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# Structural elucidation of an extracellular polysaccharide produced by the marine fungus *Aspergillus versicolor*

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#### ABSTRACT

A homogenous extracellular polysaccharide, designated AWP, was isolated from the fermented liquid of the marine fungus *Aspergillus versicolor* from the coral *Cladiella* sp. and purified by anion-exchange and size-exclusion chromatography (SEC). Chemical and spectroscopic analyses, including one- and two-dimensional nuclear magnetic resonance (1D and 2D NMR) spectroscopy showed that AWP consisted of glucose and mannose in a molar ratio of 8.6:1.0, and its average molecular weight was estimated to be 500 kDa. AWP is a slightly branched extracellular polysaccharide. The backbone of AWP is mainly composed of  $(1\rightarrow6)$ -linked  $\alpha$ -D-glucopyranose residues, slightly branched by single  $\alpha$ -D-mannopyranose units attached to the main chain at C-3 positions of the glucan backbone. The investigation demonstrated that AWP is a novel extracellular polysaccharide different from those of other marine microorganisms.

#### 1. Introduction

Marine organisms, such as sponges, sea squirts, corals, worms and algae, host diverse and abundant symbiotic microorganisms (Olson & Kellogg, 2010). The microorganisms may be saprophytic, pathogenic, or may serve important positive functions in the ecosystem, and protect their hosts from pathogens (Armstrong, Yan, Boyd, Wright, & Burgess, 2001; Harder, Lau, Dobretsov, Fang, & Qian, 2003; Harel et al., 2008; Kramarsky-Winter et al., 2006; Rohwer, Breitbart, Jara, Azam, & Knowlton, 2002; Siboni et al., 2010). Recently, the interest in extracellular polysaccharides produced by symbiotic microorganisms from marine organisms is increasing because of the microorganisms are mostly new or inadequately described species (Miranda et al., 2008). The marine bacterium Shewanella colwelliana from Crassostrea virginic excreted a kind of acidic polysaccharide, which contained mannose, glucose, galactose and pyruvic acid (Sledjeski & Weiner, 1993). Aeromonas sp. HYD154 from a deep-sea hydrothermal vent polychaete annelid produced an acidic polysaccharide, with glucose, galactose, glucuronic acid, galacturonic acid and 4,6-pyruvic acidic galactose (Cambon-Bonavita, Raguenes, Jean, Vincent, & Guezennec, 2002).

The coral-associated fungus *Aspergillus versicolor* LCJ-5-4 could produce a mannoglucan, with the backbone of  $(1\rightarrow6)$ -linked  $\alpha$ -D-glucopyranose and  $(1\rightarrow2)$ -linked  $\alpha$ -D-mannopyranose units (Chen et al., 2012). The extracellular polysaccharides produced by symbiotic microorganisms from marine organisms represent a potential source to be explored (Dobretsov & Qian, 2004). In the present work, a novel extracellular polysaccharide was obtained from the culture liquids of the marine fungus *A. versicolor* from the coral *Cladiella* sp., and its structure was investigated by a combination of chemical, chromatographic (SEC, HPGPC, GC–MS) and spectroscopic (FTIR, 1D and 2D NMR) methods.

#### 2. Methods

#### 2.1. Materials

Dextran standards ( $M_W$ : 788, 404, 112, 47.3, 22.8, 11.8 and 5.9 kDa) were from Showa Denko K.K. (Tokyo, Japan). D-Glucose, D-mannose, D-galactose, L-rhamnose, D-xylose, L-fucose and D-glucuronic acid were from Sigma–Aldrich (St. Louis, MO, USA). Dialysis membranes (flat width 44 mm, molecular weight cut off 3500) were from Lvniao (Yantai, China). Q Sepharose Fast Flow and Sephacryl S-400/HR were from Amersham Biosciences (Uppsala, Sweden).

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#### 2.2. Microbial strains and culture conditions

A. versicolor was isolated from the coral Cladiella sp. collected from Hainan, China. It was identified according to its morphological characteristics and 18S rRNA sequences. Briefly, the fungus was cultivated in a liquid medium containing sorbitol (20 g/L), maltose (20 g/L), monosodium glutamate (10 g/L), KH<sub>2</sub>PO<sub>4</sub> (0.5 g/L), MgSO<sub>4</sub>·7H<sub>2</sub>O (0.3 g/L), tryptophane (0.5 g/L), yeast extract (3 g/L), sea salt (33.3 g/L), pH 6.5 at 20 °C for 30 days, and 90 L of fermented broth were collected (Zhuang et al., 2011).

#### 2.3. Isolation and purification of the extracellular polysaccharide

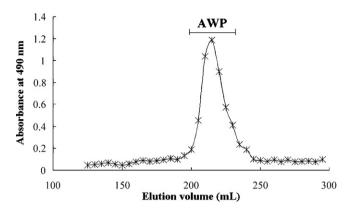
The fermented broth was filtered through cheesecloth to separate into fermentation liquids and mycelium. The fermentation liquids from A. versicolor cultures were concentrated, and three volumes of 95% (v/v) ethanol were added. The resulting precipitate was recovered by using centrifugation at 8000 rpm for 10 min, and dialyzed in a cellulose membrane (flat width 44 mm, molecular weight cut off 3500) against distilled water for 48 h. The retained fraction was vacuum-dried, and the protein in the fraction was removed by the method of Sevag (Matthaei, Jone, Martin, & Nirenberg, 1962). The crude polysaccharide was fractionated on a Q Sepharose Fast Flow column (300 × 30 mm) coupled to an AKTA FPLC system, eluting with 0, 0.3 and 2.5 M NaCl (Chen et al., 2012). The total sugar content of the fractions was determined by the phenol-sulfuric acid method. The fractions eluted with distilled water were pooled, and further purified on a Sephacryl S-400 column ( $100 \, \text{cm} \times 3 \, \text{cm}$ ) with 0.2 M NH<sub>4</sub>HCO<sub>3</sub> as eluent. The major polysaccharide fraction was gathered and freeze-dried.

#### 2.4. Analytical techniques

Total sugar content was measured by the phenol-sulfuric acid method using glucose as the standard (Dubois, Gilles, Hamilton, Rebers, & Smith, 1956). Protein content was assayed according to the modified Lowry method (Bensadoun & Weinstein, 1976). Sulfate content was measured according to Silvestri, Hurst, Simpson, and Settine (1982). Uronic acid content was determined by the carbazole-sulfuric acid method using glucuronic acid as standard (Blumenkrantz & Asboe-Hansen, 1973). The homogeneity and molecular weight was determined by high performance gel permeation chromatography (HPGPC) on a Shodex OHpak SB-804 HQ column, and the column calibration was performed with dextran standards (Li et al., 2012). The monosaccharide compositions were measured by gas chromatography (GC) (Chen et al., 2011). Sugar identification was done by comparison with reference sugars (D-glucose, D-mannose, D-galactose, L-rhamnose, D-xylose and L-fucose), and inositol was added as the internal standard. Calculation of the molar ratio of the monosaccharide was carried out on the basis of the peak area of the monosaccharide.

#### 2.5. Methylation analysis

Methylation analysis was performed according to the modified Hakomori method (Ciucanu & Kerek, 1984). Briefly, the polysaccharide in dimethyl sulfoxide was methylated using NaH and iodomethane. The completion of methylation was confirmed by FTIR spectroscopy as the disappearance of OH bands. After hydrolysis with 2 M trifluoroacetic acid at 105 °C for 6 h, the methylated sugar residues were converted to partially methylated alditol acetates by reduction with NaBH<sub>4</sub>, followed by acetylation with acetic anhydride. The derivatized sugar residues were extracted into dichloromethane and evaporated to dryness, dissolved again in 100 μL dichloromethane, and then analyzed by gas chromatography–mass spectrometry (GC–MS) on a HP6890II/5973



**Fig. 1.** Purification of the extracellular polysaccharide AWP from the marine fungus *A. versicolor* on a Sephacryl S-400 column.

instrument (Agilent Technologies Co. Ltd., USA) using a DB 225 fused silica capillary column (0.25 mm  $\times$  30 m). Identification of partially methylated alditol acetates was carried out on the basis of the retention time ( $t_R$ ) and its mass fragmentation patterns (Chen et al., 2012; Hung et al., 2012).

#### 2.6. IR spectroscopy analysis

For FTIR spectroscopy, the polysaccharide was mixed with KBr powder, ground and then pressed into a 1 mm pellets for FTIR measurement in the frequency range of 4000–500 cm<sup>-1</sup> with the resolution of 4.0 cm<sup>-1</sup> and 320 scans co-addition. FTIR spectrum of the polysaccharide was measured on a Nicolet Nexus 470 spectrometer by using the Nicolet Omnic software.

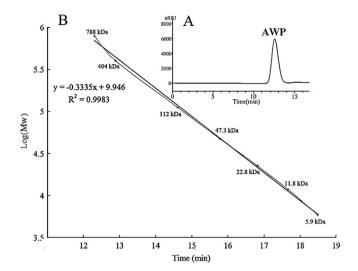
#### 2.7. NMR spectroscopy analysis

Seventy mg of polysaccharide was deuterium exchanged by two successive freeze-drying steps in 99%  $D_2O$  (Sigma–Aldrich, Canada) and then dissolved in 0.5 mL of 99.98%  $D_2O$ .  $^1H$  nuclear magnetic resonance (NMR) and  $^{13}C$  NMR spectra were measured at 23  $^{\circ}C$  using a JEOL JNM-ECP 600 MHz spectrometer by using Delta NMR processing and control software.  $^1H$ - $^1H$  correlated spectroscopy (COSY),  $^1H$ - $^{13}C$  heteronuclear multiple quantum coherence spectroscopy (HMQC) and  $^1H$ - $^{13}C$  heteronuclear multiple bond correlation spectroscopy (HMBC) experiments were also carried out. Chemical shifts are expressed in ppm using acetone as internal standard at 2.225 ppm for  $^1H$  and 31.07 ppm for  $^{13}C$ .

#### 3. Results and discussion

## 3.1. Purification and chemical composition of the extracellular polysaccharide AWP

The two crude extracellular polysaccharides from the fermented broth of coral-associated fungus A. versicolor were isolated on a Q Sepharose Fast Flow column, and the polysaccharide fraction eluted with 0.3 M NaCl was reported (Chen et al., 2012). Here, the polysaccharide fraction eluted with distilled water was further purified by a Sephacryl S-400 column (Fig. 1), and an extracellular polysaccharide AWP was obtained. The yield of AWP from crude extracellular polysaccharide (0.24 g/L) was about 25%. As shown in Fig. 2, AWP appeared as a single peak in the HPGPC chromatogram, and its average molecular weight was about 500 kDa. AWP did not contain any sulfate esters and protein, and its uronic acid content was below the detection limit. GC analysis demonstrated that AWP was composed of glucose and mannose in a molar ratio of 8.6:1.0. It is noted that the chemical composition of AWP was different from that of



**Fig. 2.** (A) HPGPC chromatogram of AWP on a Shodex Ohpak SB-804 column and (B) calibration curve using molecular weight standards.

the extracellular polysaccharide AVP isolated from coral-associated fungus *A. versicolor* (Chen et al., 2012).

#### 3.2. Methylation analysis

The methylation analysis may offer valuable information about the position of the glycosidic linkage (Vieira, Mulloy, & Mourao, 1991). The identification and the proportions of the methylated alditol acetates of AWP are listed in Table 1. Large amounts of 1,5,6-tri-O-acetyl-2,3,4-tri-O-methyl-glucitol and small amounts of 1,3,5,6-tetra-O-acetyl-2,4-di-O-methyl-glucitol and 1,5-di-Oacetyl-2,3,4,6-tetra-O-methyl-mannitol were detected in AWP, suggesting that AWP was mainly composed of  $(1\rightarrow 6)$ -linked glucopyranose with small amounts of  $(1\rightarrow3,6)$ -linked glucopyranose and  $(1\rightarrow)$  mannopyranose residues. AWP could be recognized as a slightly branched polysaccharide with a  $(1\rightarrow 6)$ -linked glucopyranose backbone, like a polysaccharide dextran main chain (Naessens, Cerdobbel, Soetaert, & Vandamme, 2005), and the mannopyranose side chains attached to C-3 position of  $(1\rightarrow 6)$ -linked glucopyranose residues. The molar ratios were in good agreement with the monosaccharide composition of AWP obtained from GC analysis. The linkage pattern of the polysaccharide AWP was different from those of other extracellular polysaccharides from marine microorganisms (Chen et al., 2012; Guo et al., 2010).

#### 3.3. IR spectrum analysis

IR spectrum of AWP was presented in Fig. 3. The broad and intense band at 3383 cm<sup>-1</sup> was the result of valent vibrations OH groups and valent vibration of H<sub>2</sub>O constitutional molecules (Mitić, Nikolić, Cakić, Premović, & Ilić, 2009), and the band at 2927 cm<sup>-1</sup> was attributed to the stretching vibration of C—H bond (Chen et al., 2011; Mitić et al., 2011). The band at 1650 cm<sup>-1</sup> was due to the bending vibrations of HOH, and the band at 1419 cm<sup>-1</sup> originated from the bending vibrations of O—H bond (Cakić, Mitić, Nikolić,

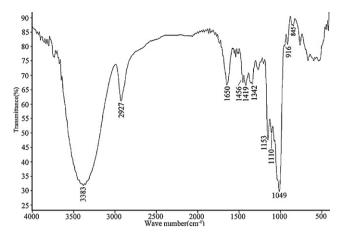


Fig. 3. IR spectrum of AWP.

Ilić, & Nikolić, 2008; Mitić, Cakić, & Nikolić, 2010; Mitić et al., 2009, 2011). The bands at 1456 and 1342 cm $^{-1}$  were the absorption peaks of variable angle vibration of C–H bond. The band at 1153 cm $^{-1}$  was assigned to valent vibrations of the C–O–C bond and glycosidic bridge. The band at 916 cm $^{-1}$  was due to the asymmetric telescopic vibration of glucopyranose ring, and the band at 845 cm $^{-1}$  indicated the presence of α-anomeric configuration. The bands at 916 and 845 cm $^{-1}$  also indicated that the glucopyranose units in AWP had a  $C_1$  chair conformation (Mitić et al., 2009, 2010; Shingel, 2002).

#### 3.4. NMR spectroscopy analysis

In the  $^{13}$ C NMR spectrum of AWP (Fig. 4A), six major signals among 98–65 ppm were attributed to C-1–C-6 of glucopyranose units. The chemical shift of C-6 was displaced to the low magnetic field at 66.7 ppm, which indicated that the glucopyranose was substituted at C-6 (Uzochukwu et al., 2002). It was deduced that the backbone of AWP was composed of  $\rightarrow$ 6)- $\alpha$ -D-Glcp-(1 $\rightarrow$  units. In the anomeric region of the  $^1$ H NMR spectrum (Fig. 4B), the resonance at 5.02 ppm was due to the H-1 of the (1 $\rightarrow$ 6)-linked D-glucopyranose units (Purama, Goswami, Khan, & Goyal, 2009). The weak resonance peak at 5.36 ppm indicated the existence of a little branched linkage (Seymour, 1979), and it was ascribed to H-1 of the (1 $\rightarrow$ 3,6)-linked D-glucopyranose units. The anomeric proton signals at 5.02 and 5.36 ppm were assigned to  $\alpha$ -configuration pyranose units.

The assignments for different resonances of 2D NMR experiments, including  $^1H^{-1}H$  COSY,  $^1H^{-13}C$  HMQC and  $^1H^{-13}C$  HMBC were used to further assign the chemical shifts of the main spin system.  $^1H$  NMR spin systems of the polysaccharide were assigned by  $^1H^{-1}H$  COSY (Fig. 4C). Combined with the analysis of the  $^1H^{-13}C$  HMQC (Fig. 4D), the assignment of all the signals of the major unit could be completed. The presence of strong cross peak H-1/C-6 and H-6/C-1 on  $^1H^{-13}C$  HMBC (Fig. 4E) confirmed that  $(1\rightarrow 6)-\alpha-p-Glcp$  was the main pattern of linkage. The correlation signals assignment of  $^1H^{-1}H$  COSY,  $^1H^{-13}C$  HMQC and  $^1H^{-13}C$  HMBC further confirmed the conclusion (Maina, Tenkanen, Maaheimo, Juvonen, & Virkki, 2008; Purama et al., 2009; Yan, Li, Wang, & Wu, 2010). The  $^1H$  and  $^{13}C$  NMR chemical shifts are summarized in Table 2.

**Table 1**Results from methylation analysis of AWP.

Methylated sugara	Primary mass fragments $(m/z)$	Molar ratio	Deduced linkage type
2,3,4,6-(Me) <sub>4</sub> -Man	87, 101, 117, 145, 161, 205	1.0	Manp-(1→
2,3,4-(Me) <sub>3</sub> -Glc	87, 101, 117, 129, 161, 189, 233	7.7	$\rightarrow$ 6)-Glcp-(1 $\rightarrow$
2,4-(Me) <sub>2</sub> -Glc	87, 101, 117, 129, 189, 233, 305	1.1	$\rightarrow$ 3,6)-Glcp-(1 $\rightarrow$

 $<sup>^{</sup>a}~2,3,4,6-(Me)_{4}-Man,2,3,4-(Me)_{3}-Glc~and~2,4-(Me)_{2}-Glc~correspond~to~1,5-di-O-acetyl-2,3,4,6-tetra-O-methyl-p-mannitol,~1,5,6-tri-O-acetyl-2,3,4-tri-O-methyl-p-glucitol~and~1,3,5,6-tetra-O-acetyl-2,4-di-O-methyl-p-glucitol~and~1,3,5,6-tetra-O-acetyl-2,4-di-O-methyl-p-glucitol~and~1,3,5,6-tetra-O-acetyl-2,4-di-O-methyl-p-glucitol~and~1,3,5,6-tetra-O-acetyl-2,4-di-O-methyl-p-glucitol~and~1,3,5,6-tetra-O-acetyl-2,4-di-O-methyl-p-glucitol~and~1,3,5,6-tetra-O-acetyl-2,4-di-O-methyl-p-glucitol~and~1,3,5,6-tetra-O-acetyl-2,4-di-O-methyl-p-glucitol~and~1,3,5,6-tetra-O-acetyl-2,4-di-O-methyl-p-glucitol~and~1,3,5,6-tetra-O-acetyl-2,4-di-O-methyl-p-glucitol~and~1,3,5,6-tetra-O-acetyl-2,4-di-O-methyl-p-glucitol~and~1,3,5,6-tetra-O-acetyl-2,4-di-O-methyl-p-glucitol~and~1,3,5,6-tetra-O-acetyl-2,4-di-O-methyl-p-glucitol~and~1,3,5,6-tetra-O-acetyl-2,4-di-O-methyl-p-glucitol~and~1,3,5,6-tetra-O-acetyl-2,4-di-O-acetyl$ 

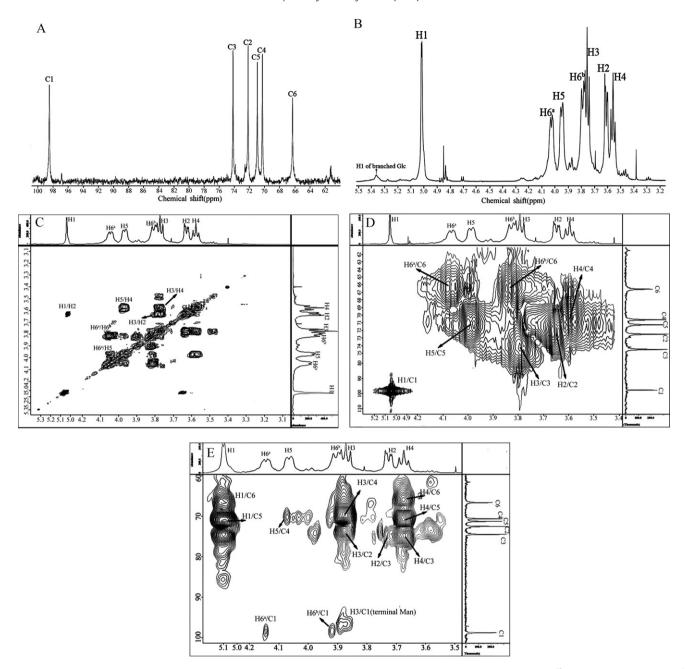


Fig. 4. NMR spectra of AWP. Spectra were performed at 23 °C on a JEOL ECP 600 MHz spectrometer using acetone as internal standard. (A)  $^{13}$ C NMR and DEPT spectra, (B)  $^{1}$ H NMR spectrum, (C)  $^{1}$ H $^{-1}$ H COSY spectrum, (D)  $^{1}$ H $^{-13}$ C HMQC spectrum, and (E)  $^{1}$ H $^{-13}$ C HMBC spectrum.

Because of the strong signals of (1 $\rightarrow$ 6)-linked  $\alpha$ -D-glucopyranose units, other signals were relatively weak and overlapped. It is hard to assign these signals clearly. The weak resonance at 97.2 ppm could be attributed to the anomeric carbon

of non-reducing terminal mannopyranose. The peak at 61.5 ppm was attributed to C-6 of the terminal mannopyranose. The correlation signals of  $^1\mathrm{H}^{-13}\mathrm{C}$  HMBC at 3.59/97.2 ppm were presumed to be the correlation of C-1 of terminal mannopyranose and H-3 of

**Table 2** <sup>1</sup>H and <sup>13</sup>C chemical shifts for AWP.

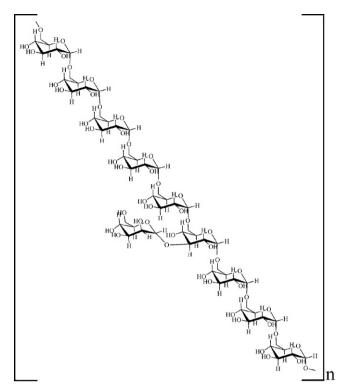
Sugar residues <sup>d</sup>	<sup>1</sup> H/ <sup>13</sup> C chemical	<sup>1</sup> H/ <sup>13</sup> C chemical shifts (ppm) <sup>c</sup>						
	H1/C1	H2/C2	H3/C3	H4/C4	H5/C5	H6 <sup>a</sup> , H6 <sup>b</sup> /C6		
$\rightarrow$ 6)- $\alpha$ -D-Glcp-(1 $\rightarrow$ $^3J_{H,H}$ (Hz)	5.02/98.80 3.0	3.61/72.48 9.75	3.75/74.82 9.36	3.54/70.59 9.36	3.95/71.26 8.28	4.03,3.82/66.0 6.9		

<sup>&</sup>lt;sup>a</sup> Equatorial bond.

b Axial bond.

<sup>&</sup>lt;sup>c</sup> The spectra were recorded using a JEOL JNM-ECP 600 MHz spectrometer. Chemical shifts are referenced to internal acetone at 2.225 ppm for <sup>1</sup>H and 31.07 ppm for <sup>13</sup>C.

d Glcp: glucopyranose.



**Fig. 5.** Proposed structure of the extracellular polysaccharide AWP produced by *A. versicolor* ( $n \approx 300$ ).

the  $(1\rightarrow6)$ -linked D-glucopyranose. Thus, it was supposed that the mannopyranose was linked to C-3 position of the  $(1\rightarrow6)$ -linked D-glucopyranose backbone.

All the results demonstrated that the backbone of AWP contained a large number of consecutive  $(1\rightarrow 6)$ -linked glucopyranose units with minor amount of branches. The branching was composed of single non-reducing-end mannopyranose units which were linked to C-3 position of  $(1\rightarrow 6)$ -linked p-glucopyranose units. On average, there was one branching point for every 9 units on the backbone. The proposed structure of AWP was presented in Fig. 5.

The extracellular polysaccharide AWP from coral-associated fungus A. versicolor had different structural characteristics from other extracellular polysaccharides from Aspergillus species (Gómez-Miranda et al., 2003; Tischer, Gorin, de Souza, & Barreto-Bergter, 2002). The polysaccharides from mycelia of A. versicolor were characterized to be phosphonogalactomannans, which contained a mannan core, substituted at O-2 and O-6 with side chains of (1 $\rightarrow$ 5)-linked  $\beta$ -galactofuranosyl units, phosphodiester groups were present principally as bridges between C-1 of units of  $\alpha$ -mannopyranose and C-6 of those of 5-0-substituted β-galactofuranose (Tischer et al., 2002). The polysaccharide from Aspergillus wentii corresponded to the following repeating unit:  $[\rightarrow 3)$ - $\beta$ -D-Galf- $(1\rightarrow 5)$ - $\beta$ -D-Galf- $(1\rightarrow ]_n\rightarrow$ mannan core. The mannan core is constituted by a  $(1\rightarrow 6)$ - $\alpha$ -mannan backbone, substituted at positions 2 by chains from 1 to 7 residues of  $(1\rightarrow 2)$ linked α-mannopyranoses (Gómez-Miranda et al., 2003). In addition, it is noted that the structure of AWP was different from that of the extracellular polysaccharide AVP isolated from the fermented broth of A. versicolor (Chen et al., 2012). The backbone of AVP was composed of  $(1\rightarrow 6)$ -linked  $\alpha$ -D-glucopyranose and  $(1\rightarrow 2)$ -linked  $\alpha$ -D-mannopyranose units, and the mannopyranose residues in the backbone were substituted mainly at C-6 by the side chain of  $(1\rightarrow 2)$ -linked  $\alpha$ -D-mannopyranose trisaccharides units. Symbiotic microorganisms have close relationship with the hosts, and may be involved in organism biosynthesis, metabolism and other

life activities of large hosts (Croft et al., 2005; Dobretsov & Qian, 2004). The present result suggested that symbiotic microorganisms from marine organisms could be a potential source of extracellular polysaccharides with unique structures, and be worth being further investigated.

#### 4. Conclusion

AWP produced by the marine fungus *A. versicolor* from the coral *Cladiella* sp. is a novel extracellular polysaccharide with different structural characteristics from extracellular polysaccharides excreted by other marine microorganisms. The main chain of AWP consists of  $(1\rightarrow 6)$ -linked  $\alpha$ -D-glucopyranose residues with minor amounts of substitutions at C-3 by  $(1\rightarrow)$ -linked  $\alpha$ -D-mannopyranose. Further studies are underway to characterize its physical and chemical properties.

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