



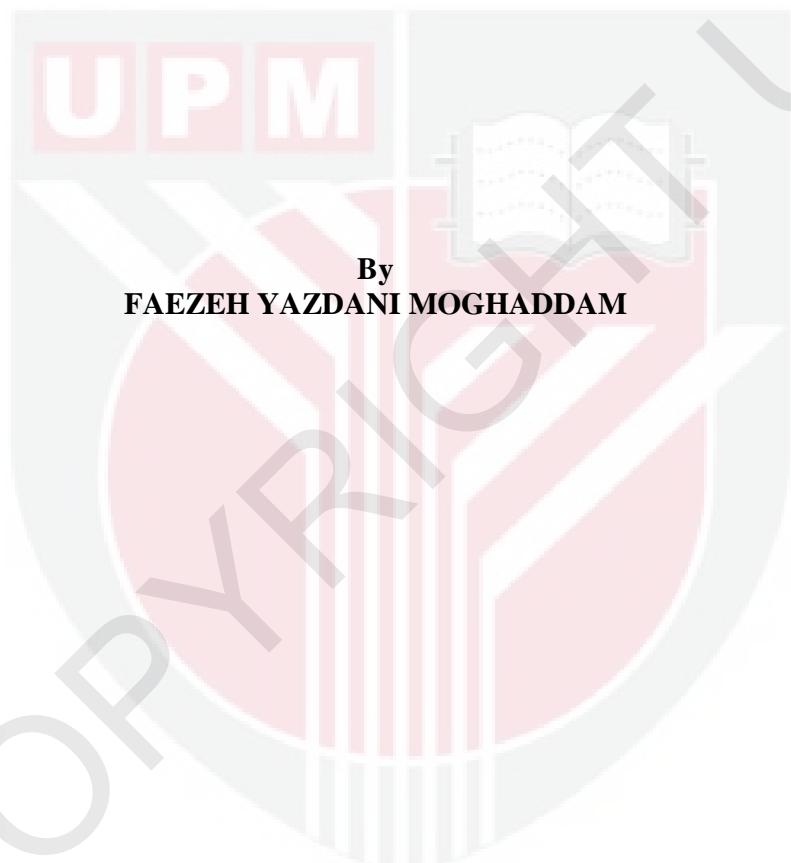
**UNIVERSITI PUTRA MALAYSIA**

**PHYLOGENETIC RELATIONSHIPS AMONG MALAYSIAN  
*PUNTIUS* AND ITS ALLIES (PISCES: CYPRINIDAE)**

**FAEZEH YAZDANI MOGHADDAM**

**FS 2012 4**

**PHYLOGENETIC RELATIONSHIPS AMONG MALAYSIAN *PUNTIUS* AND ITS  
ALLIES (PISCES: CYPRINIDAE)**



**Thesis Submitted to the School of Graduate Studies, Universiti Putra Malaysia, in Fulfilment  
of the Requirements for the Degree of Doctor of Philosophy  
January 2012**

Abstract of thesis presented to the Senate of Universiti Putra Malaysia  
in fulfilment of the requirement for the degree of Doctor of Philosophy

**PHYLOGENETIC RELATIONSHIPS AMONG MALAYSIAN *PUNTIUS* AND  
ITS ALLIES (PISCES: CYPRINIDAE)**

By

**FAEZEH YAZDANI MOGHADDAM**

**January 2012**

**Chairman: Associate Professor Siti Khalijah Daud, PhD**

**Faculty: Science**

The variation of criteria among authors and the discrepancies between morphological and molecular data generated a troublesome situation concerning the delimitation of the genus *Puntius*, which remains as yet unresolved. This is the first documentation on phylogenetic relationships among *Puntius* species in Malaysia integrating classical morphometrics, geometric morphometrics and molecular techniques. Morphological and phylogenetic studies were conducted on 10 Malaysian *Puntius* species and their allies, namely *P. bulu*, *P. fasciatus*, *P. schwanenfeldii*, *P. binotatus*, *P. everetti*, *P. tetrazona partipentazona*, *P. tetrazona hexazona*, *P. daruphani*, *P. lateristeriga* and *P. gonionotus* obtained from five locations in Peninsular Malaysia.

A total of 312 samples were analysed. The quantitative data were analysed using analysis of variance (ANOVA), Principal Component Analysis (PCA) and Discrimination Function Analysis (DFA). Discriminating efficiency was highest in the

truss network morphometric followed by classical morphometric and meristic techniques. In geometric morphometric Canonical Variate Analysis was performed on the total shape matrix. Results were further supported by sequencing of the cytochrome c oxidase subunit I (Cox1, 501 bp) and cytochrome b (Cytb, 857 bp) mitochondrial genes, and recombination activating gene (Rag2, 860 bp ) and beta actin ( $\beta$ -actin, 911) nuclear genes. A combination of mitochondrial and nuclear data for phylogenetic relationships analysis was conducted from 39 representative samples of *Puntius* species. The phylogeny among taxa was constructed through Neighbor joining (NJ), Maximum parsimony (MP), Maximum likelihood methods (ML) and Bayesian inference (BI). The result of genetic distances and haplotype diversities among species were high indicating a complex genus of *Puntius*. In an attempt to confirm the phylogenetic position of *Puntius* within common cyprinids in Southeast Asia, dataset of *Puntius* in Peninsular Malaysia was complemented by Cytb and Rag2 sequences for different species of *Puntius* and common cyprinids belonging to Southeast Asia in GenBank. The tree with Maximum likelihood scores was chosen as the best tree. Clades retrieved from different trees strongly supported that subfamily Cyprininae and four selected genera within *Puntius* complex were monophyletic group, while tribe Barbinini, Cyprinini and genus *Puntius* were not monophyletic. Results of the combined data set were similar with the taxonomic hierarchy based on geometrical morphometric analyses. Divergence time estimated using a combination of molecular data from multiple DNA loci and fossil evidence for *Puntius* among Cyprininae was about 26.85 million years ago (26.08-28.22) and supported their early Oligocene origin.

Abstrak tesis yang dikemukakan kepada Senat Universiti Putra Malaysia sebagai  
memenuhi keperluan ijazah Doktor Falsafah

**HUBUNGAN FILOGENETIK ANTARA *PUNTIUS* DI MALAYSIA DAN  
YANG BERKAITAN (PISCES: CYPRINIDAE)**

Oleh

**FAEZEH YAZDANI MOGHADDAM**  
**Januari 2012**

**Pengerusi: Profesor Maddy Siti Khalijah Daud, PhD**

**Fakulti: Sains**

Variasi kriteria di kalangan penulis dan percanggahan antara data morfologi dan molekul menghasilkan situasi yang menyukarkan persempadanan genus *Puntius* dan masih lagi belum dapat diselesaikan sehingga kini. Kajian ini merupakan dokumentasi pertama bagi perhubungan filogenetik di kalangan sepuluh spesies *Puntius* yang pertama di Malaysia yang mengintergrasikan teknik morfometrik konvensional, morfometrik geometrik and teknik molekular. Kajian morfologi dan filogenetik telah dijalankan ke atas 10 spesies *Puntius* Malaysia dan yang berkaitan, iaitu *P. bulu*, *P. fasciatus*, *P. schwanenfeldii*, *P. binotatus*, *P. everetti*, *P. tetrazona partipentazona*, *P. tetrazona hexazona*, *P. daruphani*, *P. lateristeriga* dan *P. gonionotus* yang di perolehi dari lima lokasi di Semenanjung Malaysia.

Sejumlah 312 sampel telah dianalisis. Kuantitatif data dianalisis menggunakan analisis varians (ANOVA), Analisis Komponen Utama (PCA) dan Analisis Diskriminasi (DFA).

Kecekapan diskriminasi adalah paling tinggi pada teknik morfometrik Truss. diikuti dengan morfometrik biasa dan meristik. Bagi morfometrik geometri, analisis varians kanonikal di jalankan ke atas matriks bentuk total. Keputusan telah di sokong oleh jujukan gen mitokondrial DNA pada sitokrom c oksida subunit I (CoxI, 501bp) dan sitokrom b (Cyt b, 857 bp), serta gen nukleus iaitu gen rekombinasi teraktif 2 (Rag2, 860bp) dan beta aktin ( $\beta$ -actin, 911 bp). Kombinasi data mitokondria dan nukleas dilakukan ke atas 40 sampel spesies *Puntius*.

Filogeni antara taksa dibina dengan menggunakan ‘Neighbour Joining’ (NJ), dan ‘parsimoni maksimum’ (MP), kaedah ‘Maximum likelihood’ (ML), dan inferensi Bayesian (BI). Jarak genetik dan diversiti haplotip antara spesies adalah tinggi, menunjukkan *Puntius* adalah genus yang kompleks. Cubaan untuk mengesahkan posisi filogenetik *Puntius* dalam siprinid di Asia Tenggara, data *Puntius* dari Semenanjung Malaysia telah dilengkapkan dengan jujukan Cytb dan Rag2 bagi spesis *Puntius* yang berlainan dan siprinid bias ayang berasal dari Asia Tenggara yang terdapat di Genbank. Pohon dengan skor Parsimoni maksimum dipilih sebagai pohon terbaik. Klad yang diperolehi daripada pohon yang berbeza menyokong kuat kewujudan subfamili Cyprininae dan empat genus dalam kompleks *Puntius* yang terpilih adalah kumpulan monofiletik., manakala suku Barbinini, Cyprinini dan genus *Puntius* adalah bukan monofiletik. Keputusan daripada kombinasi set data menunjukkan persamaan dengan hirarki taksonomi berasaskan analisis morfometri geometrik. Masa pencapahan yang dianggarkan dengan menggunakan gabungan data molekul daripada lokus DNA berganda dan bukti fosil DNA *Puntius* di antara Cyprininae adalah lebih kurang 26.85 juta tahun yang lalu (26.08-28.22) dan ini menyokong ia berasal daripada zaman Oligocene awal.

## **ACKNOWLEDGEMENT**

First and foremost, I would like to express my utmost gratitude to my highly respected supervisor, Associate Professor Dr Siti Khalijah Daud. Thanks for your fantastic open door policy, faith, positivity, interest, tolerance, encouragement and patience.

I would like to express my deepest thanks to Dr Mansour Aliabadian my Supervisory committee for his advice, invaluable guidance, hospitality, support and encouragement throughout the period of the study.

I would like to acknowledge other members of my Supervisory committee, Prof. Dr Siti Shapor Siraj and Prof Dr Jothi Malar Panandam for their guidance, persistence encouragement, and associated aid throughout this study.

Thank you to all my international collaborators, who not only assisted with collection but also made me welcome in their home country and revealed to me a side of South east Asia that I could never have experienced on my own.

I extend my thanks to, Dr. Eleanor A S Adamson, Dr. Mahvash Seifali, Dr. Arash Javanmard and Nadiatul Hafiza, who assisted me.

Lastly, to my husband Siamak: without your love and support this project would have been a very different and much harder road to travel. Thanks for enduring the long hours I have spent away from home in the field, at conferences, in the lab, and in front of my computer, for tolerating my total absorption, procrastination, lack of income and general lack of sanity.

I certify that a Thesis Examination Committee has met on 20/Jan/2011 to conduct the final examination of FAEZEH YAZDANI MOGHADDAM on her Doctor of Philosophy thesis entitled "The Systematics of Malaysian *Puntius* and Its Allies (Pisces: Cyprinidae)" in accordance with the Universities Pertanian Malaysia (Higher Degree) Act 1980 and Universiti Pertanian Malaysia (Higher Degree) Regulations 1981. The committee recommends that the student be awarded the Doctor of Philosophy.

Members of the Thesis Examination Committee were as follows:

Chairman, PhD

Faculty of Science.  
Universiti Putra Malaysia  
(Dr. Hishamuddin b Omar)

Examiner 1, PhD

Faculty.of of Science  
Universiti Putra Malaysia  
(Prof Madya Dr. Faridah binti Qamaruz Zaman)

Examiner 2, PhD

Faculty of Agriculture  
Universiti Putra Malaysia  
(Prof Madya Dr. Maheran bt Abd Aziz)

External Examiner 2, PhD  
Faculty of Science  
Aburn University United stated

---

(Assoc. Prof. Dr Jonatan W. Armbruster)  
**NORITAH OMAR, PhD**  
Associate Professor and Deputy Dean  
School of Graduate Studies

Universiti Putra Malaysia

This thesis was submitted to the Senate of Universiti Putra Malaysia and has been accepted as fulfilment of the requirement for the degree of Doctor of Doctor of Philosophy.

The members of the Supervisory Committee were as follows:

**Siti Khalijah Daud, PhD**

Associate Professor

Faculty of Science

Universiti Putra Malaysia

(Chairman)

**Siti Shapor Siraj PhD**

Professor

Faculty of Agriculture

Universiti Putra Malaysia

(Member)

**Jothi Malar Panandam PhD**

Professor

Faculty of Agriculture

University Putra Malaysia

(Member)

**Mansour Aliabadian PhD**

Associate Professor

Faculty of science

Ferdowsi university of Mashhad

(Member)

---

**BUJANG BIN KIM HUAT, PhD**

Professor and Dean

School of Graduate Studies

Universiti Putra Malaysia

## **DECLARATION**

I declare that the thesis is my original work except for quotations and citations which have been duly acknowledged. I also declare that it has not been previously and is not concurrently submitted for any other degree at Universiti Putra Malaysia or other institutions.

**FAEZEH YAZDANI MOGHADDAM**

Date: 20 January 2012



## TABLE OF CONTENTS

	<b>Page</b>
<b>DEDICATION</b>	ii
<b>ABSTRACT</b>	III
<b>ABSTRAK</b>	v
<b>ACKNOWLEDGEMENTS</b>	VIII
<b>APPROVAL</b>	X
<b>DECLARATION</b>	XI
<b>LIST OF TABLES</b>	XVII
<b>LIST OF FIGURES</b>	XIX
<b>LIST OF ABBRIVIATIONS</b>	xxiii
 <b>CHAPTER</b>	
<b>1 GENERAL INTRODUCTION</b>	1
1.1 Objectives of the Study	5
<b>2 LITERATURE REVIEW</b>	6
2.1 Cypriniformes	6
2.2 Family Cyprinidae	6
2.3 Sub family Cyprininae	7
2.4 The Biology of Malaysian <i>Puntius</i>	8
2.5 Taxonomic Classification	8
2.5.1 <i>Puntius binotatus</i> (Valenciennes, 1842)	9
2.5.2 <i>Puntius everelli</i> (Boulenger 1894)	10
2.5.3 <i>Puntiusjasciatus</i> (Jerdon, 1849)	11
2.5.4 <i>Puntius lateristriga</i> (Valenciennes 1842)	12
2.5.5 <i>Puntius tetrazona partipentazona</i> (Fowler, 1934)	13
2.5.6 <i>Puntius tetrazona hexazona</i> (Weber and de Beaufort, 1912)	14
2.5.7 <i>Puntius .s·chwanel?·eldii</i> (Bleeker, 1853)	15
2.5.8 <i>Puntius gonionotus</i> (Bleeker 1850)	16
2.5.9 <i>Puntius bulu</i> (Bleeker, 1851)	17
2.5.10 <i>Puntius daruphani</i> (1934)	18
2.6 The Taxonomic Problem of <i>Puntius</i>	19
2.7 Morphological Study	25
2.8 DNA Marker	27
2.9 Molecular Phylogenetic	27
2.10 Inferring Phylogenetic Trees	30
2.10.1 Neighbor Joining Method (NJ)	31
2.10.2 Character-Based Methods (Optimality)	32
2.10.3 Maximum Parsimony (MP)	32
2.10.4 Maximum Likelihood (ML)	33

2.10.5 Bayesian inference (Bl)	34
2.11 Mitochondrial DNA (mtDNA) Markers	34
2.12 Nuclcar Genes	35
<b>3 MORPHOLOGICAL VARIATION AMONG TEN <i>Puntius</i> SPECIES</b>	<b>37</b>
3.1 Introduction	37
3.2 Material and Methods	39
3.2.1 Sampling	39
3.2.2 Conventional Morphometric (CM)	40
3.2.3 Meristic Characters (M)	42
3.2.4 Traditional Truss-Based Morphometrics (TM)	43
3.2.5 Geometric Morphometric	45
3.3 Results	46
3.3.1 Morphometric Data Conventional Morphometric Analysis (CM)	46
3.3.2 Meristic Data Analysis (M)	53
3.3.3 Truss Morphometric TM	57
3.3.4 Geometric Morphometric	61
3.4 Discussion	65
<b>4 <i>Puntius</i> GENETIC VARIATION AND PHYLOGENETIC ANALYSES BASED ON MITOCHONDRIAL DNA</b>	<b>69</b>
4.1 Introduction	69
4.2 Materials and methods	71
4.2.1 Sample Description and Location	71
4.2.2 DNA Extraction and Isolation	74
4.2.3 Measurement of DNA Quality	75
4.2.4 Gene Screening and Amplification	76
4.2.5 DNA Purification and Sequencing	77
4.2.6 Sequenee Alignment and Phylogenetic Analysis	78
4.2.7 Data analysis for barcoding gap	79
4.3 Results	79
4.3. I Sequences Variation of Cytb Mitochondrial Genes	79
4.3.2 Genetic Distance and Barcoding gap for Cytb	88
4.3.3 Sequences Variation of Cox I Mitochondrial Genes	90
4.3.4 Phylogenetic Relationships among Sample	92
4.3.5 Genetic Distance and Barcoding gap for Cox I	97
4.4 Discussion	99
<b>5 <i>Puntius</i> GENETIC VARIATION AND PHYLOGENETIC ANALYSES BASED ON NUCLEAR DNA</b>	<b>102</b>
5.1 Introduction	102
5.2 Materials and Methods	104
5.2.1 Sample Description and Location	104
5.2.2 DNA Extraction and Isolation, Measurement ofDNA Quality, Gene Screening	107
5.2.3 DNA Purification, Sequencing and Phylogenetic Analysis	109

<b>9 CONCLUSIONS AND RECOMMENDATIONS</b>	179
9.1 Conclusions	179
9.2 Future Recommendations	180
<b>REFERENCES</b>	182
<b>APPENDICES</b>	203
<b>BIODATA OF STUDENT</b>	209

