UNIVERSITI TEKNOLOGI MARA

GENETIC IDENTIFICATION OF Amphibalanus amphitrite (Darwin, 1854) FROM BAGAN PASIR AND PASIR PENAMBANG, KUALA SELANGOR

NURUL NASUHA BINTI ROSEMAN

MSc

March 2019

UNIVERSITI TEKNOLOGI MARA

GENETIC IDENTIFICATION OF Amphibalanus amphitrite (Darwin, 1854) FROM BAGAN PASIR AND PASIR PENAMBANG, KUALA SELANGOR

NURUL NASUHA BINTI ROSEMAN

Dissertation submitted in partial fullfilment of the requirements for the degree of **Master of Science** (Applied Biology)

Faculty of Applied Science

March 2019

ABSTRACT

Barnacle is a marine sessile crustacean that inhabits the intertidal areas of tropical and temperate waters worldwide. Barnacles in family Balanidae belongs to superorder Thoracica and they consist of acorn barnacles and stalkless barnacles. In Malaysia, only a few studies have been done on barnacles in family Balanidae. In this present study, the molecular identification (16S rDNA) of barnacles in family Balanidae from Bagan Pasir and Pasir Penambang were investigated. Molecular identification is more accurate and faster compared to morphological classification as the morphology of the barnacles might differ due to geographical isolation. The objectives of this study were to assess the genetic information of barnacles in family Balanidae and to identify the specific species of barnacles in family Balanidae based on 16S rDNA identification. Four samples from Bagan Pasir and three samples from Pasir Penambang were studied to identify the specific species of the barnacles. The mitochondrial 16S molecular marker was used to identify the species and 500 bp band was observed on 1.5% agarose gel electrophoresis gel. The results showed that only Amphibalanus amphitrite (Darwin, 1854) was identified from both sampling sites. The phylogenetic tree constructed also stated that all the samples were identified as A.amphitrite (Darwin, 1854) and they were clustered in one clade. However, one sample from Bagan Pasir showed some difference in their DNA sequence as it was located at a different node from the other samples under the same cluster. In future, the molecular identification of the barnacles could have been done by using other molecular marker such as 12S rDNA, COI and the nuclear molecular marker 18S rDNA. The sampling site should also include other places in Selangor coast as to obtain different species of barnacles.

ACKNOWLEDGEMENT

Alhamdulillah. First and foremost my praise to Allah S.W.T for the strength and patience for me to complete my Masters study. My greatest gratitude and appreciation go to my supervisor Prof. Dr. Farida Zuraina bt Md Yusof for all the support and encouragement throughout this journey. I would also like to give my heartfelt thank you to her in helping and leading me in writing this thesis as well as conducting labwork. Without her patience and kindness, it would not be possible for me to complete my Masters study and thesis.

I would also like to extend my appreciation to my co-supervisor, Dr. Wan Razarinah bt Wan Abdul Razak for guiding me throughout the writing and completing my study. In addition, I would also like to say thank you to Puan Rafidah Rasol for her support and advice throughout this study.

Finally, I would like to thank my family who always encourage me to finish my Masters study and not to forget my friends who are always there to give their support.

TABLE OF CONTENTS

	Page		
CONFIRMATION BY PANEL OF EXAMINERS AUTHOR'S DECLARATION ABSTRACT	ii iii iv		
		ACKNOWLEDGEMENT	v
		TABLE OF CONTENTS	vi
LIST OF TABLES LIST OF FIGURES LIST OF SYMBOLS	viii ix xi		
		LIST OF ABBREVIATIONS	xii
		CHAPTER ONE: INTRODUCTION	1
1.1 Background of Study	1		
1.2 Problem Statement	2		
1.3 Significance of Study	2		
1.4 Objectives of the Study	2		
1.5 Scope and Limitations of Study	3		
CHAPTER TWO: LITERATURE REVIEW	4		
2.1 Family Balanidae	4		
2.2 Barnacle Identification	7		
2.3 Molecular Markers	9		
2.4 Economic Value of Barnacles	12		
CHAPTER THREE: RESEARCH METHODOLOGY	14		
3.1 Sampling Materials	14		
3.2 DNA-based Protocols	15		
3.2.1 DNA extraction	15		
3.2.2 DNA quantification and qualification	16		

3.3 Polymerase Chain Reaction

16