

DNA FINGERPRINTING OF LOKAN, *Polymesoda eximia*
USING RAPD - PCR TECHNIQUE

LEE SAU YEE

DEPARTMENT OF BIOLOGICAL SCIENCE
FACULTY OF SCIENCE AND TECHNOLOGY
KOLEJ UNIVERSITI SAINS DAN TEKNOLOGI MALAYSIA
KUSTEM
2003

1100024989

Universiti Sains Dan Teknologi Malaysia

LP 9 FST 1 2003



1100024989

DNA fingerprinting of Lokan, Polymesoda expansa using
RAPD-PCR technique / Lee Sau Yee.

1100025055

PERPUSTAKAAN
KOLEJ UNIVERSITI SAINS & TEKNOLOGI MALAYSIA
(KUSTEM)

c/n 1645

Pengarang	LEE SAU HOOI	No. Panggilan	LP 8
Judul	NERUS RIVER WATER - -		PST 13
Tarikh	Waktu Pemulangan	Nombor Ahli	Panda tangan
2003/05/10	6/15PM	UK6709	Sau

29/2/10

-P
9
PST
1
2003

DNA Fingerprinting of Lokan, *Polymesoda expansa*
Using RAPD-PCR Technique

By:

Lee Sau Yee

This project report is submitted in partial Fulfillment of the requirement for the
Bachelor of Applied Science
(Conservation and Management of Biodiversity)

PUSAT PERLAUTAN DAN SULTANAH NUR ZAHIRAH

Department of Biological Sciences
Faculty of Science and Technology

Kolej Universiti Sains Dan Teknologi Malaysia, KUSTEM

2003

1100024989

This project report should be cited as:

Lee, S.Y. 2003. DNA Fingerprinting of Lokan, *Polymesoda expansa* Using RAPD-PCR Technique. Report of Final Year Academic Project, Bachelor of Applied Science-Conservation and Management of Biodiversity, Faculty of Science and Technology, Kolej Universiti Sains Dan Teknologi Malaysia. 48p.

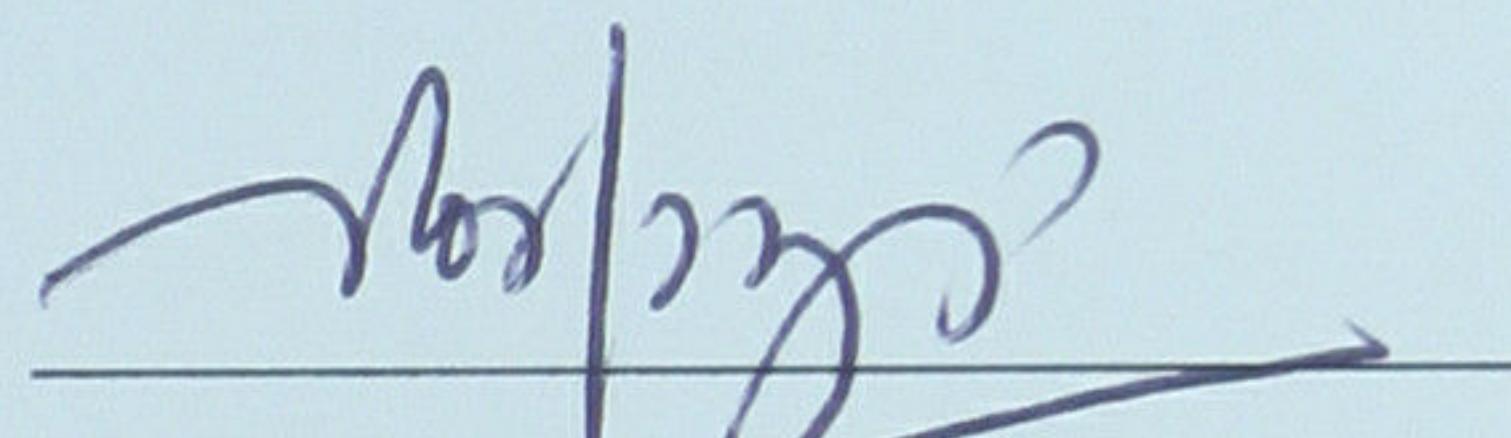
No part of this project report may be reproduced by any mechanical, photographic, or electronic process, or in the form of photographic recording, nor mat it be stored in a retrieval system, transmitted, or otherwise copied for public or private use, without written permission from the author and the supervisor of the project.

KOLEJ UNIVERSITI SAINS DAN TEKNOLOGI MALAYSIA

PENGAKUAN DAN PENGESAHAN LAPORAN PENYELIDIKAN ILMIAH TAHUN AKHIR

Adalah ini diakui dan disahkan bahawa laporan penyelidikan ilmiah tahun akhir bertajuk DNA Fingerprinting of Lokan, *Polymesoda expansa* Using RAPD-PCR Technique oleh Lee Sau Yee, no. matrik UK 4090 telah diperiksa dan semua pembetulan yang disarankan telah dilakukan. Laporan ini dikemukakan kepada Jabatan Sains Biologi sebagai memenuhi sebahagian daripada keperluan memperolehi ijazah Sarjana Muda Sains – Pengurusan dan Pemuliharaan Biodiversiti, Fakulti Sains dan Teknologi, Kolej Universiti Sains dan Teknologi Malaysia.

Disahkan oleh:



(CIK NORAZNAWATI ISMAIL)

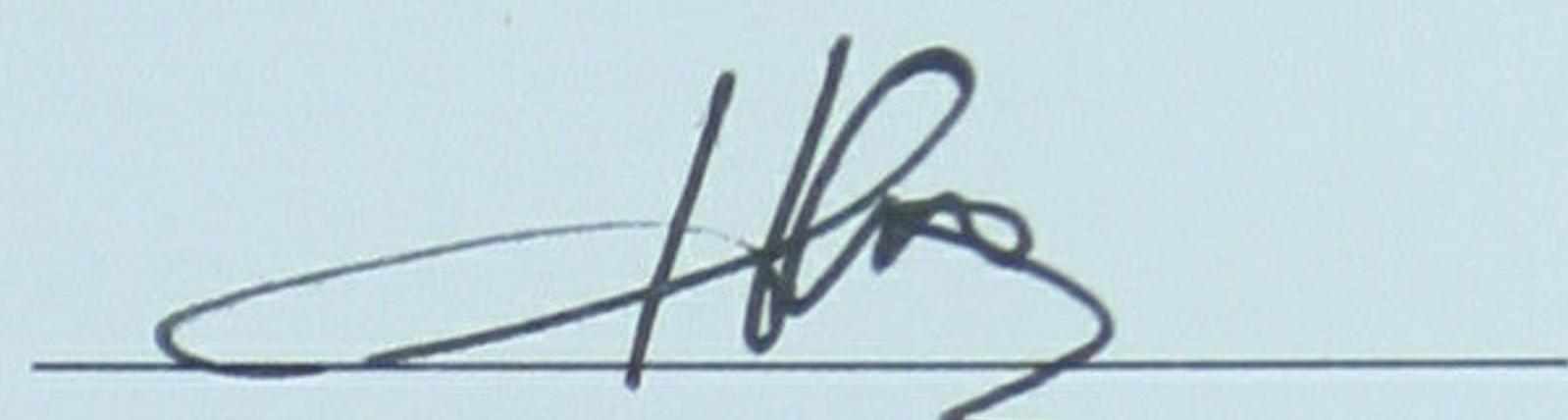
Penyelia Utama

Nama: Cik Noraznawati Binti Ismail

Cop:

Noraznawati Ismail
Lecturer
Department of Biological Science
Faculty of Science and Technology
KUSTEM 21300 K. Terengganu

Tarikh: 27/02/03



(PROF. DR. CHAN ENG HENG)

Ketua Jabatan Biologi

PROF. DR. CHAN ENG HENG

Cop:

Ketua
Jabatan Sains Biologi
Fakulti Sains dan Teknologi
Kolej Universiti Sains dan Teknologi Malaysia
(KUSTEM)
21030 Kuala Terengganu.

Tarikh:

3/3/03

ACKNOWLEDGEMENT

Here I would like to take this opportunity to express my appreciation to all the people who have helped me finish this project report. First of all, I would like to thank my supervisor, Miss Noraznawati Binti Ismail who willfully taught me the correct way to run the laboratory work and helped in the writing of this report. The second person I would like to thank is Miss Wan Bayani who helped to guide and teach me whenever my supervisor was out for a meeting or on leave. My thanks also goes to the laboratory assistants and research assistants, Pn. Faridah, Kak Ella, Kak Ina, Mr. Sayed, Mr. Rizal and others that provided information about the laboratory, instructions on using the equipments and also their cooperation for letting me use the laboratory on weekends and holidays.

I am also grateful to my family for encouraging me and giving me the permission to stay in Terengganu to do the laboratory work during the holidays when I'm supposed to accompany them at home. Talking about encouragement and support, I would like to thank Eng Oon who cheers me up whenever the laboratory result is not that good and also to my groupmates, Shen, Sik Loo and Mei Yen who tolerated and cooperated with each other. Last but not least thank you to my friends and roommate who have kept on supporting me. I appreciate your friendship and hope to keep in touch in the future. May all of you find the light in the darkness and guide you all to success in life.

ABSTRACT

Lokan, *Polymesoda expansa* (*P. expansa*) formerly known as *Geloina coaxans* is a filter feeder and can be found in Southern Asia and Malaysia. Besides playing an important role as a bioindicator, it is also a food resource for the east coast of Peninsular Malaysia. Random Amplified Polymorphic DNA (RAPD) based on Polymerase Chain Reaction (RAPD-PCR) technique was used to amplify the genomic DNA of *P. expansa*. The aims of this study were to establish the DNA fingerprinting of *P. expansa*, to verify the genetic variability and to analyze the degree of polymorphisms on DNA level marker of *P. expansa*. The genomic DNA of 12 samples (six samples per population) was extracted from muscular tissue using Phenol-Chloroform Protocol. The purity of DNA ranged from 1.453-1.731, estimated from the ratio between the reading of absorbance at 260 nm and 280 nm (OD_{260}/OD_{280}) in UV-spectrophotometer. The purity of DNA can be observed through the single band formed on a 0.8% agarose gel stained with 0.5 μ g/mL ethidium bromide. DNA quantifications ranged between 235.00-580.00 ng/ μ L. 40-50% of 20 primers with 60-70% GC content were able to amplify the fragments from each genomic DNA. The fragments of PCR products were observed when electrophoresised in 2.0% (w/v) agarose gel stained with 0.5 μ g/mL ethidium bromide. The three primers selected (OPA 08, 09 and 12) generated a total of 51-55 scorable loci (bands) with 90.91-93.98% polymorphic loci. RAPD banding for Kg. Pulau Tok Aji and Kg. Laut populations ranged from 3-12 bands and 3-9 bands with 250-2200 bp and 240-2080 bp. The similarity index among individuals between populations was 0.263 ± 0.152 while the genetic distance value was 0.737 ± 0.152 . From this study, a monomorphic band (650 bp in OPA 09) was found to be reliable as a diagnostic marker of *P. expansa*. Further studies on this species should be carried out in order to conserve and maintain the gene pool of *P. expansa* which have the commercial potential in aquaculture.

ABSTRAK

Lokan, *Polymesoda expansa* (*P. expansa*) pada mulanya dikenali sebagai *Geloina coaxans* merupakan organisma jenis pemakan hasil tapisan dan ditemui di benua Asia Selatan dan Malaysia. Selain daripada sebagai organisma penunjuk bio, ia juga merupakan sumber makanan bagi penduduk di pantai timur Semenanjung Malaysia. Teknik “Random Amplified Polymorphic DNA” (RAPD) yang berasaskan “Polymerase Chain Reaction” (RAPD-PCR) digunakan untuk mengamplifikasi genomik DNA *P. expansa*. Tujuan kajian ini dijalankan adalah untuk menerbitkan satu pencapjarian DNA untuk *P. expansa*, untuk mengenalpasti perubahan genetik dan untuk menganalisis tahap polimorfik berdasarkan penanda DNA *P. expansa*. Genomik DNA untuk 12 sampel (Enam sampel untuk satu populasi) diekstrak daripada tisu otot dengan menggunakan kaedah Fenol-Kloroform. Julat ketulenan DNA yang diperolehi daripada nisbah bacaan penyerapan pada 260 nm dan 280 nm (OD_{260}/OD_{280}) pada spektrofotometer ialah 1.453-1.731. Ketulenan DNA yang diperolehi juga boleh diperhatikan dengan pembentukan satu jalur tunggal pada 0.8% gel agaros yang telah diwarnai dengan 0.5 μ g/mL etidium bromida. Julat kepekatan DNA ialah antara 235.00-580.00 ng/ μ L. Daripada 20 primer yang mengandungi 60-70% GC, hanya 40-50% dapat mengamplifikasi fragmen daripada setiap genomik DNA. Hasil fragmen daripada PCR dapat diperhatikan apabila di elektroforesiskan dalam 2.0% gel agaros yang diwarnai dengan 0.5 μ g/mL etidium bromida. Sejumlah 51-55 jalur dicatatkan dengan 90.91-93.98% jalur polimorfik serta julat penjaluran RAPD antara 3-12 jalur dan 3-9 jalur dengan 250-2200 bp dan 240-2080 bp bagi populasi Kg. Pulau Tok Aji dan Kg. Laut diperolehi daripada tiga primer (OPA08, 09 dan 12) yang dipilih. Indek kesamarataan sesama individu antara populasi adalah 0.263 ± 0.152 dengan jarak genetik antara 0.737 ± 0.152 . Satu jalur monomorfik iaitu 650 bp dalam OPA09 yang diperolehi daripada kajian ini dapat digunakan sebagai penanda diagnostik untuk *P. expansa*. Kajian yang lebih mendalam harus dijalankan untuk memelihara dan mengekalkan kolam genetik *P. expansa* yang mempunyai potensi untuk dikultur dalam bidang akuakultur.