Structural Characterization of Anti-Complementary Polysaccharides from the Leaves of *Artemisia princeps*

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**Abstract:** Structural characterizations of the anti-complementary acidic heteroglycans, AAF IIb-2 and IIb-3, obtained from the leaves of *Artemisia princeps* PAMP have been studied. AAF IIb-2 consists of rhamnose, xylose, arabinose, galactose, glucose and uronic acids (glucuronic acid and galacturonic acid) in the molar ratio of 13.0:3.0:57.9, and AAF IIb-3 consists of the same sugars in the ratio of 19.7:2.6:24.7:2.6:46.5. Methylation analysis including carboxyl-reduction and also selective enzymolysis using exo-α-L-arabinofuranosidase suggested that AAF IIb-3 has a main chain consisting of (1→4)-linked galacturonic acid and (1→2)-linked rhamnose mostly substituted at the O-4 position. AAF IIb-3 also contained arabinono-3,6-galactan moiety and most of the arabinose was present as an α-L-furanosyl residue in the non-reducing terminals and highly branched side chains which mostly attached to the O-3 position of (1→6)-linked galactopyranosyl residue. The basic structure of AAF IIb-2 is similar to that of AAF IIb-3, but IIb-3 has a higher arabinogalactan content than IIb-2.

**List of Abbreviations**

- C₅H₅N: pyridine
- THF: tetrahydrofuran
- TFA: trifluoroacetic acid
- ¹H-NMR: proton magnetic resonance
- NaBH₄: sodium borohydride
- NaBD₄: sodium borodeuteride
- Araf: arabinofuranose
- Gal: galactopyranose
- Ara: arabinopyranose
- Xyl: xylopyranose
- Rha: rhamnopyranose
- GalA: galacturonic acid

**Materials and Methods**

**Materials**

The leaves of *Artemisia princeps* Pamp (Japanese name = Gaiyo) were purchased from Uchida Wakanayaku Co. Ltd., Tokyo, Japan. α-L-Arabinofuranosidase from *Rhodotorula flava* (2) was a kind gift from Dr. Naoto Shibuya (National Food Research Institute, Ministry of Agriculture, Forestry and Fisheries, Japan). Bio-gel P-2 (200—400 mesh) was obtained from Bio-Rad. Sephadex LH-20 was purchased from Pharmacia and Sep-Pak C18 cartridge from Waters Associates Inc.

**Purification of anti-complementary polysaccharides, AAF IIb-2 and IIb-3**

The crude polysaccharide from the leaves of *A. princeps* Pamp was prepared by hot-water extraction and dialysis (1). The crude polysaccharide was purified by ion exchange chromatography on DEAE-Sepharose, affinity chromatography on *Ricinus communis*-agglutinin conjugated Sepharose and gel filtrations on Sephadex G-100 and Sepharose CL-4B as reported previously (1).

**Methylation analysis of polysaccharides**

Before per-O-methylation, AAF IIb-2 and AAF IIb-3 were passed through the column of AG 50W-X8 resin (H⁺ form) in order convert all of the carboxyl groups into protonated form because this treatment permits more complete O-methylation of uronic acid containing polysaccharides (9). Then AAF IIb-2 and IIIb-3 (2 mg) were methylated by the Hakomori procedure (10) and the completeness of the methylation was checked by using tricarboxylic acid (11). The per-O-methylated polysaccharides were purified by gel filtration on Sephadex LH-20 (12) or reversed-phase chromatography on Sep-Pak C18 cartridge (9). Per-O-methylated polysaccharides were eluted from the Sep-Pak C18 cartridge with 100% CH₃CN, followed by 100% EtOH as described by Waeghe et al. (9).

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The fully methylated polysaccharides were reduced with NaBD₄ in 27:73 (v/v) 95% EtOH-THF (13). Methyl-esterified hexosyluronic acid residues reduced by this procedure were converted into the corresponding 6,6-dideuterioluminated hexosyl residue (9). Per-O-methylated and reduced polysaccharides were hydrolyzed with 2 M TFA at 121°C for 1 h. The hydrolyzates were reduced to the corresponding partially O-methylated alditols with NaBH₄ in 1 M ammonia in 95% EtOH. For the detection of uronic acid residues, the hydrolyzates were subjected to carboxyl reductase digestion of polysaccharides (16). The enzyme digested polysaccharide (yield, 75.4%) was recovered in the void volume by gel filtration on a column (2.0 x 52 cm) of Bio-gel P-2 at 55°C or in the non-dialyzable portion reported previously (16). The enzyme digested polysaccharide (yield, 75.4%) was recovered in the void volume by gel filtration on a column with splitless injection. The flow rate of the carrier gas, helium, was 0.9 ml/min, The gas chromatograph was programmed at 150°C for 1 min, followed by a rate of 2°C/min to 210°C. All GLC flame ionization responses to partially methylated alditol acetates were corrected to mol masses of derivatives.

Results

AAF IIb-2 and IIb-3 were composed of rhamnose, xylose, arabinose, galactose, glucose, galacturonic acid, and glucuronic acid in the molar ratios of 7.6:7.6:13.0:10.9:3.0:51.2:6.7 and 3.9:2.6:24.7:19.7:2.6:15.0:31.5, respectively. AAF IIb-2 and IIb-3 showed positive specific rotations ([α]D: +114.3° and +67.0°). The 1H-NMR spectrum of major anti-complementary polysaccharide, AAF IIb-3, showed signals due to the methyl protons of the rhamnose residue at δ = 1.25, O-acetyl protons at δ = 2.0 and 2.13, O-methyl protons as carboxylic acid methyl esters at δ = 3.80, and anomeric protons due to β- and α-configurations at δ = 4.5—4.8 and 4.9—5.3. AAF IIb-2 and IIb-3 were methylated by the methylsulphinyl-carbanion and methyl iodide in dimethyl sulfoxide, followed by the reduction with NaBD₄. The fully methylated and reduced polysaccharides were hydrolyzed with acid, and the products were analyzed as the alditol acetates by GLC and GLC-MS. The results suggest that both polysaccharides contained arabinofuranosyl residues as non-reducing end groups, (1→5)-linked arabinofuranosyl or (1→4)-linked arabinopyranosyl residues and (1→3,5)-linked arabinofuranosyl residues, and galactopyranosyl residues as non-reducing end groups and (1→3)-, (1→4)-, (1→6)-, and (1→3,6)-linked galactopyranosyl residues, xylopyranosyl residues as non-reducing end groups, and (1→2)- and (1→2,4)-linked rhamnopyranosyl residues in the chain (Table I). The identification of 2,3-di-O-methyl-1, 4, 5, 6-tetra-O-acetylglactitol labelled with deuterium at the O-6 position in the methylated and reduced polysaccharides confirmed that the compound was derived from the (1→4)-linked galacturonic acid residue in the original polysaccharide. To confirm the structure of uronic acid moiety of AAF IIb-3, carboxyl reduced polysaccharide was also subjected to methylation analysis. The presence of glucuronic acid as non-reducing end groups and (1→4)-linked galacturonic acid are suggested by the increase of 2, 3, 4, 6-tetra-O-methylglucose and 2, 3, 6-tri-O-methylgalactose in the carboxyl-reduced AAF IIb-3 (Table II). The yield of the reduction product, 2,3-di-O-methylgalactose, derived from fully methylated and reduced AAF IIb-3 was very low value in comparison with expected value based on the uronic acid content of the original polysaccharide. It was

Table I. Methylation analysis of AAF IIb-2 and IIb-3.

<table>
<thead>
<tr>
<th>Methylated alditol acetate derivatives</th>
<th>T°</th>
<th>Mol %</th>
<th>Major mass spectral fragments (m/z)</th>
<th>Linkage</th>
</tr>
</thead>
<tbody>
<tr>
<td>1,4-Ac2, 2, 3, 5-Me2-arabinitol</td>
<td>100</td>
<td>11.8</td>
<td>43, 45, 71, 87, 101, 117, 129, 161</td>
<td>Ara1→2→</td>
</tr>
<tr>
<td>1,5-Ac2, 2, 3, 4-Me2-xylitol</td>
<td>95</td>
<td>4.0</td>
<td>43, 71, 87, 101, 117, 161</td>
<td>Xyl→2→</td>
</tr>
<tr>
<td>1,2, 5-Ac3, 3, 4-Me2-xylitol</td>
<td>100</td>
<td>8.3</td>
<td>43, 87, 99, 129, 131, 189</td>
<td>Xyl→2→</td>
</tr>
<tr>
<td>1,4, 5-Ac2, 2, 3-Me2-arabinitol</td>
<td>2.1</td>
<td>13.4</td>
<td>43, 87, 101, 117, 129, 189</td>
<td>Ara1→2→</td>
</tr>
<tr>
<td>1,5-Ac2, 2, 3, 4-Me2-galactitol</td>
<td>4.9</td>
<td>7.9</td>
<td>43, 45, 71, 87, 101, 117, 129, 145</td>
<td>Gal1→</td>
</tr>
<tr>
<td>1,2, 4, 5-Ac3, 3-Me2-xylitol</td>
<td>1.2</td>
<td>10.3</td>
<td>43, 87, 101, 129, 143, 189, 207</td>
<td>Xyl→2→</td>
</tr>
<tr>
<td>1,3, 5-Ac2, 2, 3-Me2-arabinitol</td>
<td>1.7</td>
<td>15.7</td>
<td>45, 85, 117, 201, 261</td>
<td>Ara1→2→</td>
</tr>
<tr>
<td>1,3, 5-Ac2, 2, 4, 6-Me2-galactitol</td>
<td>1.9</td>
<td>3.6</td>
<td>45, 45, 87, 101, 113, 117, 129</td>
<td>Gal1→</td>
</tr>
<tr>
<td>1,4, 5-Ac2, 3, 6-Me2-galactitol</td>
<td>2.1</td>
<td>14.3</td>
<td>43, 45, 87, 101, 117, 129, 161, 233</td>
<td>Gal1→</td>
</tr>
<tr>
<td>1,5-Ac2, 3, 4-Me2-arabinitol</td>
<td>2.8</td>
<td>4.7</td>
<td>43, 87, 99, 101, 117, 129, 189</td>
<td>Gal1→</td>
</tr>
<tr>
<td>1,4, 5-Ac2, 3, 4-Me2-galactitol</td>
<td>4.5</td>
<td>4.2</td>
<td>43, 85, 87, 99, 101, 117, 127, 261</td>
<td>Gal1→</td>
</tr>
<tr>
<td>1,3, 5, 6-Ac2, 2, 4-Me2-galactitol</td>
<td>4.9</td>
<td>1.8</td>
<td>43, 87, 117, 129, 139, 189, 233</td>
<td>Gal1→</td>
</tr>
</tbody>
</table>

a Retention times of the corresponding alditol acetates relative to that of 1,5-Ac2, 3, 4, 6-Me2-glucitol on an OV-225 column.
b Calculated from peak areas and mol masses of derivatives.
c Assigned in the literature (14).
suggested that the low recovery of the reduction product may be caused by the low efficiency of the reduction step and the loss of uronic acid residues during Hakomori-methylation by β-elimination. To obtain more detailed information about the structure of the side chains of AAF IIb-3, AAF IIb-3 was digested with exo-α-L-arabinofuranosidase from R. flava. Then the reaction products were fractionated on a column of Bio-gel P-2 (data not shown), and two carbohydrate fractions that were eluted in the void and retained volumes were obtained. Arabinose was the only sugar detected in the retained volume by TLC and GLC. The material recovered in the void volume consisted of rhamnose, arabinose, xylose, galactose, and glucose in a molar ratio of 0.35:0.28:0.16:1:0.1 as the neutral sugars. These results lead to the conclusion that most of arabinosyl residues in AAF IIb-3 were linked in an α-configuration in the furanose form. The ¹H-NMR spectrum of exo-α-L-arabinofuranosidase digested AAF IIb-3 showed that the signal at 5 = 2.0 had decreased in intensity, suggesting that a part of O-acetyl group may be linked to a part of arabinofuranosyl residues. The results of methylation analysis of the arabinofuranosidase digested AAF IIb-3 showed that the molar ratio of the terminal, (1→5), and (1→3,5)-linked arabinofuranosyl residues, and (1→3) and (1→3,6)-linked galactosyl residues had decreased significantly, and that the molar ratio of the terminal and (1→6)-linked galactosyl residues had increased compared to AAF II-3 (Table II). When AAF IIb-2 was digested with the same enzyme, similar results with AAF IIb-3 were obtained, but further studies of AAF IIb-2 were not possible because of the small quantity available.

**Discussion**

The major anti-complementary polysaccharide, AAF IIb-3, was mainly composed of arabinose and galactose while the minor one, AAF IIb-2, was mainly composed of rhamnose, xylose, arabinose, and galactose (1). Previously, a significant amount of galacturonic acid was detected in these heteroglycans by TLC of the acid hydrolyzates (1), but in the present paper, galuronic acid was also identified by GLC analysis. AAF IIb-2 contained more substantial amounts of (1→4)-linked galacturonic acid and (1→2)-linked rhamnopyranosyl residues than AAF IIb-3. These results indicate that AAF IIb-2 and IIb-3 have a rhamnogalacturonan main chain as shown in pectic polysaccharides similar to those isolated from cotyledon meals and hulls of Glycine max (18, 19), leaves and stems of Medicago sativa (20), suspension cultured sycamore cell walls (21), rice endosperm cell walls (22), the fruits of Zizyphus jujuba Mill var. inermis Reh (23) and suspension-culture cells and leaves of Nicotiana tabacum (24, 25). Darvill et al. classified that there are two types of rhamnogalacturonan I and II (26). AAF IIb-2 and IIb-3 appeared to be rhamnogalacturonan I type polysaccharides because apiose, 2-methylfucose, and 2-methylxylose were not detected in these acidic heteroglycans. The results of methylation analysis of AAF IIb-3 before and after α-L-arabinofuranosidase digestion, suggest that the highly branched arabinogalactan moiety is a (1→3)- and (1→6)-linked galactan, possessing attached highly branched arabinofuranosyl side chains at O-3 of some of the (1→6)-linked galactosyl residues in the main chain or the interior chains of galactose residues mutually joined by (1→3)- and (1→6)-linkages. This structure moiety was very similar to the anti-complementary arabinono-3,6-galactans from the roots of

**Table II. Methylation analysis of AAF IIb-3, carboxyl-reduced IIb-3 and IIb-3 α-L-arabinofuranosidase (α-L-Arafase) digest**

<table>
<thead>
<tr>
<th>Methylated alditol acetate derivatives</th>
<th>Mol %</th>
<th>Reduced IIb-3</th>
<th>α-L-Arafase treated</th>
<th>Linkage</th>
</tr>
</thead>
<tbody>
<tr>
<td>1,4-Ac4-2,3,5-Me4-arabinitol</td>
<td>19.2</td>
<td>8.7</td>
<td>6.7(5.1)↓</td>
<td>Ara6→</td>
</tr>
<tr>
<td>1,5-Ac5-2,3,4-Me4-xylitol</td>
<td>1.6</td>
<td>6.2</td>
<td>2.1(1.6)↓</td>
<td>Xyl1↓</td>
</tr>
<tr>
<td>1,2,5-Ac5-3,4,6-Me4-rhamnitol</td>
<td>4.1</td>
<td>3.0</td>
<td>6.9(5.2)↓</td>
<td>→Rha1↓</td>
</tr>
<tr>
<td>1,5-Ac5-2,3,4,6-Me4-glucitol</td>
<td>6.2↑</td>
<td></td>
<td>(~)</td>
<td></td>
</tr>
<tr>
<td>1,4,5-Ac5-2,3,5-Me5-arabinitol</td>
<td>16.0</td>
<td>7.0</td>
<td>10.2(7.7)↓</td>
<td></td>
</tr>
<tr>
<td>1,5-Ac5-3,4,6-Me4-galactitol</td>
<td>15.1</td>
<td>12.9</td>
<td>25.6(19.5)↑</td>
<td></td>
</tr>
<tr>
<td>1,2,4,5-Ac5-3-Me4-rhamnitol</td>
<td>4.0</td>
<td>1.1</td>
<td>6.4(4.8)↓</td>
<td></td>
</tr>
<tr>
<td>1,3,4,5-Ac5-2-Me4-arabinitol</td>
<td>14.6</td>
<td>4.2</td>
<td>3.6(2.7)↓</td>
<td></td>
</tr>
<tr>
<td>1,3,5-Ac5-2,3,4,6-Me4-galactitol</td>
<td>7.4</td>
<td>3.8</td>
<td>9.2(7.0)↓</td>
<td></td>
</tr>
<tr>
<td>1,4,5-Ac5-3,5-Me5-galactitol</td>
<td>8.2</td>
<td>4.27↑</td>
<td>11.2(8.5)↑</td>
<td></td>
</tr>
<tr>
<td>1,5-Ac5-2,6,3,4-Me4-galactitol</td>
<td>4.6</td>
<td>2.3</td>
<td>12.6(9.6)↑</td>
<td></td>
</tr>
<tr>
<td>1,3,5-6-Ac5-2,4-Me4-galactitol</td>
<td>5.2</td>
<td>1.9</td>
<td>5.4(3.6)↓</td>
<td></td>
</tr>
</tbody>
</table>

* Methylated different IIb-3 preparation from Table I.
* Molar ratio in parenthesis calculated relative to the molar ratio of 1,4-Ac4-2,3,5-Me4-arabinitol in IIb-3.
↓ decreased, ↑ increased.
Angelica acutiloba Kitagawa (AR-arabinogalactan IIa and IIb-1) (16, 27, 28) and the arabinogalactans from Zea shoots (29).

When AAF IIb-2 and IIb-3 were digested with exo-α-L-arabinofuranosidase, a significant amount of the arabinosyl residue still remained in the enzyme resistant polysaccharide. The results of methylation analysis of the enzyme digested AAF IIb-3 indicate that the terminal and (1→5)-linked arabinofuranosyl residue had been retained significantly compared to AAF IIb-3. These results suggest that arabinogalactan moiety of AAF IIb-3 contained a highly branched araban moiety, possessing attached arabinofuranosyl side chains at O-3 of some of the (1→5)-linked arabinofuranosyl residues (Fig. 1), and the enzyme may be cleaved only at outer side chains of the branching point. The structure of this branched arabinan moiety was also similar to that of AR-arabinogalactan IIb-1 (28) from A. acutiloba. The rhamnogalacturonan moieties in AAF IIb-2 and IIb-3 may be present in the inner core region because when AAF IIb-3 was subjected to partial acid hydrolysis, galacturonic acid, rhamnose, and galactose were detected as major component sugars in the acid resistant polysaccharide (unpublished results). This result also suggested that the arabinogalactan side chain may be attached to the rhamnogalacturonan backbone through position 4 of (1→2)-linked rhamnose residue in AAF IIb-2 and IIb-3 (Fig. 1). AAF IIb-2 was shown to have more potent anti-complementary activity than IIb-3 (1), but both contained similar glycosidic linkages with different molar ratios.

Although rhamnogalacturonan moieties and non-reducing end groups of xylosyl and glucuronic acid residues were more predominant than the arabinogalactan moiety in AAF IIb-2, the elucidation of the structural difference between AAF IIb-2 and IIb-3 in association with the activity must await further detailed analysis.

Acknowledgements

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References