

DETERMINATION OF ISOELECTRIC FOCUSING METHOD
FOR PROTEOMIC ANALYSIS OF
PURPLE VARIETY OF *Orthosiphon stamineus*

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ABSTRACT

An ancient medicinal plant known as *Orthosiphon stamineus* purple variety exhibiting multiple remedial effects by the bioactive compound had been reported. However, no study on its protein separation work was documented. Indeed, two-dimensional (2D) electrophoresis systems have few major issues such as complex time-consuming procedure and poor reproducibility result. Therefore, this proteomic research can be utilized as preliminary platform in identification of protein responsible for the production of bioactive compound and its application as remedies. Different sample require different voltage of isoelectric focusing (IEF) to separate the protein due to its isoelectric point (pI) hence this study was conducted in order to determine the best IEF method producing reproducible protein spots and to determine the protein spot distribution and pattern from 2D proteomics of *O. stamineus*. Initially, protein extraction utilizing unmodified and modified (1 mL of 80% methanol in 0.1M ammonium acetate, 0.6 mL of each phenol and SDS buffer in 5 % β -mercaptoethanol, and 0.15mL lysis buffer) phenol extraction with three preliminary washes were performed. The extracted protein was then quantified by Bradford Assay whereas its quality was checked via Sodium Dodecyl Sulfate- Polyacrylamide Gel Electrophoresis (SDS-PAGE). The modified extraction method resulted in higher spots distribution compared to the unmodified method. In this study, three different IEF method of method 1 (2000V), method 2 (4000V), and method 3 (8000V) were applied before the protein spots pattern and distribution were analysed further. It was found that method 2 was the best IEF method for this plant as 2-fold higher reproducible protein spots were produced. In fact, 15 proteins were expected to be associated with the protein spots obtained that may have high relation with photosynthesis and antioxidant activity. The findings suggested that numerous valuable proteins are expressed in *O. stamineus*. Hence, the proteins mechanism and properties should be revealed to serve the medical sector in the future.

ABSTRAK

Pokok perubatan purba iaitu *Orthosiphon stamineus* berwarna ungu mempunyai pelbagai kesan pemulihan kesan daripada sebatian bioaktif telah dilaporkan sebelum ini. Tetapi, tiada sebarang kajian tentang pemisahan protein pokok ini didokumenkan. Sistem elektroforesis dua dimensi (2D) mempunyai beberapa kelemahan iaitu prosedur yang rumit dan kadar kebolehulangan yang rendah. Oleh itu, kajian proteomik terhadap pokok ini dapat digunakan sebagai platform kerja awal dalam mengenal pasti protein yang bertanggungjawab menghasilkan sebatian bioaktif dan aplikasinya sebagai penawar. Voltan isoelectric focusing” (IEF) yang diperlukan untuk memisahkan protein berbeza bagi setiap sampel kerana “isoelectric point” (pI). Kajian ini dijalankan bagi mengenal pasti kaedah IEF terbaik yang menghasilkan titik protein berulang dan mengenal pasti corak dan taburan titik protein bagi pokok ini. menggunakan kaedah proteomik 2D. Pada mulanya, protein diekstrak dengan kaedah pengekstrakan menggunakan fenol yang asal dan yang diubah suai (1 mL 80% methanol dalam 0.1M ammonium acetate, 0.6 mL phenol dan SDS buffer dalam 5 % β -mercaptoethanol, dan 0.15mL lysis buffer) sebelum kuantiti dan kualiti protein dikenal pasti menggunakan Bradford assay dan sodium dodecyl sulfate-poliakrilamid gel elektroforesis (SDS-PAGE). Kaedah pengekstrakan menggunakan fenol yang telah diubah suai menghasilkan lebih banyak bilangan titik protein berbanding kaedah yang asal. Selain itu, tiga kaedah IEF yang digunakan dalam kajian ini ialah kaedah 1(2000V), kaedah 2(4000V) dan kaedah 3(8000V). Kaedah 2 menghasilkan corak titik protein dua kali ganda lebih tinggi berbanding kaedah IEF lain. Seperkara lagi, 15 protein yang dijangka mempunyai hubungkait dengan titik protein dari elektroforesis 2D telah dikenal pasti dan mungkin mempunyai kaitan dengan proses fotosintesis dan aktiviti antioksidan pokok ini. Kesimpulannya, terdapat pelbagai protein berharga dihasilkan oleh pokok ini. Oleh itu, sifat dan mekanisma protein ini harus dikaji agar dapat dimanfaatkan dalam bidang perubatan di masa hadapan.

TABLE OF CONTENTS

CHAPTER	TITLE	PAGE
	DECLARATION	ii
	ACKNOWLEDGEMENTS	iii
	ABSTRACT	iv
	ABSTRAK	v
	TABLE OF CONTENTS	vi
	LIST OF TABLES	ix
	LIST OF FIGURES	x
	LIST OF SYMBOLS	xi
	LIST OF APPENDICES	xii
	LIST OF ABBREVIATIONS	xiii
1	INTRODUCTION	
1.1	Background of Study	1
1.2	Problem Statement	2
1.3	Objectives	3
1.4	Significance of Work	3
1.5	Scope of Work	3

2	LITERATURE REVIEW	
2.1	<i>Orthosiphon stamineus</i>	5
2.2	Morphology of <i>Orthosiphon stamineus</i>	7
2.3	Pharmacological Properties of <i>Orthosiphon stamineus</i>	8
2.3.1	Diuretic and Antidiabetic Properties of <i>Orthosiphon stamineus</i>	9
2.3.2	<i>Orthosiphon stamineus</i> Versus Bacteria, Fungi, and Cancer Cell	10
2.4	Plant Proteomics	11
3	MATERIALS AND METHODS	
3.1	Experimental Design	13
3.2	Plant Material	14
3.3	Protein extraction	15
3.3.1	Phenol/SDS buffer with 3 preliminary washes	15
3.3.2	Modified Phenol/SDS buffer with 3 preliminary washes	16
3.4	Protein Quantification	16
3.5	Protein Separation	18
3.5.1	One-dimensional SDS-PAGE	18
3.5.2	Two-dimensional Gel Electrophoresis	19
4	RESULTS AND DISCUSSION	
4.1	Determination of the best IEF method in 2D proteomics of purple <i>O. stamineus</i>	21
4.2	Determination of the protein spot distribution and pattern from 2D proteomics of purple <i>Orthosiphon</i> <i>stamineus</i> .	29
4.3	Expected protein from 2D protein spots	30

5	CONCLUSION	
5.1	Conclusion	34
5.2	Future Works	35
	REFERENCES	36
	APPENDIX A	43
	APPENDIX B	44
	APPENDIX C	45

LIST OF TABLES

TABLE	TITLE	PAGE
2.2	Morphological characteristics of white and purple <i>O. stamineus</i>	7
3.1	Volume of BSA standard and distilled water for specific BSA standard concentration	17
3.2	Isoelectric focusing (IEF) method	20
4.1	Protein quantification of the extracted protein samples	22
4.2	Expected proteins of purple <i>O. stamineus</i> .	33

LIST OF FIGURES

FIGURE	TITLE	PAGE
2.1	(a) White (b) Purple variety of <i>O. stamineus</i>	6
3.1	Flowchart of the experimental design	14
4.1	Protein spot distribution pattern obtained from purple <i>O. stamineus</i> run on 7cm IPG strips using phenol extraction with 3 solvent washes using IEF (i) method 2, (ii) method 3, modified method of phenol extraction with 3 solvent washes using IEF (iii) method 2 and (iv) method 3 loaded at protein amount of 50 µg.	24
4.2	Protein spot distribution pattern obtained from purple <i>O. stamineus</i> run on 7cm IPG strips using (i) IEF method 1, (ii) method 2 and (iii) method 3 on 12% (w/v) SDS gel loaded at 50 µg.	27
4.3	Protein spot distribution pattern obtained from purple <i>O. stamineus</i> run on 7cm IPG strips using 4000V IEF producing highest number of 35 protein spots distribution. The alphabets represent expected protein spots.	31

LIST OF ABBREVIATIONS

1D	-	One dimensional
2-DE	-	Two dimensional electrophoresis
APS	-	Ammonium persulfate
BSA	-	Bovine serum albumin
CHAPS	-	3-(3-Cholamidopropyl)dimethylammonia)-1-Propanesulfonic acid
CBB	-	Coomassie Brilliant Blue
cm	-	Centimeter
DNA	-	Deoxyribonucleic acid
DTT	-	Dithiothreitol
EDTA		Ethylenediaminetetraacetic acid
<i>et al.</i>	-	And friends
HCl	-	Hydrochloric acid
IAA	-	Indole-3-acetic acid
IEF	-	Isoelectric focusing
IPG	-	Immobilised pH gradient
KPO ₄		Potassium phosphate
QB	-	Quenching buffer
RNA	-	Ribonucleic acid
Rpm	-	Rotary per minute
SDS	-	Sodium Dodecyl Sulfate
SDS-PAGE	-	Sodium Dodecyl Sulfate- Polyacrylamide Gel Electrophoresis
SEM	-	Standard Error Mean
TCA	-	Trichloroacetic acid
TEMED	-	Tetramethylethylenediamine

LIST OF SYMBOLS

A_{595}	-	Absorbance at 595 nanometer
β	-	Beta
$^{\circ}\text{C}$	-	Degree celcius
G	-	Gram
kDa	-	Kilo Dalton
μg	-	Microgram
$\mu\text{g/mL}$	-	Microgram per milliliter
$\mu\text{g/g}$	-	Microgram per gram
μL	-	Microliter
mg	-	Milligram
mL	-	Milliliter
mM	-	Milli Molar
M	-	Molar
%	-	Percent
V	-	Voltage
v/v	-	Volume per volume
w/v	-	Weight per volume

LIST OF APPENDICES

APPENDIX	TITLE	PAGE
A	Standard curve of BSA standard concentration against absorbance (A_{595}).	43
B	<i>O.stamineus</i> protein electrophoretic pattern on 12 % (w/v) 1D SDS-PAGE gel (M = marker, S1 = Sample 1, S2 = Sample 2, S3 = Sample 3, S4 = Sample 4, S5 = Sample 5, S6 = Sample 6).	44
C	<i>O.stamineus</i> protein electrophoretic pattern on 12 % (w/v) 1D SDS-PAGE gel.	45

CHAPTER 1

INTRODUCTION

1.1 Background of study

Orthosiphon stamineus Benth or Cat's Whiskers or "misai kucing" is from the family of Lamiaceae. *O. stamineus* is an ancient medicinal herb that had been widely used since many centuries before (Wiart, 2000). This herbal tea is a well-known herbal drink consumed to cure critical human disease (Basheer and Majid, 2010). It is commonly found in South East Asia used for treatment of various disorders and ailments including urinary lithiasis, edema, eruptive fever, influenza, rheumatism, hepatitis, jaundice and biliary lithiasis (Tran, 1990).

Extracts of *O. stamineus* leaves contain wide variety of compounds, including flavonoids, terpenoids or organic acids such as rosmarinic or caffeic acid (Tezuka *et al.*, 2000). The polyphenols for instance are able to provide enzyme inhibition and antioxidant activity which lead to therapeutic effect to its consumer (Hossain and Rahmanjung, 2011). The various remedial effects of this herb has attracted researcher's interest to further investigate its valuable components and mechanism of action. Proteomic analysis approach is a great option to complement previous studies as it enables the study of protein properties extracted from the valuable herbs. Indeed, the study of the plant proteomes wholly contributes to its medicinal value mainly for human sake.

Proteomics was traditionally used for protein separation utilizing the phenomenal two-dimensional gel electrophoresis technology (Celis and Bravo, 1984). This enables quantitative analysis of protein amounts available in the complex extracts according to the protein's isoelectric point and molecular weight (*et al.*, 1989). The fundamentals and principles of two dimensional (2D) proteomics have been established and practiced in numerous successful studies (Doleckova *et al.*, 2012; Barrabes *et al.*, 2010; Hartinger *et al.*, 1996; Tang *et al.*, 1997). Hence, its advantages and limitations are widely found in reviews and articles (Rogowska *et al.*, 2013).

1.2 Problem statement

O. stamineus is well known for its medicinal values contributed by the bioactive compound which attracts researchers' interest. Still, no study on its protein separation work was reported due to the complexity of the protein and the proteomic protocol itself. In fact, this plant specifically require more starting material to extract its protein sample compared to other plant since it contain high impurities. The most commonly used protein separation alternative is the 2D proteomics methods; combination of IEF protocol and molecular weight (Rabilloud *et al.*, 2010). However, 2D electrophoresis systems still have few major issues that need to be addressed, such as complex time-consuming experimental procedure and poor reproducibility of the result (Barrabes *et al.*, 2010). In this study, the experiment was conducted in order to determine the best IEF protocol producing reproducible protein spot of *Orthosiphon stamineus* thus saving time. The findings of the study can be further utilised for identification of protein responsible in producing valuable bioactive compound and application of the protein in medical area.

1.3 Objectives

The objectives of this study are as follows:

- i. To determine the best IEF method producing reproducible protein spots from purple variety of *Orthosiphon stamineus*.
- ii. To determine the protein spot distribution and pattern from 2D proteomics of purple *Orthosiphon stamineus*.

1.4 Significance of work

Various medicinal value that *O. stamineus* offers since decades ago has attracted the researchers' interest (Tran, 1990). Numerous study were done on its pharmacological and phytochemical properties. On the other hand, fewer study related to the herb's protein content was established. The findings of this project are beneficial as the best IEF method producing reproducible spots of extracted protein from *O. stamineus* was determined thus saving time consumed in performing IEF. The 2D proteomic analysis of this herbal plant can be utilized as preliminary work platform for further protein identification especially related to its medicinal potential. Hence, those valuables protein will be able to serve the mankind in the pharmaceutical industry and the medical sector according to its medicinal properties.

1.5 Scope of work

The purple variety of *O. stamineus* was chosen in this study since it is pigmented hence it was expected to contain higher protein content compared to the white variety. In order to achieve the objectives of the study, there are 3 main procedures involved. These procedures are, protein extraction from the *O. stamineus* leaves using phenol extraction with 3 solvent wash, protein quantification utilizing

Bradford assay, and protein separation technique which focuses on the 2D gel electrophoresis. The 2D gel electrophoresis involves the isoelectric focusing (IEF) and SDS-PAGE principles which separate extracted protein according to its pI and molecular weight respectively. In fact, different protein sample require different voltage of IEF method in order to separate and focus the protein spots according to its pI. In this study, three IEF methods of method 1 (2000V), method 2 (4000V), and method 3 (8000V) were applied to compare its protein spot distribution and pattern.

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