



Alexandria University
Faculty of Agric. (Saba-Bacha)

**GENETICAL STUDIES ON SOME BACTERIAL
GENERA AND ITS APPLICATION IN
BIODEGRADATION FOR WASTEWATER**

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For the Degree of

**MASTER OF AGRICULTURAL SCIENCE
(GENETICS)**

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دراسات وراثية على بعض الاجناس البكتيرية وتطبيقاتها فى الهدم الحيوى لمخلفات الصرف الصحى

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BY

SUZAN MOHAMMED HASSAN MOHAMMED HASSAN

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Table (7): DNA fragments of RAPD for treated and untreated *Bacillus* bacteria using four random primers.

Primers DNA fragments	AATCCGCAAT					CGCGGAATGC					AATCGGTACT					CTCGAATGCA					
	B	B1	B2	B3	H	B	B1	B2	B3	H	B	B1	B2	B3	H	B	B1	B2	B3	H	
1600																					
1550													+		+						
1500																					
1450																					
1350																				+	
1250					+																
1200					+	+						+									
1180												+									
1150	+	+			+								+	+	+		+		+	+	
950																+				+	
900		+					+														
850		+	+			+	+	+	+												
800	+		+						+	+											
750	+		+														+	+			
700		+	+								+	+	+	+	+						
680	+		+	+	+		+	+	+	+											
600	+					+	+														
550																+		+	+		
500			+				+														
450		+			+						+							+			
350													+								
300												+		+							
280																		+			

Where: B: *Bacillus* sp. without addition of ZnSO₄ in their growth medium, B1: *Bacillus* sp. under the treatment of 0.5 mg/L of ZnSO₄, B2: *Bacillus* sp. under the treatment of 1.0 mg/L of ZnSO₄, B3: *Bacillus* sp. under the treatment of 1.5 mg/L of ZnSO₄, and H: Hybrid between *Bacillus* sp. and *Pseudomonas* sp.

Table (6): DNA fragments of RAPD for treated and untreated *Pseudomonas* bacteria using four random primers.

Primers DNA fragments	AATCCGCAAT					CGCGGAATGC					AATCGGTACT					CTCGAATGCA				
	P	P1	P2	P3	H	P	P1	P2	P3	H	P	P1	P2	P3	H	P	P1	P2	P3	H
1600													+							
1550															+	+				
1500																				
1450													+							
1350													+							
1250					+			+												
1200					+			+	+											
1180																				
1150	+	+			+	+			+		+	+			+					+
950																				
900	+	+																		
850		+	+	+		+		+	+						+			+		
800		+	+	+						+										
750			+					+	+											
700	+			+		+					+	+		+	+			+		
680	+	+	+		+			+	+	+										
600						+	+													
550																				
500																+				
450			+		+													+		
420																				
350	+															+				
280																	+			
220																+				

Where:

P: *Pseudomonas* sp. without addition of PbNO₃ in their growth medium.

P2: *Pseudomonas* sp. under the treatment of 1.0 mg/L of PbNO₃.

H: Hybrid between *Bacillus* sp. and *Pseudomonas* sp

P1: *Pseudomonas* sp. under the treatment of 0.5 mg/L of PbNO₃.

P3: *Pseudomonas* sp. under the treatment of 1.5 mg/L of PbNO₃.

CHAPTER 1

INTRODUCTION

Wastewater treatment was the most important problem in recent years. Thus, advances in molecular biology have extended our understanding of the metabolic processes related to microbial transformation of heavy metals and petroleum hydrocarbons (**Puntambekar and Ranjeker, 1989** and **Mezzanotte, 2002**). By establishing conditions which maximize rates and extents of microbial growth, hydrocarbon access, and transformation, highly accelerated and bioreactor-based petroleum waste degradation processes have been implemented. Microbes may be exploited to break stable oilfield emulsions to produce pipeline quality oil. Molecular approaches are being used to broaden the substrate specificity and increase the rates and extents of desulfurization. Bacterial processes are being commercialized for removal of H₂S and sulfoxides from petrochemical waste streams. Microbes also have potential for use in removal of nitrogen from crude oil leading to reduced nitric oxide emissions provided that technical problems similar to those experienced in biodesulfurization can be solved. Thus, new strains of *Bacillus* were presented that produce lipase and other enzymes for the degradation of oleaginous materials such as fats, greases and cooking oils, and protease enzymes to degrade proteins. This novel strain and the enzymes produced thereby have a number of applications, including wastewater treatments, agricultural uses, laundry and dish detergents, drain cleaners and spot removers, among others (**Kozo et al., 1979; Raquel et al., 2007 and Reshma et al., 2007**).

Novel mutant strain *Pseudomonas fluorescens* 3P has been found to be capable of degrading and removing anionic and/or non-ionic surface active agents, detergents and like materials. Thus, this mutant could be cultured in wastewater from any type of industrial plant containing anionic and/or non-ionic surface active agents. The mutant strain of this invention can be employed in ion exchange resin treatment systems, in tricking filter systems, in carbon adsorption systems, in activated sludge treatment systems, in outdoor lagoons or pools, etc. (**James, 2006**). Molecular biology technique presented a huge helpful for wastewater treatment, **Wikstrom (2004)** demonstrated that microbial community changes over time in a nitroaromatic-contaminated groundwater upon amendment with hydrocarbons previously unknown to the microbial community (extrinsic) and hydrocarbons previously known to the microbial community (intrinsic).

Microbial growth, biodegradation, and community structure changes measured by random amplified polymorphic DNA (RAPD) and quantitative PCR (qPCR) targeting catechol-2,3-dioxygenase (C23O) genes were monitored over time. All amended substrates were biodegraded after both substrate amendments except for 2,4-DNT, which was only partially degraded after the second amendment. Unique microbial communities were developed in flasks amended with phenol, benzoic acid, and naphthalene. In addition protoplast fusion technique was applied for enhancement hereditary desirable genotype between different bacterial genera. **Kanevski et al. (1983)** carried out hybridization by fusion between protoplasts of *A. tumefaciens* and different *Pseudomonas* sp. All the eight strains which selected on screening media were true chromosomal hybrid and hereditary tumor formation as a character of *A. tumefaciens*, but nine strains which were selected by

mechanical isolation were resistant for two different pesticides as a remark of expressing, the *Pseudomonas* plasmids. **Gleba et al. (1986)** fused protoplasts of *Serratia sp.* with protoplasts of *Agrobacterium tumefaciens*. All the obtained 57-hybrid strains contained lysopine dehydrogenase, which is encoded by T-DNA, in both callus tissue and shoots. They found that, in six of 12 strains, chromosomes from both parents were present and found the nearly diploid chromosome numbers of both species, while in the other six cell lines, one of the parents was predominate. Esterase and amylase isoenzymes patterns and chloroplast DNA restriction analysis were detected. Furthermore, **Muller and Schieder (1987)** showed that protoplasts of a *Nicotiana paniculata* cell line could be fused with protoplasts of nitrate reductase deficient cnx68 cell line of *Pseudomonas*. From about 1.1×10^7 cultured protoplasts, 18 cell lines were obtained. Four of them showed two dominant and two recessive markers, while three cell lines lacked one or more markers.

The main objective of this study could be summarized as follow:

- 1- Isolation, purification and identification of bacterial genera characterized with their ability for wastewater biodegradation through different microbial tests.
- 2- Assay the ability of bacterial isolates from different genera for wastewater biodegradation.
- 3- Induction of a bacterial mutant which is superior in wastewater biodegradation comparing with its parents.
- 4- Improvement of wastewater biodegrading bacterial isolates under study through applied genetical technique (protoplast fusion technique).
- 5- Application of different molecular biological techniques (protein fingerprinting and Random Amplified Polymorphic DNA (RAPD) to study and compare parental and hybrid bacteria which are produced by protoplast fusion.