

Enzymatic modification of pectic polysaccharides obtained from sugar beet pulp

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Abstract

Rhamnogalacturonans and arabinans, purified from an autoclave extract of sugar beet pulp, as well as an acid extracted beet pectin (ABP) were treated with enzymes in order to modify their physico-chemical properties. The enzymes used were arabinofuranosidase B (AF), endo-arabinanase plus arabinofuranosidase (EA + AF), rhamnogalacturonase plus rhamnogalacturonan acetyl esterase (RGase + RGAE), and polygalacturonase plus pectin methyl esterase (PG + PE). During the enzyme treatments the intrinsic viscosity ($[\eta]_w$), apparent molecular weight (M_w), and radius of gyration (Rg_w) of the polysaccharides were monitored.

We found in most cases that treatment with glycanases decreased the $[\eta]_w$ of the polysaccharides investigated. However, results showed that degradation of the rhamnogalacturonan backbone of a beet pectin obtained by acid extraction had little influence on the $[\eta]_w$ which led to the conclusion that the rhamnogalacturonans present in this type of pectin are mainly located at the extremities of the pectin molecules. Some evidence was found that the rhamnogalacturonans obtained by autoclaving are linked through the arabinose side-chains, probably by diferulic acid cross-links. Removal of the side-chains of the arabinans present in the rhamnogalacturonans of ABPs with enzymes also showed that the arabinan side-chains contribute relatively little to the $[\eta]_w$.

In our experiments we found that modification with glycanases can be used to change the structural characteristics of pectic polysaccharides without a significant loss of viscosity. In this way glycanases can be used for the removal of structural elements of pectic polysaccharides, which limit their physico-chemical properties. © 2002 Elsevier Science Ltd. All rights reserved.

Keywords: Sugar beet pulp; Pectins; Rhamnogalacturonans; Enzymatic modification

1. Introduction

The structural characteristics of pectic polysaccharides from sugar beet have been the subject of many studies (Colquhoun, Ralet, Thibault, Faulds & Williamson, 1994; Guillou & Thibault, 1988, 1989a, 1989b; Oosterveld, Beldman, Schols & Voragen, 1996, 2000a; Ralet, Thibault, Faulds & Williamson, 1994; Rombouts & Thibault, 1986; Thibault, Guillou & Rombouts, 1991; Thibault, Renard, Axelos, Roger & Crépeau, 1988). Also several studies have been attended towards the physico-chemical properties of these polysaccharides (Faulds & Williamson, 1990; Fishman, Gillespie, Sondey & El-Atawy, 1991; Hwang & Kokini, 1991; Matthew, Howson, Keenan & Bolton, 1990; Thibault, 1986). However, relatively little is known about the relationship between their structure and physico-chemical properties.

Due to their specificity, enzymes are appropriate tools to modify the structure of pectins. In this way information can be obtained about the relation between chemical structure and physical characteristics. The knowledge acquired can be used to produce tailor made polysaccharides with specific (physico-chemical) properties.

Several studies have been carried out, addressing the enzymatic modification of beet pectins. Some of these studies focused on the enzymatic removal of acetyl groups from acid extracted sugar beet pectins. As a result the gelling properties of these pectins improved (Faulds and Williamson, 1990; Matthew et al., 1990; Oosterveld, Beldman, Searle-van Leeuwen & Voragen, 2000b; Pippen, McCready & Owens, 1950). McCleary et al. showed that arabinans extracted from sugar beet pulp could be used e.g. as fat replacer after debranching with the enzyme arabinofuranosidase (Cooper, McCleary, Morris, Richardson, Marrs & Hart, 1992; McCleary, Cooper & Williams, 1989). An improvement of the gel formation of pectins from sugar beet pulp through oxidative cross-linking with ammonium persulfate after pretreatment with the

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same enzyme was found by Guillon and Thibault (1987, 1990).

The current study deals with the elucidation of the chemical structure and the modification of the physico-chemical properties of three types of pectic polysaccharides obtained from sugar beet pulp by using specific enzymes. The polysaccharides used were rhamnogalacturonans and arabinans obtained from beet pulp by autoclaving (Oosterveld et al., 1996, 2000a), and a commercially available acid extracted pectin, mainly consisting of homogalacturonans.

2. Experimental

2.1. Materials

Wet beet pulp (8.9 % dry weight) was obtained from CSM Suiker bv (Breda, the Netherlands). Pectins were extracted by autoclave treatment as described previously (Oosterveld et al., 1996). For the experiments described in this manuscript the extract named Autoclave 2 was used. The acid extracted pectin obtained from sugar beet pulp (ABP) was a gift from the Copenhagen Pectin Factory Ltd.

2.2. Analytical methods

The uronic acid contents of the extracts were determined by the automated *m*-hydroxy biphenyl assay (Thibault, 1979). The neutral sugar composition was determined after hydrolysis with 2 M trifluoroacetic acid (1 h, 121°C) as described previously (Oosterveld et al., 1996). The degrees of methylation (DM) and acetylation (DA) were determined as described previously (Oosterveld et al., 1996). Feruloyl groups were determined spectrophotometrically at 375 nm in freshly prepared pectin solutions adjusted to pH 10 with a 0.1 M NaOH solution, using a molar extinction coefficient of 31,600 (Rombouts & Thibault, 1986).

2.3. Chromatography

High-performance size-exclusion chromatography (HPSEC) was performed on three Bio-Gel TSK columns in series (60XL-40XL-30XL) as described (Oosterveld et al., 2000a) using a combined RI detector and viscometer (Viscotek, model 250), a Right Angle Laser Light-Scattering detector (RALLS, Viscotek, LD 600) and a UV detector (Kratos, Spectroflow 773). Apparent molecular weights, intrinsic viscosities and radii of gyration were calculated using the light-scattering module of the Trisec software (Viscotek).

Preparative size-exclusion chromatography was performed as described previously (Oosterveld et al., 2000a).

Preparative anion-exchange chromatography was performed on a column (15 × 10 cm) of Source Q

(Pharmacia) using a Biopilot system (Pharmacia). The sample (~0.4 g) was applied onto the column at a flow rate of 15 ml/min (0.005 M NaOAc, pH 5.0). After 1575 ml the flow rate was increased to 80 ml/min and a linear gradient to 2 M NaOAc (pH 5.0) was applied for 7200 ml to elute the polysaccharides. Subsequently, the elution was proceeded using elution with 1200 ml 2 M NaOAc. The columns were washed with 0.5 M NaOH, followed by regeneration with 2 M NaOAc (pH 5.0) and equilibration with 0.005 M NaOAc (pH 5.0). Fractions (90 or 356 ml) were assayed for total neutral sugar (Tollier & Robin, 1979) and uronic acid (Thibault, 1979) content using arabinose and galacturonic acid as standards. A correction was made for the response of uronic acids in the neutral sugar test. Pooled fractions were dialyzed against distilled water (estimated molecular weight cut-off for polysaccharides ~2000 Da) and freeze-dried.

2.4. Enzymatic modification

The pools obtained by DEAE anion-exchange chromatography and preparative size-exclusion chromatography were treated with the following purified enzymes: arabinofuranosidase B (AF) (Rombouts, Voragen, Searle-van Leeuwen, Gereads, Schols & Pilnik, 1988), AF plus endo-arabinanase (EA) (Rombouts et al., 1988), endo-polygalacturonase (PG) (Pasculli, Gereads, Voragen & Pilnik, 1991) combined with pectin methyl esterase (PE) (Baron, Rombouts, Drilleau & Pilnik, 1980), and rhamnogalacturonase (RGase) (Schols, Gereads, Searle-van Leeuwen, Kormelink & Voragen, 1990) in combination with rhamnogalacturonan acetyl esterase (RGAE) (Searle-van Leeuwen, van de Broek, Schols, Beldman & Voragen, 1992).

All pools were dissolved in 0.04 M sodium acetate buffer containing 0.01% of NaN₃ (pH 5.0) to a final concentration of 5 mg/ml. Enzymes were added to a final concentration of 1 µg of protein/ml. Incubations were carried out at 20°C for 42 h. The digests were analyzed at several stages of the incubation by HPSEC.

3. Results and discussion

3.1. Composition of the substrates

3.1.1. Pectins from an autoclave extract

Two rhamnogalacturonan fractions and an arabinan fraction were purified from an autoclave extract of sugar beet pulp by preparative size-exclusion chromatography and characterized as described previously (Table 1) (Oosterveld et al., 2000a). Additionally the degree of feruloylation (DF) of these substrates was determined, as well as the apparent molecular weight (M_w), intrinsic viscosity ([η]_w), and radius of gyration (Rg_w) (Table 2).

Three fractions are distinguished (see also Table 1). Pool I mainly contains rhamnogalacturonans with a very high M_w

Table 1

Sugar compositions (mol%) of the pools of the extracts obtained by autoclaving and of the acid extract

	ABP ^a	Autoclave				
		Pool I ^b	Pool II ^b	Pool III ^b	Pool IIIA	Pool IIIB
Rha ^c	5.4	6.8	5.3	1.5	0.1	3.8
Ara ^c	7.1	58.8	51.5	36.9	91.0	19.3
Xyl ^c	1.2	0.8	0.5	0.4	0.1	1.0
Man ^c	2.0	0.0	0.7	1.9	2.2	3.3
Gal ^c	11.2	13.2	9.9	3.2	0.9	7.7
Glc ^c	0.4	0.6	0.4	1.3	2.5	2.9
UA ^c	72.9	19.9	31.8	54.9	3.2	62.0
DA ^d	15.3	51.1	56.3	25.8	nd	25.6
DM ^d	57.7	28.8	33.4	56.0	nd	60.5
DF ^e	1.2	1.7	1.4	0.9	1.0	0.8

^a Molar sugar composition from (Oosterveld et al., 2000b).

^b Molar sugar composition from (Oosterveld et al., 2000a). nd not determined.

^c Expressed as mol%.

^d Expressed as moles acetyl or methyl per 100 moles of galacturonic acid.

^e Expressed as moles ferulic acid per 100 moles of galactose + arabinose.

(1020 kDa, Table 2). The pool is highly acetylated and feruloylated. The DM is relatively low. The $[\eta]_w$ of this pool was relatively low for a molecule with such a high M_w , what can be explained by the high degree of branching.

In pool II rhamnogalacturonans with an apparent molecular weight of 225 kDa and with a relatively low intrinsic viscosity are predominantly present. Except for the galacturonic acid content, its composition is similar to that of pool I. The DF is somewhat lower than that of pool I.

Table 2

Apparent molecular weight (M_w , kDa), intrinsic viscosity ($[\eta]_w$, dl/g), and radius of gyration (Rg_w , nm) of ABP and pools obtained from the autoclave extract, before and after treatment with various enzyme combinations for 42 h (nd not determined)

	ABP	Autoclave		
		Pool I	Pool II	Pool IIIA
Blank	M_w	271	1020	225
	$[\eta]_w$	4.54	0.93	0.66
	Rg_w	33.1	30.8	15.6
Rgase + RGAE	M_w	112	157	60
	$[\eta]_w$	4.01	0.52	0.45
	Rg_w	22.7	12.7	9.2
AF	M_w	nd	712	131
	$[\eta]_w$	nd	0.99	0.67
	Rg_w	nd	28.4	14.2
EA + AF	M_w	207	99	126
	$[\eta]_w$	4.24	0.79	0.42
	Rg_w	29.2	14.0	12.2
PG + PE	M_w	5	735	156
	$[\eta]_w$	0.12	0.83	0.46
	Rg_w	2.5	26.4	12.3

Pool III is a mixture of homogalacturonans and arabinans with low M_w as determined by light-scattering (~20 kDa). This fraction is highly methylated, while the DA and DF are low. The arabinans and homogalacturonans present in pool III were separated by Source Q anion-exchange chromatography (Fig. 1). Pool IIIA consisted almost exclusively of arabinans, with a high ferulic acid content. The composition of this arabinan population was similar to the neutral fraction of the autoclave extract as obtained after DEAE anion-chromatography (Oosterveld et al., 2000a). Most of the galacturonic acid eluted in pool IIIB, together with 38 mol% of neutral sugars. This fraction had a high DM, whereas the DA was lower than in pool I and pool II.

3.1.2. Acid extracted sugar beet pectin (ABP)

The ABP consist of 72.9 mol% of galacturonic acid and galactose was the predominant neutral sugar (Oosterveld, et al., 2000b). Substantial amounts of both methyl esters and acetyl groups were present. The DF was 1.2, but since this is calculated based on the amount of galactose plus arabinose, the absolute amount of ferulic acid was relatively low. The composition of ABP was comparable to other ABPs described in the literature (Michel, Thibault, Mercier, Heitz & Pouillaude, 1985; Phatak, Chang & Brown, 1988).

3.2. Effect of enzymatic modification

Pool I, II, and IIIA of the autoclave extract, as well as ABP, were subjected to the following combinations of enzymes: AF; Rgase + RGAE, AF + EA, and PG + PE. During enzymatic treatment for 42 h, changes in the apparent molecular weight (M_w), intrinsic viscosity ($[\eta]_w$), and radius of gyration (Rg_w) were monitored in time, as shown in Figs. 2–5. The values of these parameters after 42 h are shown in Table 2. The homogalacturonan pool IIIB was not subjected to enzymatic modification, because of its low M_w , which indicates that this pool was severely degraded during autoclave treatment. ABP was used as an example of a beet pectin with relatively intact homogalacturonan regions.

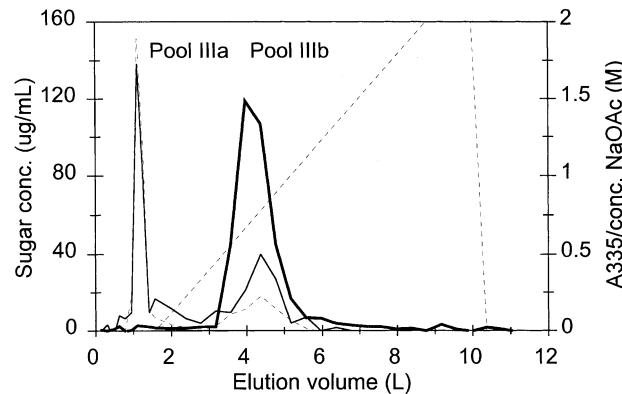


Fig. 1. Preparative anion-exchange chromatography (Source Q) of pool III obtained by S 500 size-exclusion chromatography. Thick line: uronic acid; dashed line: neutral sugars; thin line: A_{335} ; dotted line: concentration NaOAc.

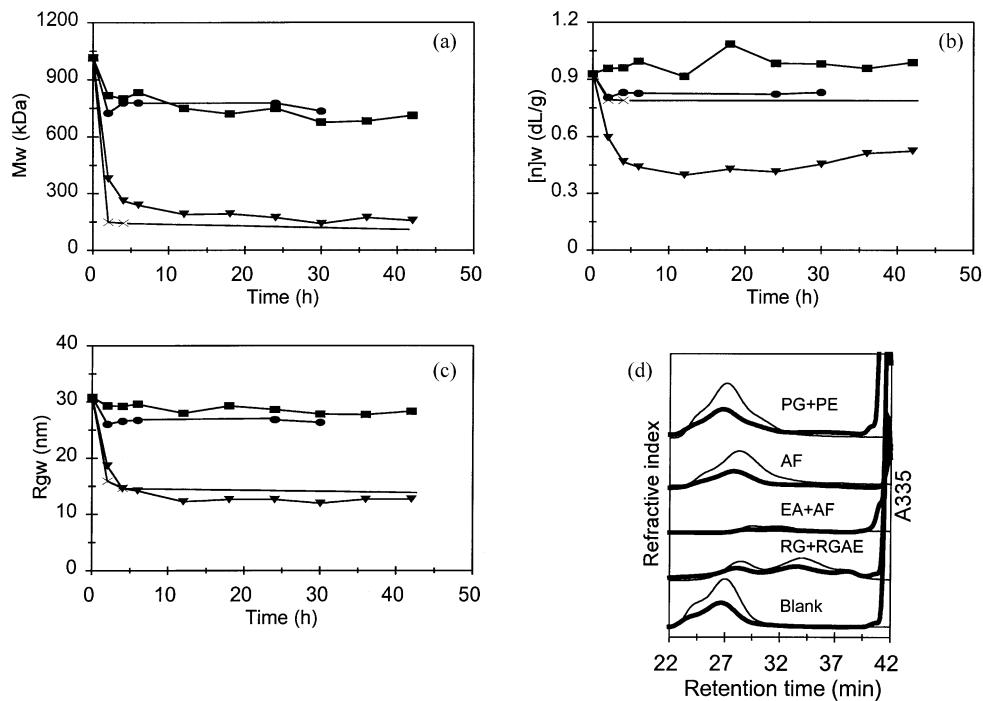


Fig. 2. Effect of enzymatic modification of pool I of the autoclave extract on M_w (A), $[\eta]_w$ (B), and R_{gw} (C) in time, and on the HPSEC elution pattern (after 42 h; D). ■: AF; ×: EA + AF; ▼: RGase + RGAE; ●: PG + PE. (D) Thick line: refractive index; thin line A_{335} .

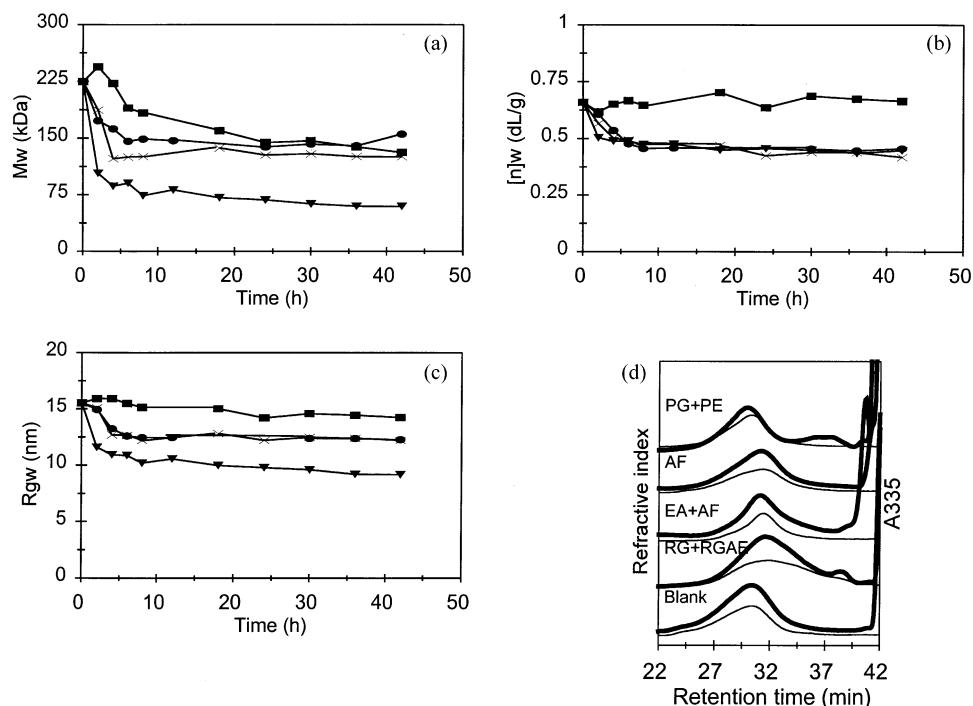


Fig. 3. Effect of enzymatic modification of pool II of the autoclave extract on M_w (A), $[\eta]_w$ (B), and R_{gw} (C) in time, and on the HPSEC elution pattern (after 42 h; D). ■: AF; ×: EA + AF; ▼: RGase + RGAE; ●: PG + PE. (D) Thick line: refractive index; thin line A_{335} .

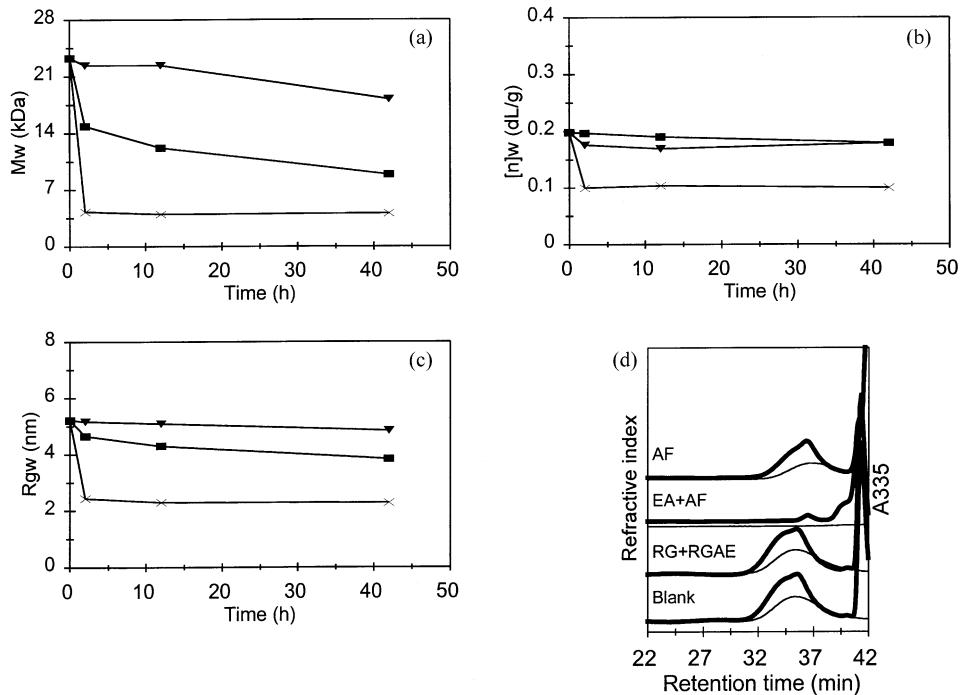


Fig. 4. Effect of enzymatic modification of pool IIIa of the autoclave extract on M_w (A), $[\eta]_w$ (B), and R_{gw} (C) in time, and on the HPSEC elution pattern (after 42 h; D). ■: AF; ×: EA + AF; ▼: RGase + RGAE. (D) Thick line: refractive index; thin line A_{335} .

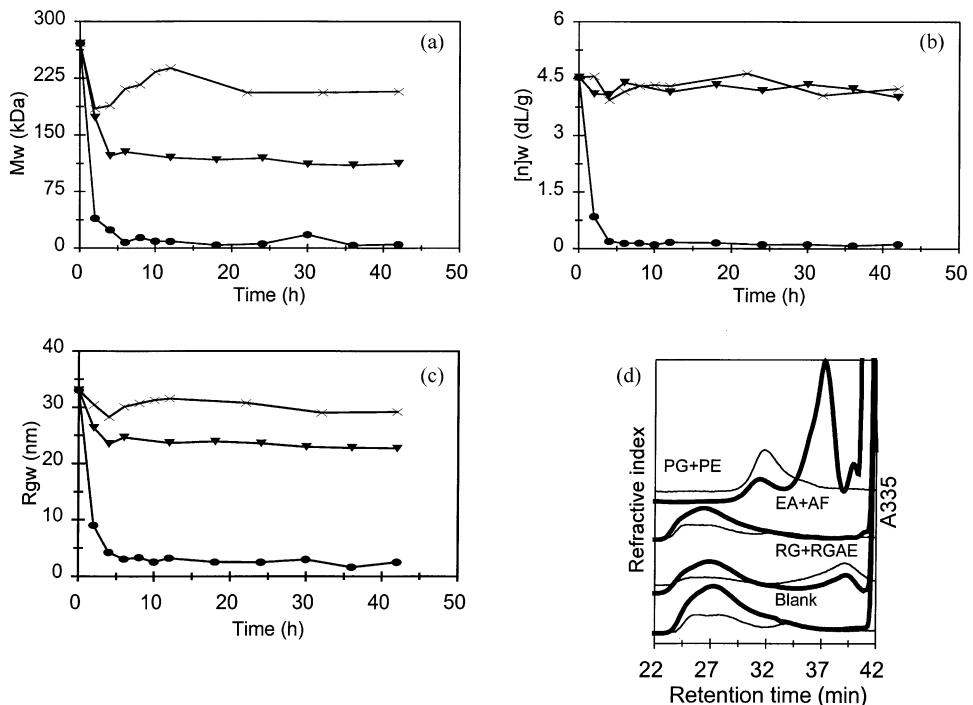


Fig. 5. Effect of enzymatic modification of ABP on M_w (A), $[\eta]_w$ (B), and R_{gw} (C) in time, and on the HPSEC elution pattern (after 42 h; D). ×: EA + AF; ▼: RGase + RGAE; ●: PG + PE. (D) Thick line: refractive index; thin line A_{335} .

3.3. Effect of enzymatic modification on pool I of the autoclave extract

The changes in M_w , $[\eta]_w$, and Rg_w during enzyme treatment of pool I of the autoclave extract are shown in Fig. 2a–c, while the HPSEC elution patterns before and after enzymatic treatment are shown in Fig. 2d.

Treatment of pool I of the autoclave extract with RGase + RGAE caused a rapid decrease of the M_w from 1020 to 157 kDa. Moreover, the $[\eta]_w$ and the Rg_w decreased approximately by a factor 2 and an additional population with a high ferulic acid content appeared upon HPSEC (34 min). The M_w of this new pool was approximately 50 kDa, whereas its $[\eta]_w$ was approximately 0.37 dl/g. Since RGase is known to degrade the more linear rhamnogalacturonan backbone only substituted with single unit galactose residues, the material which remains is assumed to consist of the rhamnogalacturonan subunit highly branched with arabinose.

The addition of AF to pool I reduced the M_w to 712 kDa. Interestingly, this resulted in a small increase of the $[\eta]_w$, whereas the Rg_w decreased slightly. As a consequence of the lower Rg_w , this pool eluted somewhat later upon HPSEC. Apparently a significant part of the arabinose can be removed from this pectin, without a loss (and in fact a small increase) of the $[\eta]_w$. AF is known to be active both on the terminal arabinofuranosyl residues in side-chains and on the non-reducing end of the arabinan backbone. Since the arabinan side-chains are present in much higher amounts than the non-reducing ends, AF degrades the side-chains in the arabinans faster than the backbone (Voragen, Rombouts, Searle-van Leeuwen, Schols & Pilnik, 1987). Because only part of the arabinose had been removed, considering the M_w after treatment with AF, it is assumed that mainly the side-chains of the arabinans had been removed and that these contribute little to the intrinsic viscosity of the rhamnogalacturonans. In the HPSEC chromatogram, the ratio A_{335} : RI increased after treatment with AF, which indicates that feruloylated arabinose is not removed, or with less preference than the non-feruloylated arabinose. Apparently, AF is hindered by the ferulic acid residues. Another possible explanation is the observation that the galactose residues in sugar beet pulp have a higher degree of feruloylation than the arabinose residues (Ralet et al., 1994).

Hydrolysis of the arabinan side-chains with EA + AF showed a decrease in M_w to 99 kDa. The $[\eta]_w$ of this population remained relatively high, apparently due to the fact that the backbone of the rhamnogalacturonan was not degraded by EA + AF. Since 58.8 mol% of this pool consisted of arabinose, which was present as side-chain of the rhamnogalacturonan, and since mono- and oligomers were not included in the calculation of the M_w , a maximum decrease in M_w of approximately 60% was expected. Therefore, the resulting low M_w after treatment with EA + AF is striking and might be ascribed to the

degradation of the backbone of arabinans that participate in cross-links between two different rhamnogalacturonan molecules in the original material. Diferulic acid bridges are present in the autoclave extract (Oosterveld, Grabber, Beldman, Ralph & Voragen, 1997), and are able to link multiple molecules together. ~10% of the ferulates present in the original autoclave extract is present as dimer (Oosterveld et al., 1997). Since the M_w of pool I is ~1000 kDa, a degree of polymerization can be estimated of 6700. About 70 mol% of the sugars is made up by arabinose and galactose and the ferulic acid content was 1.7 molecule per 100 molecules of arabinose and galactose. From this it can be calculated that on average every molecule contains four diferulates, which shows that cross-linking might occur to some extent. The presence of ferulic acid cross-links is supported by our findings that the molecular weight of this population dropped to 50% of its initial value after saponification (Oosterveld et al., 2000a).

It can be concluded that extensive degradation of the arabinan side-chains with EA + AF has little influence on the $[\eta]_w$ of a rhamnogalacturonan. These findings are in agreement with the results of Hwang and Kokini, who investigated the contribution of side-chains to the rheological properties of apple pectins with different neutral sugar contents (Hwang & Kokini, 1991), although the neutral sugar contents were relatively low in all the pectins investigated. They found that the degree of branching had little effect on the intrinsic viscosity, which is determined at low concentrations, but that the side-chains of pectins were significantly involved in entanglements of the pectin molecules in concentrated solutions.

PG + PE caused a small decrease in M_w , $[\eta]_w$, as well as in Rg_w . Also, a small shift in hydrodynamic volume was observed upon HPSEC. This limited decrease in M_w shows that this enzyme combination was only slightly active on this pool, showing that no homogalacturonan regions are present internally in this rhamnogalacturonan population. Addition of pectin acetyl esterase, which is known to increase the activity of PE (Oosterveld et al., 2000b) and therefore indirectly increases the activity of PG, besides PG + PE to the parental extract did not increase the degradation of this rhamnogalacturonan population (unpublished data), but only increased the degradation of the separately present homogalacturonan populations (pool III). These results indicate that pieces of homogalacturonan are possibly located at the extremities of the rhamnogalacturonan molecules or that some homogalacturonans are present as a separate population. Since PG + PE are active on the ‘smooth’ regions and not on the ‘hairy’ regions, the small decrease in M_w indicates that the hairy regions of sugar beet pectin may have a very high M_w .

3.4. Effect of enzymatic modification on pool II of the autoclave extract

Pool II of the autoclave extract, a highly branched

rhamnogalacturonan with a lower M_w than pool I, was incubated with the same enzyme combinations as pool I (Fig. 3). The results were rather similar to the results found for pool I. Hydrolysis with the enzymes RGase + RGAE caused a large decrease in M_w , $[\eta]_w$, and Rg_w . Degradation of the arabinan side-chains of pool II with AF decreased the M_w , after a small initial increase in M_w . We assumed that this small increase was due to inaccuracies in the measurements, and that is was not significant. The $[\eta]_w$ and the Rg_w of pool II changed relatively little after degradation with AF. Hydrolysis of the arabinan side-chains of pool II by EA + AF led to a relatively large decrease in M_w . Although the M_w decreased to a value similar to the value found after treatment with AF, the $[\eta]_w$ and Rg_w decreased to lower values. Hydrolysis of homogalacturonan regions in pool II by PG + PE led to a relatively large decrease in M_w . If possible homogalacturonans stretches are only located at the extremities of the molecule, as was suggested for pool I, this decrease seems rather large. Possibly some high M_w homogalacturonans eluted together with the rhamnogalacturonans in small amounts, as was also indicated by the higher galacturonic acid content in this pool as compared with pool I. The formation of an additional peak with no visible A_{335} absorption at 37 min upon HPSEC after degradation with PG + PE is in agreement with this assumption. The $[\eta]_w$ and Rg_w also decreased to relatively low values, which shows that this homogalacturonan fraction has a relatively large influence on these parameters.

3.5. Effect of enzymatic modification on pool IIIa of the autoclave extract

The arabinan fraction pool IIIa of the autoclave extract was also treated with the enzyme combinations mentioned above (Fig. 4).

Modification by RGase + RGAE resulted only in a small change in M_w . This is in agreement with the assumption that some arabinans are still linked to a small piece of rhamnogalacturonan backbone, too small to bind to the anion-exchange column, as was previously shown by the methylation analysis of the arabinans (Oosterveld et al., 2000a). The $[\eta]_w$ did not change, since this is determined by the longest backbone of the fragment, in this case probably the arabinan backbone. The HPSEC diagram hardly changed as compared with the blank.

Hydrolysis of the arabinans with AF decreased the M_w from 23 to 9 kDa. From these results a degree of polymerization of the arabinans can be calculated of 130–170 residues before and 60–70 after modification with AF. However, the $[\eta]_w$ did not change. This can be explained by the fact that AF predominantly removes the side-chains from the arabinans, having little effect on the length of the backbone, which determines the $[\eta]_w$ to a large extent. So, probably the backbone has a length of 60–70 residues. This would implicate that more than 45–65% of the arabinose

residues are present as single unit or oligomeric side group of the arabinan main-chain. Also, methylation analysis showed that approximately 35% is present as terminal residues indicating that on average the side-chains of the arabinans are relatively short (Oosterveld et al., 2000a). Cooper et al. (1992) described the production of linear arabinans with AF. They estimated a length of 50–80 residues after linearization, which is in good agreement with the value described in this paper.

Incubation of the arabinans with the enzymes EA + AF led to a decrease in both M_w and $[\eta]_w$ of the arabinan population. In the HPSEC elution pattern only a very small population remained. It is speculated that this population consists of glucomannans, which represent less than 5 mol% of the sugars initially present in the arabinan fraction (Table 1). The M_w of this fraction was 4 kDa.

The effect of treatment of the arabinans with PG + PE was not determined, since the galacturonic acid content of the arabinans was very low.

3.6. Effect of enzymatic modification on an ABP

Besides rhamnogalacturonans and arabinans obtained from an autoclave extract from beet pulp, also an acid extracted pectin from sugar beet pulp (ABP) was incubated with the enzyme combinations RGase + RGAE, PG + PE, and EA + AF (see Fig. 5). ABP consists primarily of homogalacturonans. As the arabinose content of this pectin is low (Table 1), and the effect of arabinan degrading enzymes on the physico-chemical properties was expected to be negligible, treatment with AF was omitted.

Treatment of ABP with RGase + RGAE decreased the M_w from 271 to 112 kDa, due to hydrolysis of the rhamnogalacturonan backbone present in the hairy regions of ABP. Surprisingly, the $[\eta]_w$ hardly changed. Most of the material eluted at the same retention time upon HPSEC. However, a low M_w population with a high A_{335} absorption appeared at 38 min in low quantities. It made us conclude that the hairy regions in this pectin are located at the extremities of the molecules. It is also possible that a pectin population is degraded by this enzyme, which is present in small amounts and with a very high M_w . Berth, Dautzenberg & Rother (1994) described a pectin population, which was present in minute amounts. This population had such a high M_w that it significantly influenced the M_w of the whole pectin extract.

Our results show that the rhamnogalacturonan backbone from ABP contributes little to the $[\eta]_w$. On the other hand treatment with RGase + RGAE causes a loss of the gelling properties with Ca^{2+} , as we showed in another study (Oosterveld et al., 2000b).

Modification of the arabinan side-chains by addition of EA + AF to ABP decreased the M_w to 207 kDa. This decrease was somewhat larger than was expected based on the low arabinose content (7 mol%). Both the Rg_w and the

$[\eta]_w$ hardly changed. Upon HPSEC no clear changes were observed.

Modification of the smooth homogalacturonan regions of ABP with PG + PE decreased the average M_w rapidly to a value of 5 kDa. Furthermore, the $[\eta]_w$, and Rg_w also decreased rapidly. This is in agreement with the results found by Guillon and Thibault, who also found a rapid decrease in M_w and $[\eta]_w$ for an ABP during treatment with PG + PE (Guillon & Thibault, 1990). After treatment of ABP with PG + PE only some hairy fragments remained. Based on our results the average M_w of these fractions is 50 kDa (Fig. 5D), which is much lower than those obtained by autoclaving (see above). On the other hand, it is somewhat higher than the results found by Guillon and Thibault, who found a viscosity average molecular weight of 12 kDa after treatment of an ABP with PG + PE (Guillon & Thibault, 1990).

4. Concluding remarks

Enzymatic modification of various types of pectic polysaccharides obtained from sugar beet pulp with specific enzymes affected structural characteristics as well as rheological characteristics of these polysaccharides. This allowed studies of the effect of structural changes on the rheological properties.

In most cases we found that the enzymatic modifications employed in this study decreased the $[\eta]_w$ of the polysaccharides investigated. Obviously, this occurred when backbone degrading enzymes were used. However, we found that degradation of the rhamnogalacturonan backbone of an ABP has relatively little influence on the $[\eta]_w$ which made us conclude that the rhamnogalacturonans present in this type of pectin are mainly located at the extremities of the pectin molecules. Some indications were found that the rhamnogalacturonans obtained by autoclave extraction are linked through the arabinose side-chains, probably by diferulic acid cross-links. However, further study is needed to confirm this. Removal of the arabinans present in these rhamnogalacturonans with enzymes had relatively little effect on the $[\eta]_w$, although the M_w of the populations decreased significantly. This shows that the arabinan side-chains contribute little to the $[\eta]_w$ of ABPs.

In our experiments we found that enzymatic modification with glycanases can be used to significantly change the structural characteristics of pectic polysaccharides without a significant loss of viscosity. In this way glycanases can be used for the removal of side-chains of the pectic molecule, which limit the physico-chemical properties, as described for the production of debranched arabinans (Cooper et al., 1994), and for the improvement of the oxidative cross-linking reaction of beet pectins by removal of the arabinan sidechains (Guillon & Thibault, 1990).

An important prerequisite for the introduction of the use of enzymes for the modification of beet pectins on industrial

scale, is the availability of sufficient amounts of enzymes at reasonable prices. The use of cloned polysaccharide degrading enzymes as described by Kofod, Kauppinen, Christgau, Andersen, Heldt-Hansen & Dörreich (1994) is assumed to be an important step towards the introduction of these enzymes on industrial scale.

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