

**IDENTIFICATION AND CHARACTERIZATION
OF PROTEINS FROM SABAH EDIBLE
SEaweeds BY USING PROTEOMICS
APPROACH**

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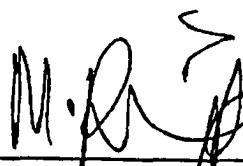
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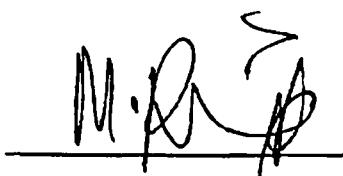
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ABSTRACT

The present study was carried out to examine the proteome profile of selected seaweed species from Sabah waters to explore their nutritional properties. Moisture content was a major component in seaweed followed by carbohydrate, mineral, and protein. The results showed significant differences ($p<0.05$) in proximate compositions and total phenolic contents among several species of red, green and brown seaweeds. Studies showed that red seaweed higher in carbohydrate (63 to 74 % dry weight) and protein (6 to 15 % dry weight) content. Brown seaweed higher in mineral (22 to 45 % dry weight) and fibre (21 to 45 % dry weight) content. Meanwhile, green seaweed higher in total phenolic content (48 to 52 mg PGE/g dry weight sample). Using *Caulerpa lentillifera*, four different protocols for protein extraction were evaluated, namely: (i) no protein precipitation with fractionation; (ii) Trichloroacetic acid (TCA)-acetone precipitation; (iii) TCA-acetone precipitation with fractionation; and (iv) Phenol extraction methanol-ammonium acetate precipitation. Overall, the phenol extraction method gave a higher protein yield and some protein spots resolved with a satisfactory 2D-PAGE image quality than the other methods. Thus this method was used to extract total soluble protein from all 15 seaweed samples. There was a strong positive correlation between crude protein and total soluble protein concentration ($r=0.952$; $p=0.007$) in these fifteen varieties of seaweed. A distinctive protein pattern was observed in the SDS-PAGE gels between three different seaweed classes of green, red and brown colours. Using this method of extraction, it managed to get 2D-PAGE proteome map for three seaweeds species, namely *Caulerpa lentillifera*, *Kappaphycus alvarezii* var. *aring-aring* and *Sargassum polycystum*. The image analysis of 2D-PAGE was performed with Progenesis SameSpots v 4.5 and four sets of comparison of the 2D-PAGE gel image were done, Set 1: *Caulerpa lentillifera* fronds-CLB vs *Caulerpa lentillifera* stolon-CLT; Set 2: *Kappaphycus alvarezii* small branches-KAK vs *Kappaphycus alvarezii* big branches-KAB; Set 3: *Sargassum polycystum* vacuoles-SPB vs *Sargassum polycystum* leaves-SPD vs *Sargassum polycystum* stolon-SPT; and Set 4: *Caulerpa lentillifera*-CLS vs *Kappaphycus alvarezii* -KAS vs *Sargassum polycystum* -SPS to detect differentially expressed protein spot. Results of image analysis revealed 77 differentially expressed proteins for experiment set of CLB vs. CLT, 98 differentially expressed proteins for experiment set of KAK vs. KAB, 93 differentially expressed proteins for experiment set of SPB vs. SPD vs. SPT and 148 differentially expressed proteins for experiment set of CLS vs. KAS vs. SPS. Only 20 differentially expressed proteins from all four sets of experiments have been successfully identified and characterized by using nano-LC-MS/MS (Orbitrap) and Bioinformatics Software PEAKS Studio 7. Most of the identified proteins were a non-enzyme protein such as light-harvesting phycobiliprotein, light-harvesting complex, histone, and elongation factor. The enzyme proteins identified were ribulose bisphosphate carboxylase and glyceraldehyde-3-phosphate dehydrogenase. Regarding health benefits, the potential evaluation of the peptide profile results showed that protein extracts from the three seaweeds species contain peptide sequences with a variety of beneficial health effects. From among 44 types of biological activity characteristics of peptides, as listed in the BIOPEP database, motifs with the Dipeptidyl Peptidase IV inhibitors and ACE inhibitory activity occur most frequently in all protein sequences from the seaweed samples. Others bioactive peptide present, such as antioxidant, stimulating, regulating, antiamnestic, antithrombotic and many other activities. Therefore, this protein raw material, which currently remains a relatively untapped reservoir, has the potential to act as a resource for the generation of bioactive peptides with potential health-promoting and disease-preventing properties. Hopefully, these findings will contribute towards seaweed protein utilization as alternative sources of peptide-based nutraceuticals for human health promotion.



ABSTRAK

PENGENALPASTIAN DAN PENCIRIAN PROTEIN DARIPADA RUMPAI LAUT SABAH DENGAN MENGGUNAKAN KAE DAH PROTEOMIK

Kajian yang telah dijalankan ini adalah untuk menilai profil protein pada spesis rumpai laut yang terpilih daripada perairan Sabah yang bertujuan untuk meneroka serta melihat ciri-ciri khasiat yang ada. Kandungan kelembapan merupakan komponen utama dalam rumpailaut diikuti oleh karbohidrat, mineral dan protein. Keputusan kajian menunjukkan perbezaan yang bererti ($p<0.05$) pada analisis proksimat dan jumlah kandungan fenolik di antara beberapa spesis rumpai laut merah, hijau dan perang. Kajian mendapati bahawa rumpailaut merah mengandungi karbohidrat (63% hingga 47% berat kering) dan kandungan protein (6% hingga 15% berat kering) yang tertinggi. Rumpailaut perang mengandungi mineral (22% hingga 45% berat kering) dan serat (21% hingga 45% berat kering) yang tertinggi. Manakala rumailaut hijau mengandungi jumlah kandungan fenol yang tertinggi (48 hingga 52 mg PGE/g berat kering sampel). Menggunakan *Caulerpa lentillifera*, empat kaedah pengekstrakan protein yang berbeza telah dikaji iaitu: (i) pembahagian protein tanpa pemendakan; (ii) asid trikloroasetik (TCA)-pemendakan aseton; (iii) Pemendakan asid trikloroasetik (TCA) dengan pembahagian; dan (iv) pengekstrakan fenol-pemendakan methanol-amonium asetat. Secara keseluruhan, pengekstrakan fenol memberikan hasil protein yang tinggi dan bilangan tompok protein yang terpisah serta gambaran 2D-PAGE yang sangat memuaskan berbanding kaedah lainnya. Oleh yang demikian kaedah ini digunakan untuk mengekstrak jumlah kandungan protein larut pada kesemua 15 sampel rumpailaut. Keputusan juga menunjukkan korelasi positif yang kuat di antara nilai protein kasar dengan kepekatan jumlah protein larut ($r=0.952$; $p=0.007$) pada limabelas varieti rumpailaut tersebut. Corak protein yang berbeza juga ditunjukkan pada gel SDS-PAGE diantara spesis rumpailaut hijau, merah dan perang. Menggunakan kaedah pengekstrakan yang sama, ianya telah menghasilkan peta protein 2D-PAGE untuk tiga spesis rumailaut iaitu, *Caulerpa lentillifera*, *Kappaphycus alvarezii* var. aring-aring dan *Sargassum polycystum*. Analisis gambar 2D-PAGE telah dibuat menggunakan Progenesis SameSpots v 4.5 dan empat set kajian perbandingan gel 2D-PAGE telah dijalankan iaitu, Set 1: *Caulerpa lentillifera* 'fronds'-CLB dengan *Caulerpa lentillifera* 'stolon'-CLT; Set 2: *Kappaphycus alvarezii* cabang halus-KAK dengan *Kappaphycus alvarezii* cabang besar-KAB; Set 3: *Sargassum polycystum* vakul-SPB dengan *Sargassum polycystum* daun-SPD dengan *Sargassum polycystum* 'stolon'-SPT and Set 4: *Caulerpa lentillifera*-CLS dengan *Kappaphycus alvarezii*-KAS dengan *Sargassum polycystum*-SPS bagi mencari tompok protein yang menunjukkan perbezaan. Keputusan analisis gambar mendapati 77 tompok protein yang berbeza pada set CLB dengan CLT, 98 tompok protein yang berbeza pada set KAK dengan KAB, 93 tompok protein yang berbeza pada set SPB dengan SPD dengan SPT and 148 tompok protein yang berbeza pada set CLS dengan KAS dengan SPS. Hanya 20 tompok protein yang berbeza telah dikenalpasti dan pencirian dilakukan dengan jayanya menggunakan nano LC-MS/MS (Orbitrap) dan perisian bioinformatik PEAKS studio 7. Kebanyakan protein yang dikenalpasti adalah bukan enzim seperti cahaya-tuai pikobiliprotein, kompleks cahaya-tuai, histona dan faktor pemanjangan. Protein enzim yang dikenalpasti adalah seperti ribulosa bifosfat karboksilase dan gliseroldihida-3-fosfat dihidrogenase. Merujuk kepada kebaikan kesihatan, penilaian potensi mendapati ekstrak protein daripada ketiga-tiga spesis rumpai laut mengandungi urutan rantaian peptida yang mempunyai pelbagai khasiat kesihatan. Daripada 44 jenis ciri aktiviti biologi yang terdapat pada rantaian peptida, seperti yang tersenarai pada pengkalan data BIOPEP, motif dengan ciri aktiviti perencat 'Dipeptidyl Peptidase-IV' dan aktiviti perencatan ACE hadir pada kekerapan yang tertinggi tinggi pada semua urutan rantaian protein pada ketiga-tiga rumpai laut. Peptida

bioaktif lain yang hadir adalah seperti, antioksida, rangsangan, pengaturan, antiamnestik, antithrombotik dan lain-lain lagi. Oleh itu, sumber protein ini, yang mana ianya masih lagi belum dibangunkan, mempunyai potensi yang baik untuk menjadi sumber penghasilan peptida bioaktif dengan ciri-ciri kebaikan kesihatan dan juga menjadi penghalang kepada penyakit. Adalah diharapkan dengan dapatan kajian ini dapat menyumbang kepada penggunaan protein daripada rumpai laut sebagai sumber alternatif untuk bekalan bahan asas nutraceutikal peptida untuk penjagaan kesihatan manusia.



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LIST OF ABBREVIATION

%	-	Percentage
°C	-	Degree celcius
µg/g	-	Microgram per gram
µg/L	-	microgram perlitre
1D	-	One dimension
2D	-	Two dimension
AcDP	-	Acetone dried powder
ACE	-	Angiotensin-converting enzyme
ACN	-	Acetonitrile
ANOVA	-	Analysis of variance
APS	-	Ammonium persulfate
BLAST	-	Basic Local Alignment Search Tool
BSA	-	Bovine serum albumin
CBB	-	Commasie brilliant blue
CHAPS	-	Chloromidopropylidemethylammonio-1-propane sulfonate
CID	-	Collision-induced dissociation
CuSO ₄	-	Copper (II) sulfate
DNA	-	Deoxyribonucleic acid
DTT	-	Dithiothreitol
ECD	-	Electron capture dissociation
ESI	-	Electrospray ionization
<i>et al.</i> ,	-	and all
ETD	-	Electron transfer dissociation
FAO	-	Food and agricultural organization
FDR	-	False discovery rates
g	-	Gram
HCl	-	Hydrochloric acid
HPLC	-	High performance liquid chromatography
IEF	-	Isoelectrofocusing



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