.0 gr.l<sup>-1</sup> PbNO<sub>3</sub>. The microorganisms employed were *Cytophaga fermentans*, *Methylomonas pelagika*, *Micrococcus tractuosa*, *Mesophilobacter marinus*, *Marinomonas comunis*, *Brevibacterium*, *Staphylococcus*, *Flavobacterium*, *Pseudomonas*, *Mocillercilla*, *Pseudomonas putida*, *Microscilla*, *Alcaligenes*, and *Bacillus*.

The experiment was designed according to Plackett and Burman method (Castro et al, 1992, Monaghan and Koupal, 1989), by using a matrix of 24 x 23 which is shown in the Table 1.

Table 1. Plackett Burman statistical design by using 24 experiments and 23 variables including 5 dummies.

Experiment	Variable																						
	A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q	R	S	T	W	X	Y
1	+	+	+	+	+	-	+	-	+	+	-	-	+	+	-	-	+	-	+	-	-	-	-
2	+	+	+	+	-	+	-	+	+	-	-	+	+	-	-	+	-	+	-	-	-	-	+
3	+	+	+	-	+	-	+	+	-	-	+	+	-	-	+	-	+	-	-	-	-	+	+
4	+	+	-	+	-	+	+	-	-	+	+	-	-	+	-	+	-	-	-	-	-	+	+
5	+	-	+	-	+	+	-	-	+	+	-	+	-	+	-	+	-	-	-	-	+	+	+
6	-	+	-	+	+	-	-	+	+	-	-	+	-	+	-	-	-	-	+	+	+	+	+

7	+	-	+	+	-	-	+	+	-	-	+	-	+	-	-	-	+	+	+	+	+	-	
8	-	+	+	-	-	+	+	-	-	+	-	+	-	-	-	-	+	+	+	+	+	-	+
9	+	+	-	-	+	+	-	-	+	-	+	-	-	-	-	+	+	+	+	+	-	+	-
10	+	-	-	+	+	-	-	+	-	+	-	-	-	-	+	+	+	+	+	-	+	-	+
11	-	-	+	+	-	-	+	-	+	-	-	-	-	+	+	+	+	+	-	+	-	+	+
12	-	+	+	-	-	+	-	+	-	-	-	-	+	+	+	+	+	-	+	-	+	+	-
13	+	+	-	-	+	-	+	-	-	-	-	+	+	+	+	+	-	+	-	+	+	-	-
14	+	-	-	+	-	+	-	-	-	-	+	+	+	+	+	-	+	-	+	+	-	-	+
15	-	-	+	-	+	-	-	-	-	+	+	+	+	+	-	+	-	+	+	-	-	+	+
16	-	+	-	+	-	-	-	-	+	+	+	+	+	-	+	-	+	+	-	-	+	+	-
17	+	-	+	-	-	-	-	+	+	+	+	+	-	+	-	+	+	-	-	+	+	-	-
18	-	+	-	-	-	-	+	+	+	+	+	-	+	-	+	+	-	-	+	+	-	-	+
19	+	-	-	-	-	+	+	+	+	+	+	-	+	-	+	+	-	+	+	-	-	+	-
20	-	-	-	-	+	+	+	+	+	+	-	+	-	+	+	-	+	+	-	+	-	+	+
21	-	-	-	+	+	+	+	+	-	+	-	+	+	-	-	+	+	-	-	+	-	+	-
22	-	-	+	+	+	+	+	-	+	-	+	+	-	-	+	+	-	-	+	-	+	-	-
23	-	+	+	+	+	+	-	+	-	+	+	-	-	+	+	-	-	+	-	+	-	-	-
24	+	+	+	+	+	-	+	-	+	+	-	-	+	+	-	-	+	-	+	-	-	-	-

Note : *Cytophaga fermentans* (variable A), *Methylomonas pelagika* (B), *Micrococcus tractuosa* (C), dummy (D), *Mesophilobacter marinus* (E), *Marinomonas comunis* (F), *Brevibacterium* (G), dummy (H), *Staphylococcus* (I), *Flavobacterium* (J), *Pseudomonas* (K), dummy (L), *Mocillercilla* (M), *Pseudomonas putida* (N), *Microscilla* (O), dummy (P), *Alcaligenes* (Q), *Bacillus* (R), initial pH (S), Pb (U), urea (V), dan sucrose (W).

The parameter used here was the amount of oil degraded. The effect of each variable were calculated from:

$$E_a = R \text{ at } (+)/n - R \text{ at } (-)/n$$

Where: R = the response (amount of oil degraded)

$$N = \text{number of experiments (24)}$$

The experimental error was determined from:

$$V_E = \Sigma(E_d)^2/d$$

Where d = number of dummy variables and  $E_d$  = the dummy effect.

The standard error of the effects was determined as:

$$S.E.\text{-eff} = \sqrt{V_E}$$

Finally, the significant level of each variable were determined using Student's t-test:

$$t\text{-value} = E/S.E.\text{eff}$$

#### *Ammonia determination*

The method used was similar to the method described by Fawcett and Scott (1960). Two reagents were prepared in dark bottles. Solution A contained 55.6 g.l<sup>-1</sup> phenol and 0.3 g.l<sup>-1</sup> sodium nitroprusside. Solution B contained 0.11 M sodium hypochlorite and 20 g.l<sup>-1</sup> sodium hydroxide. This method will give linear response at concentration between 0 and 2 mM. Equal amount of samples and each solution were mixed thoroughly and incubated in the dark at 37<sup>0</sup>C for 15 minutes. The optical density was read at 570nm.

### *Sucrose determination*

The method used was similar to the method described in Robyt and White (1990). 0.1% anthrone was dissolved in 100 ml concentrated sulfuric acid. This method will give linear response at concentration between 0 and 0,2 mg/ml glucose. 1 ml of samples or standard solutions was mixed thoroughly with 5 ml of freshly made anthron solution, heated at 100<sup>0</sup>C for 12 minutes and cooled under running water. The optical density was read at 630 nm.

### *Growth determination*

The growth level was measured as optical density at 600 nm. Samples were diluted to result an OD readings below 1.00.

### *Oil determination*

Amount of residual oil was measured simply using measuring cylinders. Level of biodegradation was counted by subtracting residual oil from initial amount of oil.

## **3. Results**

### *a. The effect of experimental variable on the degradation of diesel oil*

The results of 24 experiments using Plackett dan Burman statistical design are shown in the following tables. Table 2 shown the effect of each variables on the level of oil degradation. There are 12 variables that significantly affect (>90%) the degradation of diesel oil, 5 of

which are more than 98% significant, i.e. *Cytophaga* sp., *Micrococcus* sp., *Mesophilobacter* sp., *Staphylococcus* sp., and *Alkaligenes* sp.. The 12 variables are *Cytophaga fermentans* (variable A), *Micrococcus tractuosa* (C), *Mesophilobacter marinus* (E), *Marinomonas comunis* (F), *Staphylococcus* sp. (I), *Flavobacterium* sp. (J), *Mocillercilla* sp. (M), *Pseudomonas putida* (N), *Alcaligenes* sp. (Q), initial medium pH (S), and sucrose (W), all of which are positively affect the degradation of diesel oil, whereas *Brevibacterium* sp. (variable G) negatively affect the degradation of diesel oil. Data in Table 2 also show that the effect of control variables (dummies) are also very small, which mean that overall experimental error is quite small (%s <50%, except for dummy D, 80%). Another interesting point is that *Bacillus* sp. negatively affect the degradation of diesel oil, although it is not very significant, which is 60%. Moreover, *Brevibacterium* sp. which was isolated from soil samples in Bandung negatively affect the degradation (98%). Also, in this research, there are 2 *Pseudomonas* isolates which have different effect on diesel oil degradation. It might be that the two isolates are different species. The effect of *Pseudomonas putida* is quite

significant (90%) and the effect of the other *Pseudomonas* is only 60% significant. Many *Pseudomonas* species has been found has the ability to degrade various toxic substances

such as diesel oil (Das and Mukherjee, 2007, Pepi et al., 2003), phenol (Wang and Lie, 2007), chlorotoluene (Lehning et al., 1997).

**Table 2 The effect of each experimental variable on the degradation of diesel oil using statistical analysis on experimental results**

Variable	A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q	R	S	T	U	V	W
% S	99,9	80	99	80	99,9	95	-98	60	99	95	60	-60	95	90	60	<50	99,9	-<50	95	<50	-<50	<50	98

Notes: *Cytophaga fermentans* (variable A), *Methylomonas pelagika* (B), *Micrococcus tractuosa* (C), dummy (D), *Mesophilobacter marinus* (E), *Marinomonas comunis* (F), *Brevibacterium* (G), dummy (H), *Staphylococcus* (I), *Flavobacterium* (J), *Pseudomonas* (K), dummy (L), *Mocillercilla* (M), *Pseudomonas putida* (N), *Microscilla* (O), dummy (P), *Alcaligenes* (Q), *Bacillus* (R), initial pH (S), Pb (U), urea (V), and sugar (W).

*The effect of experimental variable on the growth of the cultures*

The results of statistical analysis on the experimental data showed that the effect of

each variables on the growth of the consortia of microbes was different from its effect on the level of oil degradation (Table 3 and Figure 1).

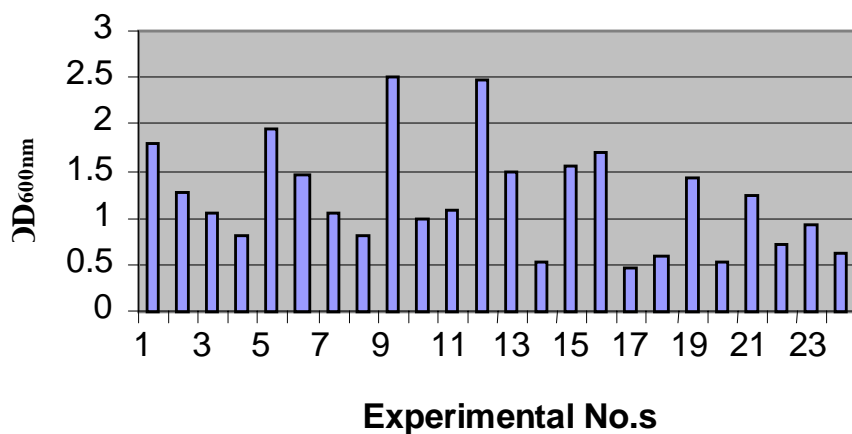
**Table 3. The effect of each experimental variable on the growth of the culture of bacteria consortia using statistical analysis on experimental data**

Variable	A	B	C	D	E	F	G	H	I	J	K	L	M	N	O	P	Q	R	S	T	U	V	W
% S	95	99,9	90	!-<50	99	60	-90	-80	95	<50	-99	-70	99	60	<50	60	90	70	98	!-<50	-60	99,9	-98

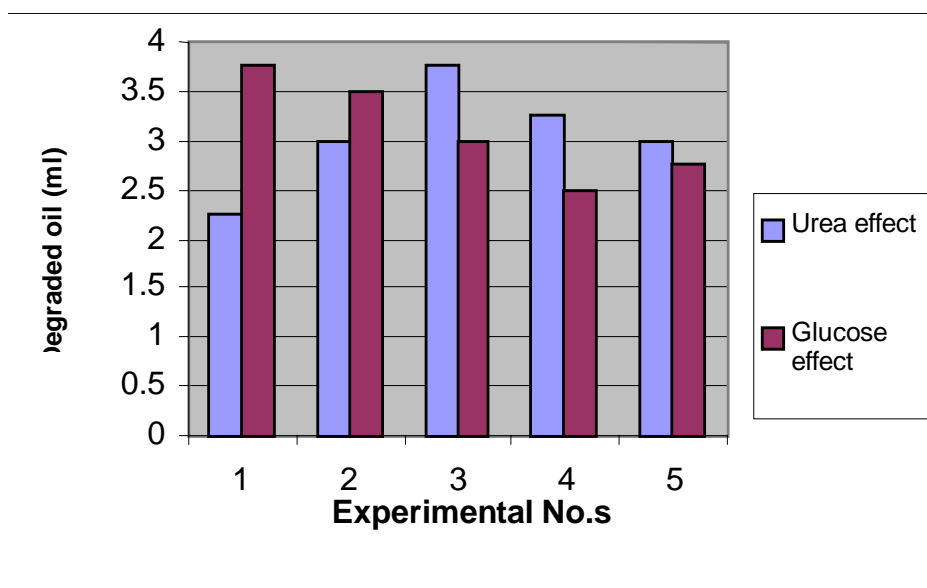
Notes: *Cytophaga fermentans* (variable A), *Methylomonas pelagika* (B), *Micrococcus tractuosa* (C), dummy (D), *Mesophilobacter marinus* (E), *Marinomonas comunis* (F), *Brevibacterium* (G), dummy (H), *Staphylococcus* (I), *Flavobacterium* (J), *Pseudomonas* (K), dummy (L), *Mocillercilla* (M), *Pseudomonas putida* (N), *Microscilla* (O), dummy (P), *Alcaligenes* (Q), *Bacillus* (R), initial pH (S), Pb (U), urea (V), and sugar (W).

The most significant variable (99%) on the growth of the consortia was *Methylomonas pelagika* (B), *Mesophilobacter marinus* (E), *Mocillercilla* (M), *Pseudomonas* (K), and urea (V). In this experiment, *Pseudomonas* sp. showed a negative effect on growth. Urea has positive effect on the growth of the consortia probably because the availability of urea provide more nitrogen in the media. In this

experiment it was 1000 mM, which is the maximum concentration for the consortia to grow, which was proven in subsequent experiment (Figure 2 and Figure 3), in the range between 0-5000 ppm, the best concentration of urea that produced the highest growth and the highest degradation level is 1000 ppm.



**Figure 1** The level of growth (measured as OD 600nm) of the bacteria consortia in 24 experiments

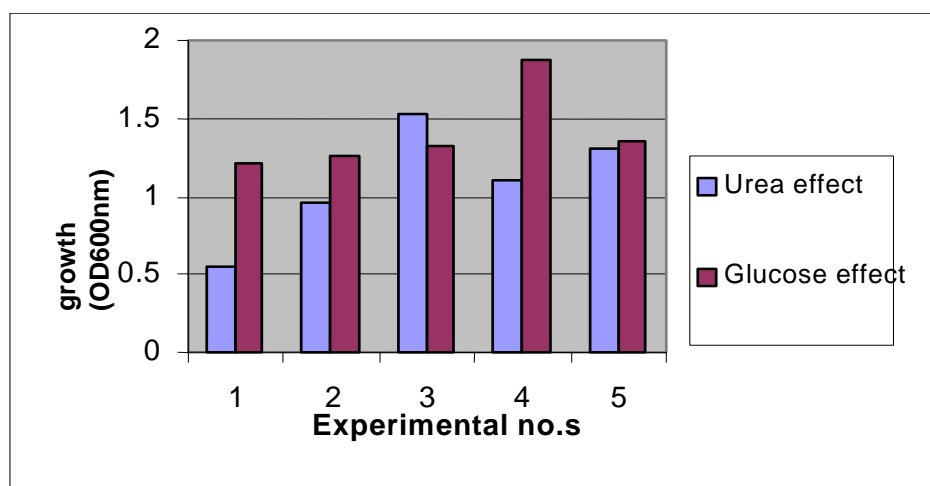


**Figure 2 The effect of urea and sucrose concentration on the degradation of diesel oil**

Note: urea, experiment 1-5, [glucose]=1000 ppm, experiment 1, urea=0 ppm, exp. 2, urea=500 ppm, exp.3, urea=1000 ppm, exp.4, urea=3000 ppm, exp.5, urea=5000 ppm. Glucose effect, [urea]= 3000 ppm, exp.1, glucose=0 ppm, exp.2 glucose=100 ppm, exp.3, glucose=500 ppm, exp.4, glucose= 1000 ppm, exp.5, glucose=5000 ppm.

The negative significant effect of sucrose on growth (-98%) is contradictory with the results of effect of glucose on growth in many experiment with other bacteria (e.g. Ciawi, 1996) and also with our previous research (Ciawi et al., 2000, Ciawi dan Santi, 2000), which showed that glucose and  $\text{NH}_4\text{Cl}$  has significant positive effect on growth as well as oil degradation. We expect a different result with the assumption that, as carbon source, sucrose is more readily to be consumed than

diesel oil. In contrast, the result of the effect of sucrose on oil degradation agreed with our previous experiment. At subsequent experiment, it showed that at the range between 0-5000 ppm sucrose, the best concentration for diesel oil degradation is 0 ppm (Figure 2) and 1000 ppm for growth (Figure 3). This research was continued by investigating the effect of the fermentation systems on the degradation level of diesel oil.



**Figure 3. The effect of urea and glucose concentration on the growth of the consortia.**

Note: urea, experiment 1-5, [glucose]=1000 ppm, experiment 1, urea=0 ppm, exp. 2, urea=500 ppm, exp.3, urea=1000 ppm, exp.4, urea=3000 ppm, exp.5, urea=5000 ppm. Glucose effect, [urea]= 3000 ppm, exp.1, glucose=0 ppm, exp.2 glucose=100 ppm, exp.3, glucose=500 ppm, exp.4, glucose= 1000 ppm, exp.5, glucose=5000 ppm.

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