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Furthermore, according to Na et al., the purified polysaccharides (glucuronogalactomannan-sulfates) from *C. fulvescens* considerably stimulated RAW264.7 cells and resulted in the secretion of cytokines and chemokines such as TNF- α , IL-6 and NO, through the expression of COX-2 and iNOS mRNAs [13]. The protective effects of the sulfated polysaccharides from *C. fulvescens* on the alcohol-induced gastric injury in rats were also reported by Hwang et al. [14]. The biological activities of the sulfated polysaccharides have been reported to be closely related with their chemical and molecular structures such as molecular weights and sulfate contents [15,16]. Therefore, a better understanding of the primary and secondary molecular structures for the polysaccharides may lead to the successful interpretation of their biological activities. However, despite the abundance and availability of *C. fulvescens*, relatively limited researches have been performed on its molecular structures and bioactivities of sulfated polysaccharides.

In this study, a sulfated polysaccharide from *C. fulvescens* was extracted in distilled water and subsequently fractionated using an ion-exchange chromatography. The purpose of this study was to investigate the chemical and molecular characteristics of the fractionated polysaccharides, to evaluate their immunomodulatory activity, and finally to obtain the relationship between their structures and bioactivities.

2. Materials and methods

2.1. Materials

The green seaweed *C. fulvescens* was collected from the coast of Wando, Chonnam province, Korea. The seaweed was thoroughly washed with tap water and air-dried at 60 °C. The dried raw material was milled using a blender, sieved (<0.5 mm) and stored in plastic bags at –20 °C before extracting the polysaccharide. All chemicals and reagents used in this work were analytical grade. RPMI-1640 medium and fetal bovine serum (FBS) were purchased from Lonza (Walkersville, MD, USA).

2.2. Extraction of crude polysaccharide

The milled green seaweed (20 g) was treated with 85% ethanol (EtOH, 200 mL) under constant mechanical stirring overnight at room temperature. The residual part was separated by centrifugation (10 °C, 8000 rpm, 10 min), rinsed with acetone, and then dried at room temperature. The dried biomass (10 g) was extracted with distilled water (200 mL) at 65 °C with stirring for 2 h. The extracts were centrifuged at 10,000 rpm for 10 min at room temperature, and the supernatants were collected. The supernatants were concentrated by evaporation under reduced pressure at 60 °C to approximately 200 mL. EtOH (99%) was added into the supernatants to obtain the final concentration of 70%, and the solution was placed at 4 °C overnight. The polysaccharide was obtained by the filtration of the solution with a membrane (0.45 μ m pore size, Whatman International, Maidstone, UK) and washed with EtOH (99%), followed by acetone, and then dried at room temperature. The precipitated polysaccharide was referred to as the crude polysaccharide and the yield was calculated according to the dried biomass obtained after treating the milled sample with 85% EtOH.

2.3. Fractionation of the polysaccharides

The crude polysaccharide (250 mg) was dissolved in distilled water (10 mL), and then the solution was injected into a DEAE Sepharose fast flow column (17-0709-01, GE Healthcare Bio-Science AB, Uppsala, Sweden). The polysaccharides were eluted with distilled water to obtain a non-adsorbed fraction and developed subsequently with a stepwise NaCl gradient (0.5–2 M) to

wash the charged polyanionic polysaccharides. Three fractions were obtained based on the phenol–H₂SO₄ assay, referring to as F₁, F₂ and F₃ [17]. These fractions were concentrated, dialyzed and lyophilized.

2.4. Chemical characterization of polysaccharides

Carbohydrate content of the polysaccharide was estimated by the phenol–sulfuric acid assay using glucose as a standard [17]. Sulfate content of the polysaccharide was determined by the BaCl₂ gelatin method using K₂SO₄ as a standard after hydrolysis of the polysaccharides with 0.5 M HCl [18]. Protein content of the polysaccharide was determined by the Lowry method using DC Protein assay kit (Bio-Rad, USA) [19]. Uronic acid content of the polysaccharide was determined by sulfamate/*m*-hydroxydiphenyl assay using glucuronic acid as standard [20].

2.5. Monosaccharide composition of the polysaccharides

The monosaccharide composition of the polysaccharides was performed using a HPLC system, which consisted of a pump (Waters 510, Waters, Milford, MA, USA), an injection valve (Model 7010, Rheodyne, Rohnert Park, CA, USA) with a 20 μ L sample loop, a column (Carbohydrate analysis column, 4.6 mm \times 250 mm, Waters, Milford, MA, USA) and a RI detector (Waters 410). After the hydrolysis of the polysaccharide (60 mg) in 2 M trifluoroacetic acid (TFA) at 120 °C for 5 h, TFA was removed by the evaporation with a dried stream of nitrogen. The hydrolysates were injected into the HPLC system. Acetonitrile (80%) was used as a mobile phase at a flow rate of 2 mL/min. The following neutral monosaccharides were used as references: rhamnose, xylose, mannose, galactose and glucose.

2.6. Deproteination of F₂ polysaccharide

The proteins in F₂ fraction were hydrolyzed using Flavourzyme (#2384, Novozymes, Tianjin, China) after dissolution of the polysaccharides in 0.1 M phosphate buffer (pH 7.0). The sample solution was incubated at 50 °C for 24 h. The resulting hydrolysate was heated in boiling water for 10 min to inactivate the enzyme, centrifuged at 7000 rpm for 10 min and dialyzed in a membrane (#3247027, Spectrum Laboratories, Compton, CA, USA) against distilled water, and eventually lyophilized.

2.7. Glycosidic linkage analysis

The glycosidic linkage analysis of the most immunoenhancing F₂ polysaccharides was carried out using the method of Hakomori with slight modifications [21]. The partially deproteinized polysaccharide (2–3 mg) was dissolved in DMSO (0.5 mL) under nitrogen, and then methylated with CH₃I (0.3 mL) and dried NaOH powder (20 mg). Partially methylated alditol acetates were prepared from fully methylated samples by acid hydrolysis with 4 M TFA at 100 °C for 6 h, followed by a reduction of the hydrolysates in water using NaBD₄ and an acetylation with acetic anhydride. The partially methylated alditol acetates were analyzed by a gas chromatography–mass spectrometry (GC–MS) (6890 N/MSD 5973, Agilent Technologies, Santa Clara, CA) using a HP-5MS capillary column (30 m \times 0.25 mm \times 0.25 μ m) (Agilent Technologies, Santa Clara, CA) under previously reported conditions [22].

2.8. Nuclear magnetic resonance (NMR) spectroscopy

The partially deproteinized fraction F₂ (30 mg) was dissolved in D₂O (0.5 mL), and subjected to the spectrometer at 50 °C. ¹H and ¹³C NMR spectra were recorded on a JEOL ECA-600 spectrometer

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