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DNA characterization of normal and damaged gills following copper exposure / Che Norihan Che Wahab.



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DNA CHARACTERIZATION OF NORMAL AND DAMAGED GILLS FOLLOWING
COPPER EXPOSURE

By
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Research Report submitted in partial fulfillment of the requirements for the degree of
Bachelor of Science (Marine Science)

**DEPARTMENT OF MARINE SCIENCE
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**JABATAN SAINS SAMUDERA
FAKULTI SAINS DAN TEKNOLOGI
KOLEJ UNIVERSITI SAINS DAN TEKNOLOGI
MALAYSIA**

**PENGAKUAN DAN PENGESAHAN LAPORAN
PROJEK PENYELIDIKAN I DAN II**

Adalah ini diakui dan disahkan bahawa laporan penyelidikan bertajuk :

DNA characterization of normal and damaged gills following copper exposure oleh **Che Norihan bt Che Wahab, No. Matrik: UK 8918** telah diperiksa dan semua pembetulan yang disarankan telah dilakukan. Laporan ini dikemukakan kepada Jabatan Sains Samudera sebagai memenuhi sebahagian daripada keperluan memperolehi Ijazah Sarjana Muda Sains – Sains Samudera, Fakulti Sains dan Teknologi, Kolej Universiti Sains dan Teknologi Malaysia.

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TABLE OF CONTENTS

ACKNOWLEDGEMENTS	ii
LIST OF TABLES	viii
LIST OF FIGURES	x
LIST OF APPENDIXS	xii
LIST OF SYMBOLS	xi
ABSTRACT	xiv
ABSTRAK	xv
CHAPTER I	
Introduction	1
CHAPTER II	
Literature review	4
2.1. Red/Nile tilapia Fish (<i>Oreochromis niloticus</i>)	4
2.2. Copper in fish	5
2.3. 96-hr LC ₅₀ acute toxicity test	7
2.4. Heavy metal in tissue gills	9
2.5. Fish as a bio-indicator	10
2.6. Heavy metal –copper	11
2.7. Fish gill	12

CHAPTER III

Material and methodology

3.1. Study site	19
3.1.1. Experimental design	19
3.2. Part A: Determination of 96 hours LC50 of Cu ²⁺ on Tilapia fingerlings	21
3.2.1. Statistical analysis	22
3.2.2. Uptake and accumulation of copper in tilapia fish	22
3.3. Part B: Determination of metal concentration in tissue of Tilapia fingerlings at 7, 14 and 21 days exposed to copper	24
3.3.1. Tissue sampling	24
3.3.2. Tissue preparation for determination of metal Concentration the gills	24
3.3.3. Open acid digestion	25
3.3.4. Metal recovery measurement	25
3.3.5. Heavy metal measurement in the tissue gills of tilapia Fingerlings	26
3.4. Part C: DNA analysis of normal and exposure tilapia	
Fingerlings at 7, 14 and 21 days exposure to copper	27
3.4.1. DNA extraction	27
3.4.2. Purification and quantification of DNA	28

3.4.2.1. Biophotometer	28
3.4.2.2. Electrophoresis method	29
3.4.3. DNA amplification by RAPD	29
3.4.4. Database establishment	31
3.4.5. Data analysis	32

CHAPTER IV

Results

4.1. Part A: Determination of 96 hours LC ₅₀ Cu ²⁺ for red nile/tilapia fish	33
4.2. Part B: Uptake and accumulation of copper on tilapia Fingerlings's gill for the period of 21 days	35
4.2.1. Metal recovery measurement	35
4.2.2. Accumulation of copper on tilapia	36
4.3. Part C: Effect of copper on the genomic of the red tilapia at 7, 14 and 21 days of exposure.	39
4.3.1. DNA extraction	39
4.3.2. RAPD screening test	41
4.3.3. PCR data analysis	42

CHAPTER V

Discussion	46
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CHAPTER VI

Conclusion	53
References	54
Appendix	60
Curriculum Vitae	71

LIST OF TABLES

2.8.1a.	Universal Primers	18
3.2.2a.	Concentration design for uptake and accumulation of Cu ²⁺ in tilapia fingerlings for 21 days period	23
3.3.4a.	Summary of DOLT-3, certified Metal concentration	26
3.4.3a.	Primer sequences selected	28
3.4.3b.	Final concentration of RAPD mixture	30
3.4.3c.	Amplification profile	31
4.1.1.	96-h LC50 using spearman-karber computer programme	33
4.1.2.	Comparison 96HLC50 of copper on tilapia by using different types of method	34
4.1.3.	96-h median lethal concentration at different percentage	35
4.2.2a.	The statistical correlation and regression for the metals accumulated in nile tilapia within 21 days	37
4.3.1a.	Absorbance reading by biophotometer at 260 and 280 wavelength and concentration of DNA	39

4.3.3a. RAPD fingerprinting pattern of tilapia fish of 7 days exposed to copper	42
4.3.3b. RAPD fingerprinting pattern of tilapia fish of 14 days exposed to copper	44
4.3.3c. RAPD fingerprinting pattern of tilapia fish of 21 days exposed to copper	45

LIST OF FIGURE

2.1.1.	Red Nile tilapia	5
2.7.1.	Diagram of gill filament Vasculature	13
2.8.2.	Illustration on PCR-RAPD technique work	17
4.2.2a.	Copper accumulation in gills tissue at 7, 14 and 21 days of exposure	36
4.2.2b.	Metal Concentration according to size for 7, 14 and 21 days Exposure.	38
4.3.1a.	The electrophoresis pattern of gills tilapia fish genomic DNA	40
4.3.2a.	RAPD screening test for selective primers	41
4.3.3a.	RAPD fingerprinting profiles of tilapia fish at 7 days exposed to copper	42
4.3.3b.	RAPD fingerprinting profiles of tilapia fish at 14 days exposed to copper	43
4.3.3c.	RAPD fingerprinting profiles of tilapia fish at 21 days exposed to copper	44

LIST OF SYMBOLS

ICP-MS	Inductively Coupled Plasma – Mass Spectrophotometry
cm	Centimeter
g	Gram
mg	milligram
μg	microgram
Cu	Copper
Cu^{2+}	Copper in ionic form
HNO_3	Nitric Acid
H_2SO_4	Sulfuric Acid
HCL	Hydrochloric Acid
H_2O_2	Hydrogen Peroxide
ppm	parts per million or equivalent to mgL^{-1}
mgL^{-1}	milligram per liter
L	Liter
d.w.	Dry weight
PCR	Polymerase Chain Reaction
PCR-RAPD	Randomly Amplified Polymerase Chain Reaction

LIST OF APPENDIXES

I	Standard stock solution of copper (Cu^{2+})	60
II	Toxicity testing-percentage of mortality of nile tilapia (<i>Oreochromis niloticus</i>) at different time exposed to different metal concentration (Copper replicate 1)	61
III	Toxicity testing-percentage of mortality of nile tilapia (<i>Oreochromis niloticus</i>) at different time exposed to different metal concentration (Copper replicate 2)	62
IV	Toxicity testing-percentage of mortality of nile tilapia (<i>Oreochromis niloticus</i>) at different time exposed to different metal concentration (Copper replicate 3)	63
V	Toxicity Testing-Summary on calculating the LC 50 values (Probit and Spearman karber method) using Spearman-karber Computer programme (Replicate 1)	64
VI	Toxicity Testing-Summary on calculating the LC 50 values (Probit and Spearman karber method) using Spearman-karber Computer programme (Replicate 2)	65
VII	Toxicity Testing-Summary on calculating the LC 50 values (Probit and Spearman karber method) using Spearman-karber Computer programme (Replicate 3)	66

VIII	Volume of copper at each concentration for the uptake copper for 21 days.	67
IX	Total Length, Body Length, Total Body Weight, Wet Weight and Dry Weight of <i>Oreochromis niloticus</i> at 7, 10, 21 Days Exposure.	68
X	Concentration of Metal in the Gills Tissue (ppb), Standard Length and Total Body Weight (g) of Nile Tilapia Fingerlings at 7, 14 and 21 Days Exposure	69
XI	Statistical Analysis (ANOVA 2 ways without replication): Mean Accumulation of copper in gill tissues of nile tilapia fingerlings for 30 days.	70

ABSTRACT

The ultimate aim of this study is to predict how genomic of aquatic organisms respond to heavy metal exposure. Red tilapia fingerlings siblings (*Oreochromis niloticus*) (0.5 cm-5.0 cm) were exposed to copper (Cu^{2+}). The study was conducted in three parts whereby each part is interrelated. The first part was the copper acute toxicity test. Using probit method, the 96-h LC50 value for tilapia fingerlings was 1.82 ppm. This value was then used to design a sub-lethal concentration (1.82 ppm, 1.36 ppm, 0.91 ppm, 0.45 ppm and 0 ppm), which the fishes were exposed to a period of 21 days. At end of 7 days, 6 fishes were killed. The changes of the genomic of tilapia's gill was detected by using the RAPD-PCR technique. Heavy metal analysis (open acid digestion technique and ICP-MS) was conducted to determine the accumulation of copper in the dry weight of the gills. No significant changes occurred in the genomic of the gills after 21 days exposure of copper in all of the exposure concentration. 0.0878 ppb and 0.00242 ppb of Cu^{2+} was found from the highest to the lowest concentration. After 7, 14 and 21 days exposure to the copper, no genomic changes were obtained. As a conclusion, the genomic of the tilapia gills shows no changes for all concentration in the exposure period.

ABSTRAK

Ciri-ciri DNA normal dan rosak akibat pendedahan terhadap kuprum.

Tujuan utama kajian ini adalah untuk meramal perubahan genomik organisma akuatik terhadap pendedahan kepada logam berat. Anak ikan tilapia merah (*Oreochromis niloticus*) (0.5 cm-5.0 cm) telah diberi kuprum. Kajian ini telah dijalankan dalam tiga peringkat yang saling berkaitan di antaranya. Bahagian pertama ialah ujian penentuan ketoksikan akut kuprum. Dengan menggunakan kaedah probit, 96-h LC₅₀ Cu²⁺ adalah 1.82 ppm. Nilai ini kemudian digunakan untuk membentuk 5 jenis kepekatan (1.82 ppm, 1.36 ppm, 0.91 ppm, 0.45 ppm dan 0 ppm) yang mana anak ikan diberi kuprum selama 21 hari. 6 ikan dibunuh pada setiap 7 hari. Perubahan pada genomik insang tilapia dikesan dengan menggunakan kaedah RAPD-PCR. Analisis logam berat (kaedah asid terbuka dan ICP-MS) juga dijalankan untuk menentukan akumulasi Cu²⁺ pada tisu insang yang kering. Tidak ada sebarang perubahan didapati setelah diberi kuprum kepada semua kepekatan selama 21 hari itu. 0.00242 sehingga 0.0878 ppb berat kering akumulasi Cu²⁺ telah dijumpai dalam ikan yang diberi kuprum dari kepekatan rendah ke kepekatan tertinggi. Selepas 21 hari diberi kuprum, didapati tiada perubahan genomik terhadap ikan tilapia. Hal ini disebabkan oleh kuprum yang merupakan logam perlu didalam semua tisu hidupan. Secara kesimpulannya, tiada perubahan genomik yang ditunjukkan dalam semua kepekatan dalam tempoh 21 hari tersebut.