



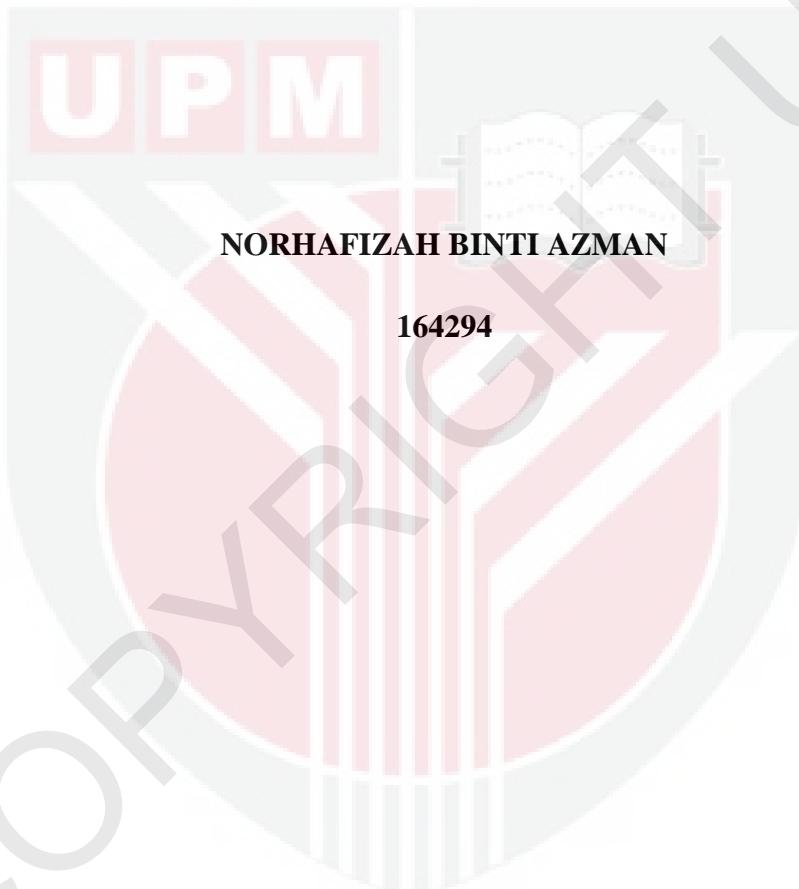
UNIVERSITI PUTRA MALAYSIA

**ISOLATION AND ANALYSIS OF THE *rhIA* AND *rhIB* FROM  
RHAMNOLIPID PRODUCING BACTERIUM *Pseudomonas aeruginosa*  
strains S5**

NORHAFIZAH AZMAN

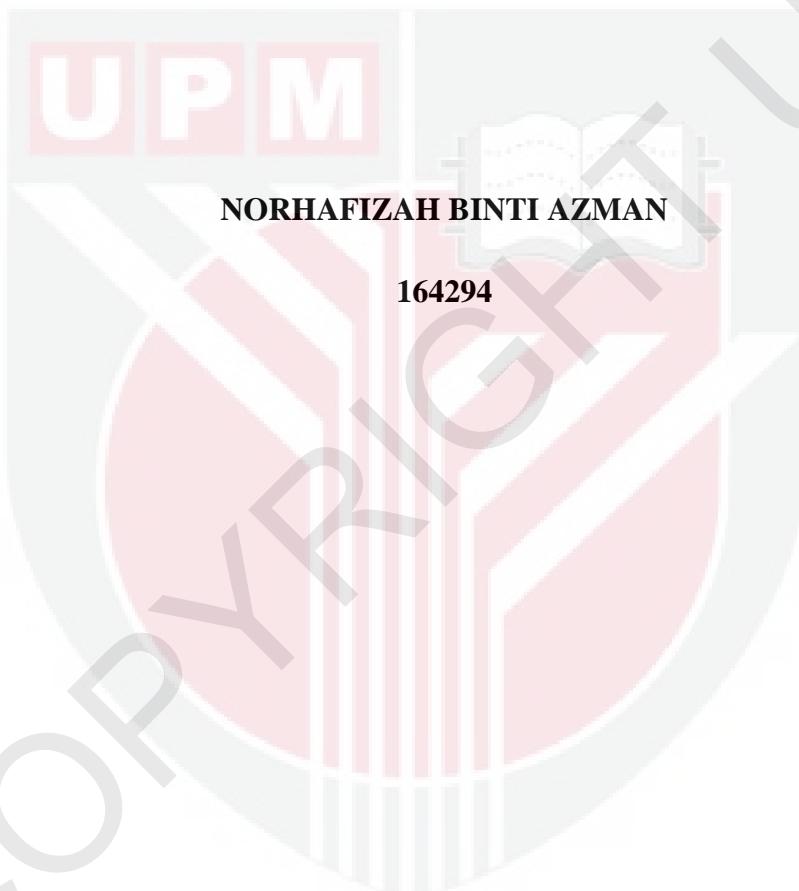
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**ISOLATION AND ANALYSIS OF THE *rhlA* AND *rhlB* FROM RHAMNOLIPID  
PRODUCING BACTERIUM *Pseudomonas aeruginosa* strains S5**



**Dissertation submitted in partial fulfillment of the requirement for the course  
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PRODUCING BACTERIUM *Pseudomonas aeruginosa* strains S5**



**DEPARTMENT OF MICROBIOLOGY  
FACULTY BIOTECHNOLOGY AND BIOMOLECULAR SCIENCES  
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## PENGESAHAN

Dengan ini adalah disahkan bahawa projek yang bertajuk “**Isolation and Analysis of the *rhlA* and *rhlB* From Rhamnolipid Producing Bacterium *Pseudomonas aeruginosa* strains S5**” telah disiapkan serta dikemukakan kepada Jabatan Mikrobiologi oleh **Norhafizah Binti Azman** dengan matrik nombor **164294** sebagai syarat untuk kursus BMY4999 projek.

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## ABSTRACT

Rhamnolipids are biotechnologically important biosurfactants, find applications in cosmetic and pharmaceutical industry, agriculture, environmental application and in bioremediation activities. *Pseudomonas aeruginosa* is one of the microbes that can produce rhamnolipids. The aim of this study is to test the rhamnolipids production of our *P. aeruginosa* strain and to analyze the genes required for the rhamnolipids biosynthesis. Based on the qualitative test on cetyltrimethyl ammonium bromide agar, *P. aeruginosa* strain S5 formed a blue halo around the culture which showed that the bacterium is a rhamnolipids producer. The emulsification index value of the rhamnolipids produced was 41%. Genes involved in rhamnolipids biosynthesis in *P. aeruginosa* strain S5; *rhlA* coding for rhamnosyltransferase 1 subunit A, and *rhlB*, coding for rhamnosyltransferase 1 subunit B were amplified, sequenced and analyzed. The sequencing results showed that the size of *rhlA* and *rhlB*, genes are 888 bp and 1281 bp, which corresponded to 295 and 426 amino acids protein, respectively. The genes sequence that were identified and analyzed could facilitate the understanding on the rhamnolipids biosynthesis in *P. aeruginosa* strain S5, hence the information can be further used in genetic modulation in order to produce natural biosurfactant in the economic yet safer manner which meets industrial and ecological prospect.

## ABSTRAK

Rhamnolipid adalah biosurfaktan yang penting dalam bioteknologi termasuklah pengunaannya dalam kosmetik, industri farmaseutikal, pertanian, alam sekitar dan aktiviti bioremediasi. *Pseudomonas aeruginosa* adalah salah satu daripada mikrob yang boleh menghasilkan rhamnolipid. Tujuan kajian ini adalah untuk mengkaji pengeluaran rhamnolipid dari strain *P. aeruginosa* kami dan untuk menganalisis gen yang diperlukan untuk biosintesis rhamnolipid itu. Melalui ujian kualitatif pada agar cetyltrimethylammonium bromide, *P. aeruginosa* strain S5 telah menghasilkan halo biru di sekeliling kultur yang menunjukkan bakteria ini adalah pengeluar rhamnolipid. Keputusan indeks pengemulsian yang terhasil daripada rhamnolipid adalah 41 %. Gen-gen yang terlibat dalam biosintesis rhamnolipid oleh *P. aeruginosa* strain S5; *rhlA* yang mengekod untuk rhamnosyltransferase 1 subunit A, dan *rhlB* yang mengekod untuk rhamnosyltransferase 1 subunit B telah diperbanyakkan, dijujukkan dan dianalisis. Keputusan penjujukan menunjukkan bahawa saiz gen *rhlA* dan *rhlB* adalah masing-masing 888 bp dan 1281 bp yang mengekod untuk masing-masing 295 dan 426 asid amino protein. Jujukan gen-gen yang telah dikenalpasti dan dianalsis akan membantu memahami biosintesis rhamnolipid dalam *P. aeruginosa* strain S5, dan seterusnya maklumat tersebut boleh digunakan dalam modulasi genetik untuk menghasilkan biosurfaktan semulajadi dengan cara yang lebih selamat dan ekonomi yang akan memenuhi prospek industri dan ekologi.

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## TABLE OF CONTENTS

PENGESAHAN	i
ABSTRACT	ii
ABSTRAK	iii
ACKNOWLEDGEMENT	iv
LIST OF ABBREVIATION	vii
LIST OF TABLES	viii
LIST OF FIGURES	ix
LIST OF APPENDICES	x
CHAPTER 1 INTRODUCTION	1
CHAPTER 2 LITERATURE REVIEW	3
2.1 Biosurfactant	3
2.2 Rhamnolipids	3
2.3 Application of rhamnolipids	4
2.3.1 Bioremediation	4
2.4 Biological control	5
2.4.1 Pharmaceutical products	5
2.5 Microbial production of rhamnolipids	5
2.6 Biosynthesis pathway of rhamnolipid	6
2.7 Genes encoding proteins for the rhamnolipids biosynthetic pathway	7
2.8 Role of RhlA and RhlB in rhamnolipid production	9
CHAPTER 3 MATERIALS AND METHODS	10
3.1 Chemical Preparation	10
3.2 Media Preparation	10
3.3 Bacterial Culture	10
3.3.1 Preparation of bacterial culture	10
3.3.2 Preparation of bacterial stocks	10
3.4 Bacterial Identification	11
3.5 Confirmation of rhamnolipid production in <i>P. aeruginosa</i> strain S5	11
3.5.1 Sample preparation	11
3.5.2 Supernatant extraction	11
3.5.3 Rhamnolipid detection test	12
3.6 Identification of rhlA and rhlB gene in <i>P. aeruginosa</i> S5	13
3.6.1 Isolation of genomic DNA from <i>Pseudomonas aeruginosa</i> strain S5	13
3.6.2 DNA Quantification	14
3.6.3 Primer Design	14
3.6.4 Amplification of <i>rhlA</i> and <i>rhlB</i> by Polymerase Chain Reaction	14
3.6.5 Electrophoresis and Detection of PCR product	15
3.6.6 DNA Sequencing	16
3.6.7 Sequence Analysis	16
CHAPTER 4 RESULTS AND DISCUSSION	17
4.1 Confirmation of <i>Pseudomonas aeruginosa</i> strain S5	17

4.2	Confirmation of rhamnolipids production	19
4.2.1	CTAB plate	19
4.2.2	Emulsification index	21
4.3	Isolation of genomic DNA	23
4.4	Successful amplification of <i>rhlA</i> and <i>rhlB</i>	24
4.5	Analysis of <i>rhlA</i> and <i>rhlB</i> sequence	25
4.6	Phylogenetic tree	33
CHAPTER 5 CONCLUSION AND RECOMMENDATIONS		35
REFERENCES		36
APPENDICES		40



## LIST OF ABBREVIATION

### Abbreviation

PAS5	<i>Pseudomonas aeruginosa</i> strain S5
PA181	<i>Pseudomonas aeruginosa</i> strain 181
PAO1	<i>Pseudomonas aeruginosa</i> PAO1
EC	<i>Escherichia coli</i>
CTAB	Cetyltrimethylammonium bromide
MB	Methylene Blue
SDS	Sodium dodecyl sulfate
MS	Mineral salt
E <sub>24</sub>	Emulsification Index
ml	mililiter
µl	microliter
ng/µl	nanogram per microliter
%	Percent
°C	Degree Celsius
h	Hour
min	Minute
PCR	Polymerase Chain Reaction
NCBI	National Center for Biotechnology Information
BLAST	Basic Local Alignment Search Tool

## LIST OF TABLES

Table	Caption	Page
1	Primers for the PCR amplification of <i>rhlA</i> and <i>rhlB</i>	14
2	PCR Reaction	15
3	CTAB plate result	20
4	The nucleotide BLAST result of <i>rhlA</i> gene	31
5	The nucleotide BLAST result of <i>rhlB</i> gene	32
6	The protein BLAST result of RhlA	32
7	The protein BLAST result of RhlB	33

## LIST OF FIGURES

Figures	Caption	Page
1	Chemical structure of rhamnolipids	4
2	Biosynthesis pathways of rhamnolipids	8
3	<i>Pseudomonas aeruginosa</i> rhamnolipid rhamnosyltransferase genes	9
4	Morphology of <i>Pseudomonas aeruginosa</i> strain S5 on plate	18
5	The Gram stain of <i>Pseudomonas aeruginosa</i> strains S5	18
6	<i>Pseudomonas aeruginosa</i> strains S5 grown on CTAB agar plate	20
7	Emulsification layer of the supernatant	22
8	Emulsification index value of different samples	22
9	Extraction of DNA	23
10	Agarose gel electrophoresis of <i>rhlA</i> gene	24
11	Agarose gel electrophoresis of <i>rhlB</i> gene	25
12	Multiple sequence alignment result for <i>rhlA</i> of <i>P. aeruginosa</i> S5 and <i>P. aeruginosa</i> PAO1	27
13	Multiple sequence alignment result for <i>rhlB</i> of <i>P. aeruginosa</i> S5 and <i>P. aeruginosa</i> PAO1	29
14	Multiple sequence alignment result for amino acid sequence of RhlA of <i>P. aeruginosa</i> S5 and <i>P. aeruginosa</i> PAO1	29
15	Multiple sequence alignment result for amino acid sequence of RhlB of <i>P. aeruginosa</i> S5 and <i>P. aeruginosa</i> PAO1	30
16	Phylogenetic tree of RhlA protein	34
17	Phylogenetic tree of RhlB protein	34

## LIST OF APPENDICES

<b>Appendix</b>	<b>Caption</b>	<b>Page</b>
1.0	Methods in preparing the chemicals	40
2.0	Methods in preparing the media	40
3.0	The nucleotide sequence of <i>rhlA</i>	42
4.0	The nucleotide sequence of <i>rhlB</i>	42
5.0	The amino acid sequence of RhlA	43
6.0	The amino acid sequence of RhlB	43
7.0	Multiple sequence alignment result when compared the RhlA protein of <i>P. aeruginosa</i> strain S5 with other <i>Pseudomonas aeruginosa</i> strain	43
8.0	Multiple sequence alignment result when compared the RhlB protein of <i>P. aeruginosa</i> strain S5 with other <i>Pseudomonas aeruginosa</i> strain	44

## CHAPTER 1

### INTRODUCTION

Rhamnolipids are surfactants that act as surface-active compounds capable of reduce interfacial tension and surface at the interfaces between liquids, solids and gases, thereby allowing them to mix or disperse readily as emulsions in water or other liquids (Banat et al., 2000). Rhamnolipids are important industrially because they are normally used as a detergent, emulsifier, foaming agent, wetting agents and also dispersant (Makkar et al., 2002).

The study about *Pseudomonas aeruginosa* strain S5 was initially performed by in the Enzyme Microbial Technology Laboratory (EMTech lab) for its ability to produce lipase (Baharum et al, 2003). It is an aerobic, gram-negative rod and motile bacterium. It was locally isolated from soil samples and known to be BTEX degrader (Baharum et al, 2003). Since rhamnolipids are normally produced by certain species of *Pseudomonas* (Pornsunthorntawee et al., 2010), the study of the potentiality of *P. aeruginosa* strain S5 for as a rhamnolipids producer is important. *Pseudomonas aeruginosa* produces primarily two forms of rhamnolipids: mono-rhamnolipids and di-rhamnolipids. Rhamnosyltransferase 1 catalyses the synthesis of mono-rhamnolipids from dTDP-L-rhamnose is encoded by the *rhlAB* operon and  $\beta$ -hydroxydecanoyl- $\beta$ -hydroxydecanoate, whereas the second rhamnosyltransferase, encoded by *rhlC* catalyses the di-rhamnolipid is produced from mono-rhamnolipid and dTDP-L-rhamnose (Deziel et al., 2003).

This study will elucidate the rhamnolipids biosynthesis pathway in *Pseudomonas aeruginosa* strain S5 and determine the availability of the *rhlA* and *rhlB* genes in this strain. The objectives of this study were (1) to determine the rhamnolipids production of *Pseudomonas aeruginosa* strain S5, (2) to isolate *rhlA* and *rhlB* from *Pseudomonas aeruginosa* strain S5 and (3) to analyze the *rhlA* and *rhlB* genes required for rhamnolipids biosynthesis.



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