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## Sequential fractionation of feruloylated hemicelluloses and oligosaccharides from wheat bran using subcritical water and xylanolytic enzymes†

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Wheat bran is a major by-product of cereal production that still has limited use for advanced nutritional and material applications. A sequential process using subcritical water, membrane filtration and selective enzymatic treatments has been designed for the combined fractionation of functional high molar mass hemicelluloses (over  $10^5$  g mol<sup>-1</sup>) and oligosaccharides from wheat bran. This process not only offers increased total solid yield compared with conventional protocols based on alkaline extraction, but it also preserves the inherent functionalities of the phenolic groups that substitute the carbohydrate structures of the extracted hemicelluloses. Feruloylated arabinoxylans (F-AX) with high molar mass and significant radical scavenging activity can be isolated from the subcritical water extract. Structurally different oligosaccharides, including mixed-linkage  $\beta$ -D-glucan oligosaccharides (BGOs) and arabinoxylo-oligosaccharides (AXOs) can be recovered from the eluent after membrane filtration. The crosslinked residue after subcritical water extraction was further treated with xylanolytic enzymes to release valuable feruloylated arabinoxylo-oligosaccharides (FAXOs). The oligo- and polysaccharide fractions isolated from this sequential process show great potential for use as prebiotic or platform chemicals, and as polymeric matrices for carbohydrate-based materials with radical scavenging properties, respectively.

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## Introduction

Cereal processing generates large volumes of by-products that have not been exploited to their full potential for advanced nutritional and material applications.<sup>1</sup> Each ton processed by the wheat milling industry generates up to 0.25 tons of wheat bran that corresponds to 14–16% of the cereal kernel.<sup>2</sup> Due to its content of valuable biomacromolecules, such as polysaccharides (mainly hemicelluloses) and phenolic compounds, wheat bran has gained renewed interest as a raw material to be used in second generation biorefineries.<sup>2–4</sup>

Xylan is one of the most abundant biopolymers in nature as a major non-cellulosic component of plant cell walls. In par-

ticular, arabinoxylan (AX) constitutes the main hemicellulose in grasses and cereal grains, with an average composition of 20–40% depending on the species and tissue considered.<sup>5,6</sup> AX structure is based on a linear backbone of (1→4)-linked  $\beta$ -D-xylopyranose (Xylp) units, mainly 2-O-, 3-O- and/or 2,3-di-O-substituted by  $\alpha$ -D-arabinofuranosyl (Araf) units. The xylan backbone can be 2-O-substituted to a lesser extent by  $\alpha$ -D-glucuronic acid (GlcA) or its 4-O-methyl derivative (4-O-Me-GlcA). The extent and distribution of the glycosyl substitutions depend on the biological source and tissue.<sup>7</sup> Phenolic compounds, mainly ferulic acid (FA), can further substitute the Araf residues in the C-5 position.<sup>5,8</sup> These phenolic compounds can dimerize with other phenolic moieties from neighboring AX molecules, creating a complex covalent network structure. This heterogeneous and crosslinked molecular architecture is responsible for the recalcitrance of bran, and imposes major challenges for the biotechnological valorization of the constituent biomolecules.

Pretreatments using harsh alkaline conditions have been traditionally used for the extraction of hemicelluloses from cereal and other plant biomasses.<sup>9,10</sup> However, these harsh conditions cause the loss of valuable functional groups, promoting the removal of acetyl, uronic acid and phenolic

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substitutions.<sup>9</sup> Other methods have also been developed for the extraction and purification of AX from cereal by-products, including acid pretreatment,<sup>11</sup> microwave assisted extraction,<sup>12</sup> ultrasound assisted extraction,<sup>13,14</sup> and steam explosion extraction.<sup>10,15</sup> As an alternative, subcritical water extraction (SWE), also called pressurized hot water extraction (PHWE), appears as a promising green method for the isolation of hemicellulose fractions with preserved molecular functionalities and high molecular weight. The use of subcritical conditions (at temperature below critical point and pressure high enough to maintain liquid state) changes dramatically the properties of water, including density surface tension, polarity, viscosity and diffusion. Mass transfer during the extraction process is favored due to the decrease in the viscosity and an increase on diffusivity.<sup>16</sup> Subcritical water has proven to be effective for the extraction of diverse bioactive and nutraceutical compounds, including polysaccharides from plants and food matrices.<sup>16–19</sup> Moreover, subcritical water has been successfully used for the extraction of hemicelluloses from wood, with the possibility for scaling up.<sup>20–23</sup> In parallel, the use of carbohydrate-active enzymes during biomass pretreatment further facilitates the extraction and fractionation of the polysaccharide components, showing good specificity.<sup>24</sup> Xylanolytic enzymes that act on the xylan backbone and on the glycosyl decorations can be used to deconstruct xylan polysaccharides into mono- and oligosaccharides that can be further converted into bioethanol and platform chemicals.<sup>25–28</sup> However, this process drastically reduces the molar mass of the hemicelluloses and restricts their use in biopolymeric material applications.

A sequential process that combines subcritical water and selective carbohydrate-active enzymes is here proposed to fractionate polymeric hemicelluloses and oligosaccharides from cereal by-products while preserving their native phenolic functionalities. The extraction conditions were optimized under subcritical water and compared to the alkaline processes. Finally, the effect of different carbohydrate-active enzymes was evaluated, as a mean to disrupt the crosslinked residues and further release of functional biomolecules.

## Results and discussion

### Composition of the defatted wheat bran

The summarized and extended compositions of the defatted wheat bran (wb) and destarched wheat bran (Dwb) are shown in Table 1 and ESI Table S1,<sup>†</sup> respectively. Polysaccharides represent the main constituents of wheat bran, with arabinoxylans (30%) and glucans (27%), including starch (17%), mixed-linkage  $\beta$ -D-glucan (6%) and cellulose (Fig. 1a), being the main populations, in agreement with previous reports.<sup>3,8,29,30</sup> Residual presence of other hemicelluloses and pectins can be also detected by the relative abundance of other sugars in minor amounts (ESI Table S1<sup>†</sup>). Proteins and phenolic compounds represent the non-carbohydrate fractions of wb and Dwb. The protein content measured based on nitrogen quantification (Dumas method) was 19.7%, which is slightly higher

**Table 1** Summarized chemical composition of the wheat bran

	wb	Dwb
Moisture content <sup>a</sup> (%)	22.0 (1.1)	18.5 (1.1)
Total carbohydrate <sup>b</sup> (mg per g DW)	595.9 (0.6)	682.4 (0.1)
AX (as % of total carbohydrate) <sup>c</sup>	48.6 (0.9)	59.4 (0.4)
AX <sup>c</sup> (mg per g DW)	289.6 (0.5)	405.3 (0.1)
A/X <sup>c</sup>	0.59	0.66
Glc (as % of total carbohydrate) <sup>c</sup>	45.5 (0.7)	35.7 (0.9)
Starch <sup>d</sup> (mg per g DW)	169.4 (0.6)	28.9 (2.9)
Mixed-linkage $\beta$ -D-glucan (mg per g DW)	57.0 (3.0)	60.8 (1.0)
Other polysaccharides <sup>e</sup> (mg per g DW)	35.2 (0.4)	33.5 (0.1)
Protein <sup>e</sup> (mg per g DW)	202	197
Hydroxycinnamic acids <sup>f</sup> (mg per g DW)	2.2 (0.1)	2.4 (0.16)
Lignin <sup>g</sup> (mg per g DW)	90.0 (3.8)	117.8 (1.4)
Other compounds <sup>h</sup> (mg per g DW)	109.9 (0.8)	0.4 (0.05)

The values represent the average of triplicate measurements. The standard deviations are presented in parentheses. <sup>a</sup> Moisture determined gravimetrically after oven drying (110 °C, 24 h). <sup>b</sup> Carbohydrate content determined by phenol-sulfuric acid method.<sup>38</sup> <sup>c</sup> Determined from the monosaccharide composition (Table S1). <sup>d</sup> Starch and  $\beta$ -D-glucan content determined enzymatically (Megazyme). <sup>e</sup> Protein content determined using Dumas method.<sup>39</sup> <sup>f</sup> Hydroxycinnamic acid content determined by GC-MS.<sup>33</sup> <sup>g</sup> Klason lignin after acid hydrolysis (H<sub>2</sub>SO<sub>4</sub>). <sup>h</sup> Determined based on the difference between the sum of total carbohydrate, protein, phenolic and lignin and the initial dry weight.

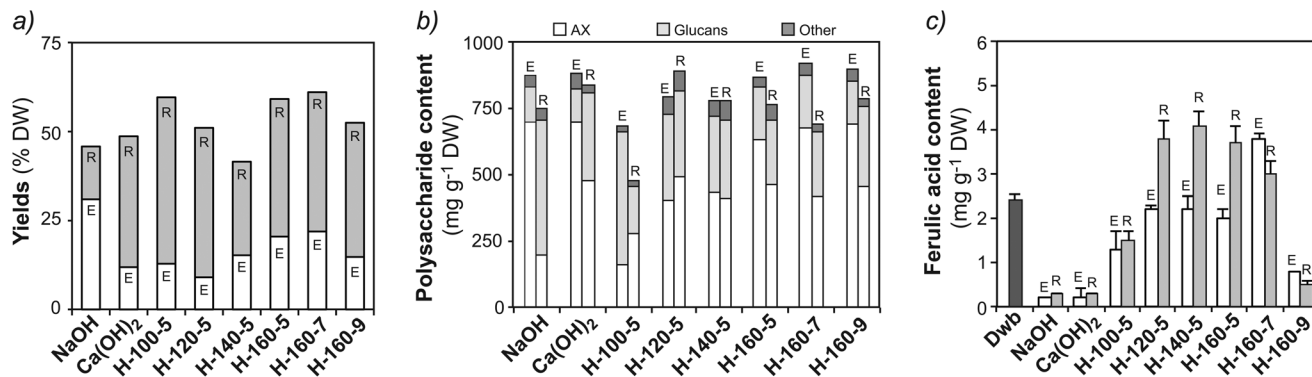
compared to other studies (13.2–18.4%).<sup>3,29,31,32</sup> The total hydroxycinnamic acid content was 2.2–2.4 mg g<sup>-1</sup>, with ferulic acid (FA) being the dominating compound. Similar or lower amounts of FA were reported for insoluble cell wall fractions of wheat endosperm<sup>33</sup> and extracted AXs from wheat bran.<sup>8,34</sup> Phytic acid (9.6 ± 0.5 mg g<sup>-1</sup>) and acetyl content (degree of acetylation, DA = 0.03) showed lower values than previously reported for wheat bran.<sup>8,35</sup> Destarching effectively removed starch to below 3%, while enriching arabinoxylan (405.3 ± 0.1 mg per g DW) with an arabinose over xylose ratio (A/X) of 0.66, as previously reported for wheat bran (A/X ~ 0.5–0.8).<sup>36,37</sup>

### Proposed scheme for the cascade fractionation of hemicelluloses and oligosaccharides from wheat bran

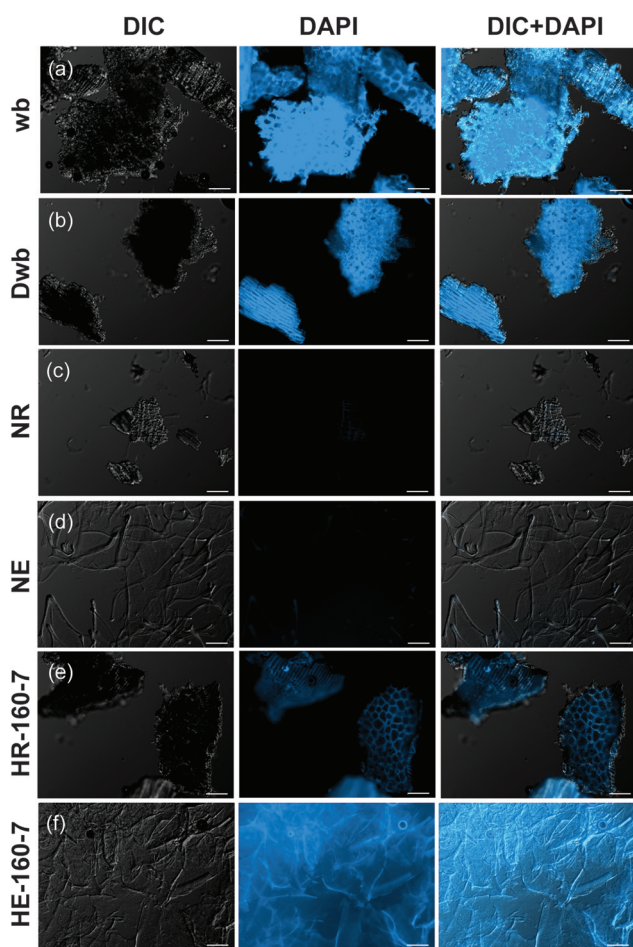
The sequential process combining subcritical water extraction, membrane filtration and xylanolytic enzymes was proposed in order to fractionate simultaneously feruloylated arabinoxylans in polymeric form and oligosaccharides from wheat bran (Fig. 1a). Before extraction, defatted wheat bran was destarched (Dwb) and no other pretreatments were required. The proteins were not removed from Dwb, since they may play important structural roles when preparing hybrid carbohydrate–protein biomaterials. Further enzymatic removal of proteins by proteases could be implemented in conjunction with destarching if required. Subcritical water (SWE) under pressurized conditions was selected as the extraction system due to its efficiency (short extraction times), process sustainability (green solvent system) and the expected ability to maintain the phenolic functionalities in the hemicellulose fractions. Membrane fractionation was introduced to separate the extracted biomolecules into high and low molar mass fractions. Finally, selective enzymatic treatments that target







**Fig. 2** Comparison of alkaline (NaOH and Ca(OH)<sub>2</sub>) and subcritical water (SWE) extractions at different pH and temperature conditions: (a) total solid yields (based on % DW of Dwb); (b) polysaccharide composition (mg per g DW) of the extracts and residues in terms of arabinoxylan (AX), glucans and minor components. The composition was determined from the monosaccharide analysis after total acid hydrolysis (TFA – extracts and H<sub>2</sub>SO<sub>4</sub> – residues) presented in ESI Table S2;† (c) ferulic acid content (mg per g DW). Note: H: subcritical water; E: extract; R: residue.



**Fig. 3** Fluorescence microscopy of wheat bran samples: (a) defatted wheat bran (wb); (b) destarched wheat bran (Dwb); (c) residue from NaOH extraction (NR); (d) extract from NaOH extraction (NE); (e) residue from SWE at 160 °C pH 7.0 (HR-160-7); (f) extract from SWE at 160 °C, pH 7.0 (HE-160-7). The images were obtained using differential interference contrast (DIC) (left column), DAPI filter (middle column) and DIC merged with DAPI filter (right column); optical objective 40x; scale bar 100 μm.

(Fig. 3c and d), which demonstrates the cleavage of the ferulic acid moieties under such pH conditions.

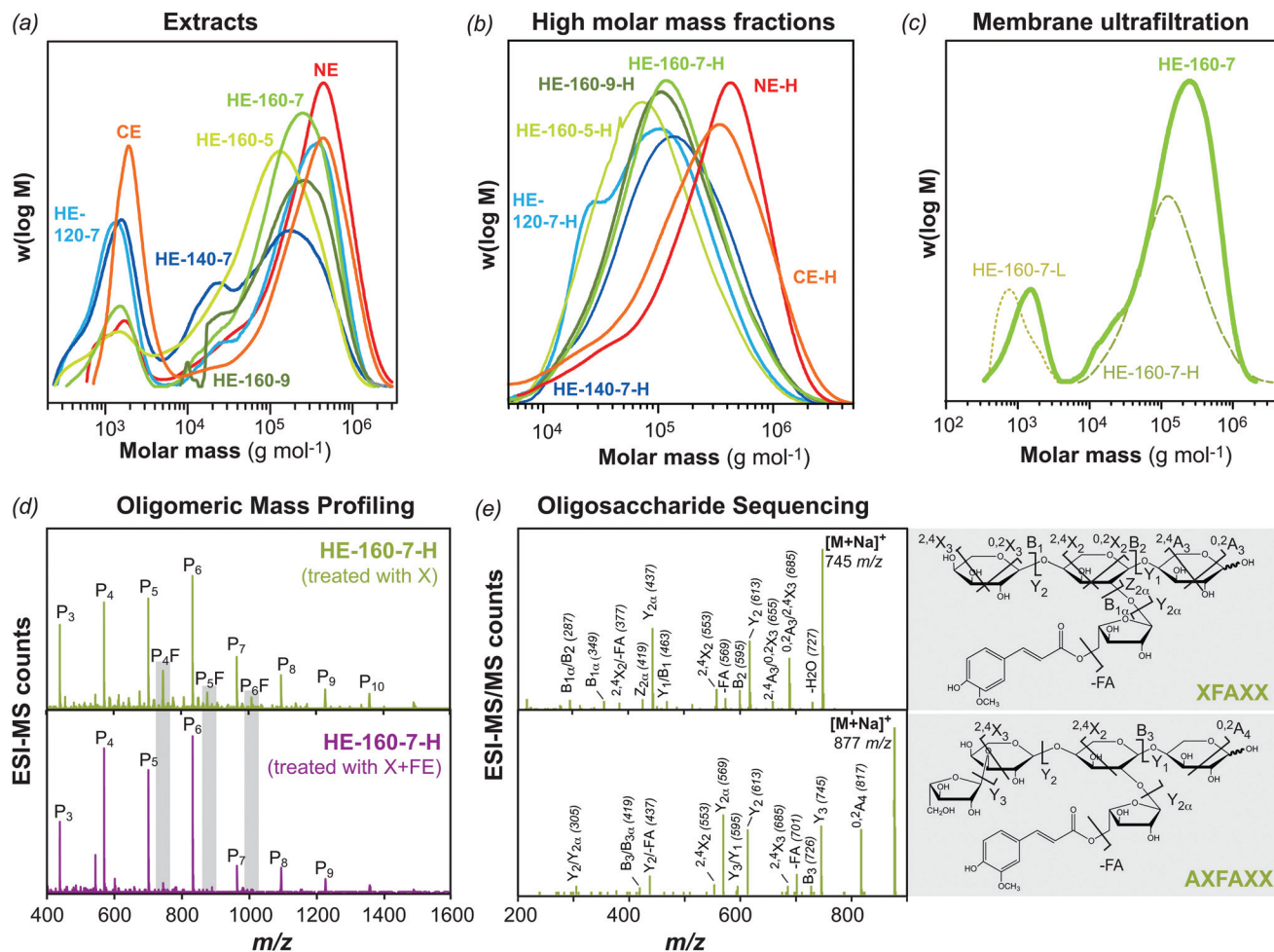
Conversely, auto-fluorescence can be observed in both the extract and residue after SWE, indicating the preservation of ferulic acid while using this method (Fig. 3e and f). The extracts from SWE exhibit evenly distributed autofluorescence, with higher intensity along the observed fibrillar morphologies.

On the other hand, specific localization of the autofluorescence in the outer layers can still be observed in the residues after SWE (Fig. 3e), which indicates that non-extractable cross-linked populations with high ferulic acid densities remain in the outer bran layers. This suggests that subcritical water extracts preferentially AXs from the inner aleurone and intermediate layers, in agreement with the obtained A/X ratios. Such evidence is reinforced by the compositional analyses of the residues after SWE, which exhibit significant content of non-extractable AX (416 mg g<sup>-1</sup>; Fig. 2b) with high phenolic content (3.0 mg g<sup>-1</sup> of FA; Fig. 2c and ESI Table S2†). These resilient AX populations remaining in the residue were the targets for the enzymatic treatments in the next steps.

### Fractionation of oligo- and polysaccharide populations

The alkaline and subcritical water extracts exhibited bimodal molar mass distributions with two main (macro)molecular populations, a high molar mass polymeric fraction (10<sup>5</sup>–10<sup>6</sup> g mol<sup>-1</sup>) and a low molar mass fraction (10<sup>3</sup>–10<sup>4</sup> g mol<sup>-1</sup>) (Fig. 4a). The number-average molar mass ( $M_n$ ), weight-average molar mass ( $M_w$ ) and the dispersity index of the extracted fractions are presented in ESI Table S2.† Closed dialysis (Fig. 4b) and membrane ultrafiltration (Fig. 4c) were applied to fractionate the low molar mass (L) from the high molar mass (H) fractions. Both procedures efficiently separated the oligosaccharides from the polysaccharide populations. After membrane fractionation, the total solid yields; the polysaccharide composition and the molecular structure were evaluated for the polymeric and oligomeric fractions (Table 2).





**Fig. 4** Molecular properties of the hemicellulose fractions obtained from destarched wheat bran (Dwb). (a) Molar mass distributions  $w(\log M)$  of the extracts after alkaline (NE and CE) and subcritical water extraction (HE); (b) molar mass distributions of the high molar mass fractions (H) after closed dialysis; (c) molar mass distributions of the polymeric (H) and oligomeric (L) fractions after membrane ultrafiltration of HE-160-7; (d) oligomeric mass profiling by ESI-MS of the polymeric fraction HE-160-7-H after enzymatic depolymerization with a xylanase (X) and a feruloyl esterase (FE); (e) oligosaccharide sequencing by ESI-MS/MS of selected feruloylated ions ( $m/z$  745 and  $m/z$  877) and proposed structures from the fragmentation pattern. The mass-to-charge ( $m/z$ ) values of the fragments are presented in parenthesis.

In general, the total yields for the high molar mass fractions were higher than for the oligomeric fractions, with ranges between 60–80% and 10–40%, respectively. The polysaccharide fractions were composed mainly of AX, with contents ranging between 70–90%. The alkaline (NE-H) fraction presented the highest AX content, whereas the AX content in the fractions extracted by subcritical water (HE-H) increased progressively with the extraction temperature, as reported earlier. The minor polysaccharide components were assigned to mixed-linkage  $\beta$ -D-glucan (12–20%) and arabinogalactan (AG) (5–8%), based on the results from glycosidic linkage analysis (ESI Table S3<sup>†</sup>). The presence of  $\beta$ -D-glucans and AG was higher for the fractions extracted at lower temperatures, which indicates that these populations are easier to extract than AX.

The A/X ratio and the substitution pattern determine the solubility, macromolecular conformation, physico-chemical and functional properties of these macromolecules.<sup>5,6</sup> The A/X

ratio for the NE-H AX fraction was 0.74, and on average 0.49 for the high molar mass fractions extracted by subcritical water. Again showing that the alkaline extracted AX is more substituted than the hydrothermal counterpart (ESI Table S3<sup>†</sup>). The average substitution pattern of the AX fractions was analyzed by glycosidic linkage analysis. Four basic building blocks can be seen in the backbone structure of AXs, which are responsible for determining the AX substitution pattern (Fig. 1b). The D-Xylp residues along the backbone can be unsubstituted (X), mono-substituted in the C2 (X<sup>2</sup>A), mono-substituted in the C3 (X<sup>3</sup>A), or di-substituted in both, C2 and C3 (X<sup>2,3</sup>A). The ratio of substituted D-Xylp units in the backbone was normalized towards the unsubstituted Xylp residues (X) and presented in Table 2. Di-substitutions (X<sup>2,3</sup>A) and unsubstituted D-Xylp residues (X) are more abundant than mono-substituted Xylp residues, whereas O-3 substitutions (X<sup>3</sup>A) are preferred over O-2 substitutions (X<sup>2</sup>A).<sup>5,6,48</sup>



Table 2 Molecular characterization of the polysaccharide (H) and the oligosaccharide (L) fractions after membrane filtration

	NE		HE-120-5		HE-140-5		HE-160-5		HE-160-7		HE-160-9	
	H	L	H	L	H	L	H	L	H	L	H	L
Yields <sup>a</sup> (% DW)	79.1	9.2	68.8	28.8	78.3	19	58.6	40.6	65.9	32.9	76.8	21.8
Carbohydrate content <sup>b</sup> (mg per g DW)	871.9 (3.4)	750.0 (4.5)	839.9 (24.9)	616.3 (1.2)	715.8 (7.6)	743.3 (9.9)	815.4 (1.8)	564.2 (27.5)	986.8 (5.2)	598.1 (7.1)	863.1 (0.4)	660.6 (26.4)
AX <sup>c</sup> (%)	92.0	54.6	77.5	50.4	77.0	73.6	76.4	70.5	85.6	80.4	85.9	75.2
X <sup>d</sup>	100	100	100	100	100	100	100	100	100	100	100	100
X <sup>2</sup> A <sup>d</sup>	14	23	9	8	10	11	11	11	15	12	10	14
X <sup>3</sup> A <sup>d</sup>	30	45	19	14	16	25	22	24	24	19	20	30
X <sup>2,3</sup> A <sup>d</sup>	45	40	32	28	27	41	25	34	27	26	23	37
β-Glucan <sup>c</sup> (%)	1.6	10.7	14.1	28.1	19	22.9	18.6	21.9	12.0	18.1	10.9	18.1
(Arabino)galactan <sup>c</sup> (%)	5.4	5.5	8.3	1.3	3.4	1.8	4.8	4.5	2.4	0.0	2.1	3.0
Protein content <sup>e</sup> (mg per g DW)	104.1 (2.2)	51.6 (1.6)	109.7 (32.5)	155.1 (24.9)	88.9 (6.2)	71.1 (10.1)	78.0 (5.3)	51.7 (2.4)	81.0 (1.9)	46.3 (14.8)	71.2 (1.5)	49.2 (3.4)
Ferulic acid content <sup>f</sup> (mg per g DW)	0.2 (0.04)	0.2 (0.04)	3.2 (0.14)	2.2 (0.14)	3.5 (0.13)	1.0 (0.10)	3.7 (0.14)	1.8 (0.13)	4.9 (0.13)	2.2 (0.12)	0.3 (0.1)	0.2 (0.10)
M <sub>n</sub> <sup>g</sup> (g mol <sup>-1</sup> )	135 640	n.d.	69 622	1494	67 330	1048	82 419	1763	135 040	1048	106 100	1920
M <sub>w</sub> <sup>g</sup> (g mol <sup>-1</sup> )	458 340	n.d.	182 170	1060	138 410	833	133 920	1314	193 980	839	187 840	1551

<sup>a</sup>Total yields (as % of initial DW of Dwb used for each extraction condition) determined gravimetrically after membrane filtration and freeze-drying. <sup>b</sup>Carbohydrate content determined by the phenol-sulfuric acid method. <sup>c</sup>Polysaccharide composition (as % of the total carbohydrate content) determined from the glycosidic linkage analysis (Tables S3 and S4). <sup>d</sup>Average backbone substitution pattern of AX. The values are normalized by the unsubstituted Xylp (X) units (see Fig. 1b for nomenclature). <sup>e</sup>Protein content determined using Dumas method. <sup>f</sup>Ferulic acid content determined by GC-MS. <sup>g</sup>Average molar masses determined by SEC-DRI. Note: The standard deviation is shown in parentheses.

Interesting differences can be observed in the substitution pattern of the polymeric extracts from subcritical water at different pH and temperature conditions. The presence of di-substituted X<sup>2,3</sup>A units progressively decreases and the relative presence of mono-substituted X<sup>3</sup>A and X<sup>2</sup>A units increases with higher extraction temperatures. Moreover, mild acidic (pH 5.0) and alkaline (pH 9.0) conditions result in AX fractions with lower substitution, compared to neutral (pH 7.0) conditions. These results suggest that it may be possible to modulate the substitution pattern of the AXs by tuning the pH and temperature conditions during SWE. Further pH and temperature conditions should be tested to verify this hypothesis.

The high molecular weight fractions showed a two-fold increase in the FA content compared to the native wheat bran. This indicates that the presence of feruloylated arabinoxylan (F-AX) populations was enriched during the extraction and fractionation processes. To confirm that the ferulic moieties are indeed covalently linked to the AX polysaccharide structures, the hydrothermal extract (HE-160-7-H) was enzymatically hydrolyzed with an *endo*-1,4-β-xylanase (X) alone and in combination with a feruloyl esterase (FE) and the resulting oligosaccharides were profiled by ESI-MS (Fig. 4d). The presence of FA in the oligosaccharide fragments was confirmed by the difference of *m/z* 176 between the non-feruloylated and the feruloylated sodium adducts ([M + Na]<sup>+</sup>). The ions assigned to the feruloylated oligosaccharides at *m/z* 745 and *m/z* 877 (corresponding to P<sub>4</sub>F and P<sub>5</sub>F, respectively) drastically decrease when treated with the FE, demonstrating the covalent linkage between the carbohydrate and phenolic moieties. The feruloylated motifs present in polymeric F-AX were further sequenced by tandem ESI-MS/MS. The fragments were assigned as pro-

posed by Domon and Costello.<sup>49</sup> The structures can be proposed as β-D-Xylp-(1→4)-[(5-*O-trans*-feruloyl)-α-L-Araf-(1→2)-β-D-Xylp-(1→4)-D-Xylp (XFAXX) and α-L-Araf-(1→3)-β-D-Xylp-(1→4)-[(5-*O-trans*-feruloyl)-α-L-Araf-(1→2)-β-D-Xylp-(1→4)-D-Xylp (AXFAXX) (Fig. 4e). The presence of FA moieties in the polysaccharide structure is crucial for the scavenging activity of the polymeric fractions and for their further applicability, as discussed later.

The molar mass distributions and average molecular weights of the polymeric fractions were investigated by SEC analyses (Fig. 4b and Table 2). The polysaccharides in the alkaline extract exhibit higher molar mass ( $4.5 \times 10^5$  kg mol<sup>-1</sup>) than those in the subcritical water extracts (*M<sub>w</sub>* between  $1.3$ – $1.8 \times 10^5$  g mol<sup>-1</sup>). These differences arise from the different extractability of the macromolecular populations under both processes. The harsh alkaline conditions disrupt the ferulic crosslinks in wheat bran, releasing AX populations with higher molar masses but without the covalently attached phenolic functionalities. Conversely, subcritical water is capable of extracting AX fractions from the most accessible parts of wheat bran with maintained feruloylated functionalities, but showing lower molar mass.

Another important issue is the potential degradation induced by subcritical water extraction at high temperatures. In a previous study on the extraction of hemicelluloses from wood (spruce), the combination of higher temperatures and acidic pH during subcritical water caused autohydrolysis and thus a reduction of the molar mass of the extracted fractions.<sup>21</sup> In this study, a buffered aqueous solution was used to avoid such pH variations and to minimize degradation. The combination of an increased temperature (from 140 °C and 160 °C) and mild acidic conditions (pH 5.0) during SWE causes a



slight reduction in the molecular weight of the polymeric fractions, which can be attributed to acid-catalyzed hydrolytic processes. Additionally, the acidic conditions during SWE may induce, as well, degradation of the Araf substitutions, which will affect the A/X ratio and the substitution pattern of the AX extracts. The fractions buffered at neutral (pH 7.0) and mild alkaline (pH 9.0) conditions at 160 °C exhibited the highest molar masses, comparable to the fractions extracted at 120 °C (Table 2). This demonstrates again the possibility of using pH and temperature to modulate the composition and substitution pattern of the AX fractions as well as the molecular weight.

The eluted fractions after fractionation are mainly composed of oligosaccharides with average molecular weights between 1000–2000 g mol<sup>-1</sup> (Table 2). Arabinoxylan oligosaccharides (AXOs) and β-D-glucan oligosaccharides (BGOs) represent the main oligosaccharide populations, as evidenced by glycosidic linkage analyses (ESI Table S4†). In general the content of BGOs in these low molar mass fractions is enriched compared to the polymeric fractions. The AXOs are generally more substituted than their polymeric counterparts, with an A/X ratio of 0.94 and 0.45–0.65 for the alkaline and SWE fractions, respectively. The oligomeric mass profiling of these low molar mass fractions by ESI-MS showed the presence of BGOs and AXOs predominantly with 4–8 sugar units, although larger oligosaccharides can be detected as well (ESI Fig. S1†). This recovered oligosaccharide fractions could be used as prebiotic agents or platform chemicals, as discussed below.

#### Valorization of the residue by sequential enzymatic treatment and subcritical water extraction

The residues after SWE still exhibited approximately 50% of non-extractable AX with high ferulic acid content (Fig. 2B and C; ESI Table S2†). This crosslinked AX constitutes a valuable resource for further recovery using enzymatic treatments. The residue after extraction at 160 °C pH 7.0 (HR-160-7) was selected to develop the process methodology. In a first attempt, auxiliary enzymes, lichenase and cellulase, that degrade mixed-linked β-D-glucan and cellulose, respectively, were tested in order to release AX populations that could be in close contact with the β-D-glucans trapped in the hydrothermal residue. Although the activity of the auxiliary enzymes was verified by reducing sugar assay (ESI Fig. S2a†), the enzymatic treatments did not result in increased total yields after subsequent SWE (ESI Fig. S2b†). This could be due to the fact that the residual AX is not tightly associated with D-glucans (mixed-linkage β-D-glucan or cellulose), or due to the reduced enzymatic accessibility as a consequence of the presence of FA dehydrodimer crosslinks.<sup>50</sup>

In a further attempt, a feruloyl esterase (FE, from family 1A) and an *endo*-1,4-β-D-xylanase (X, from family GH10) were used alone and in synergy to disrupt the phenolic crosslinked structures of the residue. The FE alone does not show significant activity towards the residue, but its activity increased after 16 h of incubation when combined with the xylanase (Fig. 5a).

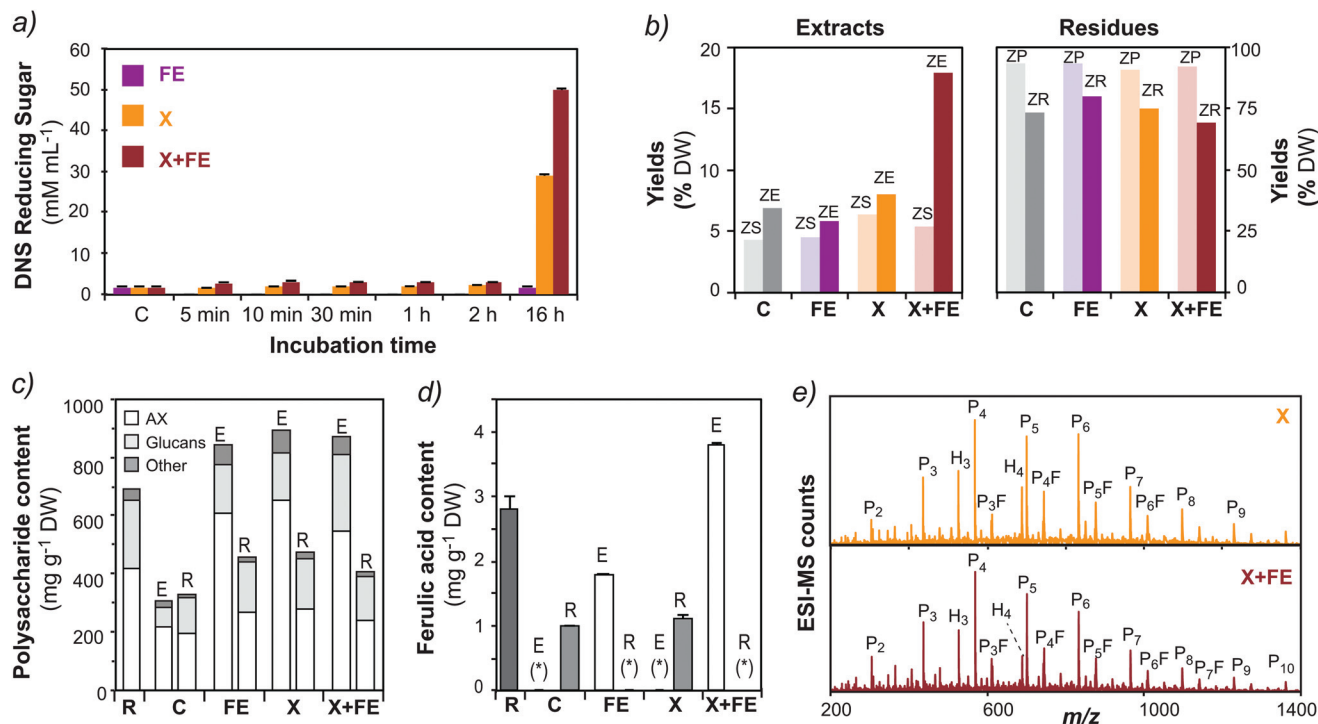
Additionally, the combination of both enzymes results in a significant increase of total solid yields after a subsequent cycle of subcritical water extraction (Fig. 5b). The polysaccharide composition and the phenolic acid content of the fractions after enzymatic treatments followed by SWE were analyzed (Fig. 5c and ESI Table S6†). The extracts after xylanase treatment (X), alone and in combination with feruloyl esterase (X + FE), present a higher A/X ratio (0.9–1.2) in comparison to HE-160-7, indicating the release of highly branched F-AX populations. Ferulic acid could be determined only in the samples that were treated with FE, alone or in combination with X (Fig. 5d).

The polysaccharide composition and the phenolic acid content of the fractions after enzymatic treatments followed by SWE were analyzed (Fig. 5c and ESI Table S6†). The extracts after xylanase treatment (X), alone and in combination with feruloyl esterase (X + FE), present a higher A/X ratio (0.9–1.2) in comparison to HE-160-7, indicating the release of highly branched F-AX populations. Ferulic acid could be determined only in the samples that were treated with FE, alone or in combination with X (Fig. 5d). The oligosaccharide fractions released after the combined enzymatic treatment and SWE were characterized by mass spectrometric techniques. The oligomeric mass profiling of the extracts treated with X and X + FE shows a range of AXOs from 2 to 12 pentose units, together with a profile of F-AXOs containing 4–7 pentose (P) units plus ferulic acid (F) (Fig. 5e). The structural diversity of the released oligosaccharides will be the subject of further investigations. This is a proof of concept of the effectiveness of the enzymatic treatment using xylanolytic enzymes to disrupt the crosslinked feruloylated network and to valorize the non-extractable residue into valuable functional biomolecules. This procedure opens the possibility to test a wide variety of FE from different families, alone or in combination with other xylanolytic enzymes (β-xylanases, α-L-arabinofuranosidases) to release AX populations from the crosslinked residues. Indeed, FEs constitutes a heterogeneous family of carbohydrate esterases (CEs) based on protein homology, their specificity towards phenolic substrates, and the ability to release phenolic dimers.<sup>51–53</sup> Few studies have focused on the specificity and substrate recognition of such FE, alone or in combination with other xylanolytic enzymes, towards complex crosslinked substrates as the ones we encounter in wheat bran.<sup>54</sup> Different enzyme families may show distinct activity and selectivity towards the ferulic crosslinked network, which may allow further optimization of the yields of released biomolecules and tuning of their molecular structures.

#### Antioxidant activities of the AX fractions

The radical scavenging activity of the alkaline and the SWE fractions was assessed against 2,2-diphenyl-1-picrylhydrazyl (DPPH<sup>•</sup>) and 2,2'-azino-bis(3-ethylbenzothiazoline-6-sulphonic acid) (ABTS<sup>•+</sup>) radicals. Both assays are based on electron transfer, involving the reduction of a colored oxidant. Ascorbic acid and ferulic acid were used as positive controls, whereas wheat endosperm AX (WE-AX; Megazyme) was used as a negative control.<sup>55</sup> The radical scavenging activity of selected





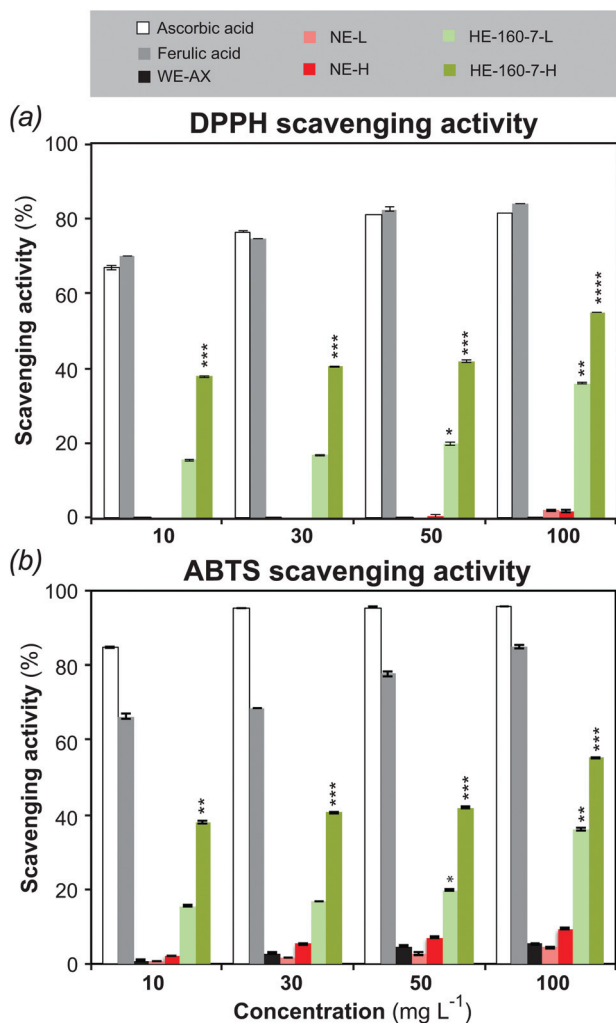
**Fig. 5** Valorization of the crosslinked residue by enzymatic treatments and subsequent subcritical water extraction. (a) Activity of the enzymatic treatments evaluated by reducing sugar assay: C: control, FE: feruloyl esterase; X:  $\beta$ -xyylanase; (b) total yields of the extracts and the residue after enzymatic treatment (ZS: supernatant; ZP: precipitate) and subsequent subcritical water extraction (ZE: extract; ZR: residue). (c) Polysaccharide content and (d) ferulic acid content in the extracts (E) and residues (R). Note: (\*) not detected. (e) Mass profiling of the oligosaccharides released after enzymatic treatments by ESI-MS. Note: H: hexoses; P: pentoses; F: ferulic acid. Peak assignment as reported in ESI Table S5.†

samples is presented in Fig. 6, whereas the results on all the samples together with the half maximal effective concentration ( $EC_{50}$ ) values are presented in the ESI Table S7.†

The alkaline extract did not show statistically significant scavenging activity compared with the controls. Contrarily, both the low molar mass (HE-160-7-L) and the high molar mass (HE-160-7-H) fractions from SWE showed significant and increasing scavenging activity with higher doses, with this effect being more marked in the high molar mass fraction. HE-160-7-H shows the highest scavenging activity against DPPH radicals ( $55.1 \pm 0.06\%$ ) with an  $EC_{50}$  value of  $53.6 \pm 1.34 \mu\text{g ml}^{-1}$  for the highest concentration ( $100 \mu\text{g ml}^{-1}$ ). At the same concentration, the  $EC_{50}$  value for the alkaline fraction NE-H is  $294.9 \pm 1.34 \mu\text{g ml}^{-1}$ , which represents a 6-fold lower potency. For the ABTS<sup>•+</sup> scavenging assay, the whole range of concentrations was tested only for the positive and negative controls (ascorbic acid, ferulic acid and WE-AX) and for the polymeric fractions (H) of the NE and HE-160-7, whereas only the highest concentration ( $100 \mu\text{g ml}^{-1}$ ) was tested for all the different fractions (ESI Table S7†). A good quantitative correlation is obtained between the antioxidant capacities measured by the ABTS and DPPH assays for the positive controls. The qualitative trends between both methods are maintained for the alkaline and SWE extracts, with fraction HE-160-7-H showing the highest ABTS scavenging activity (Fig. 6b).

As the commercial wheat endosperm AX and the alkaline fractions did not show significant scavenging activity in both assays, the ferulic acid present in the SWEs can be considered as the main contributor to the antioxidant properties.<sup>46,56–59</sup> Phenolic compounds are reported to have strong antioxidant activity that can be associated to their ability to destroy oxygen-derived free radicals, break radical chain reaction and/or chelate metals.<sup>60–62</sup> The DPPH free radical scavenging of FA probably occurs due to its hydrogen-donating ability.<sup>56,61</sup> The DPPH or any reactive radical colliding with FA phenolic nucleus can easily abstract a hydrogen atom to form a resonance-stabilized phenoxy radical, accounting for its antioxidant activity.<sup>59</sup> Many publications have reported the antioxidant activity of free FA or conducted studies to evaluate the free radical scavenging capacity of wheat, wheat bran or other cereal extractives and oligosaccharides.<sup>47,61,63–66</sup> However, just a handful number of studies have demonstrated the scavenging activity of isolated feruloylated AXs in polymeric form. Hydroxycinnamic acid bound AXs extracted using mild alkali conditions from different sources of millet bran,<sup>67</sup> maize fiber<sup>59</sup> and sorghum bran<sup>60</sup> showed antioxidant activity. These extractions exhibit a lower total phenolic content (between  $1\text{--}2.4 \text{ mg g}^{-1}$ ) resulting in a lower potency of the scavenging activity when compared to the high molar mass subcritical water extracts obtained in the present study. The high efficiency of the polymeric AX fractions with bound ferulic





**Fig. 6** Radical scavenging activity of the alkaline and SWE high and low molar mass fractions from wheat bran. (a) DPPH (b) ABTS. Ascorbic acid and ferulic acid were used as positive control; wheat endosperm arabinoxylan (WE-AX) was used as negative control. Values were expressed as mean  $\pm$  SD of triplicate measurements. Asterisks indicate a significant difference (\* $P \leq 0.05$ ; \*\* $P \leq 0.01$ ; \*\*\* $P \leq 0.001$ ; \*\*\*\* $P \leq 0.005$ ) relative to control.

acid as radical scavenging agents aligns with a recent re-evaluation of the antioxidant activity of complex carbohydrates, which indicate the important role of minor phenolic components in the antioxidant properties.<sup>55</sup> However, the nature of the correlation between ferulic acid content and antioxidant properties is still not fully understood. Further studies of the antioxidant activity of F-AXs with a range of ferulic acid content should cast more light on this issue.

#### Final mass balances and potential of the hemicellulose and oligosaccharide fractions from wheat bran

The proposed cascade bioprocess generates enriched hemicellulose polymeric populations (mainly F-AX) and oligosaccharides from wheat bran. Table 3 presents the total mass

balance of the designed process in comparison with the alkaline extraction, in terms of total solid, total AX and ferulic acid yields. The proposed bioprocess offers higher total yields and similar AX content compared to traditional alkaline extraction. As an advantage, it only uses mild aqueous solvents combined with biocatalysis (carbohydrate-active enzymes) preserving the phenolic substitutions (ferulic acid) attached to the polymeric and oligomeric hemicellulose fractions. The phenolic functionalities confer valuable antioxidant and radical scavenging properties to these fractions, which may be extremely valuable for food applications and in the design of functional materials and platform chemicals.

Arabinoxylans and their derived AXOs are important from a nutritional point of view, since they both have been associated with a great variety of health benefits as dietary fibers, including prebiotic and immunomodulatory activities.<sup>58</sup> The mechanisms of dietary fiber polysaccharides, and of arabinoxylans in particular, in modulating the gut microbiome are starting to be understood.<sup>68,69</sup> However, further research is required to better correlate the fine structure–function relationships that regulate the health-promoting mechanisms of dietary fibers.<sup>70</sup> From a technological point of view, AXs have shown great potential as biomaterials due to their biocompatibility, biodegradability, and barrier properties to oxygen.<sup>71,72</sup> The extracted polymeric F-AX fractions show excellent potential for the design of bio-based materials, combining the good rheological and viscoelastic properties of the polysaccharide core with the inherent antioxidant properties of the phenolic moieties. Moreover, the FA could offer further interesting functionalities, such as UV absorbance<sup>73</sup> and the possibility for enzymatic crosslinking.<sup>74,75</sup> This opens new routes for the development of multifunctional films, hydrogels, and texturizing agents to be used in food packaging, drug delivery, and other biomedical applications. These inherent radical scavenging materials may contribute to an extended shelf life of unstable foods and drugs (preventing oxidation of less stable components) and to mitigate the presence of reactive oxygen species (ROS) in inflammatory processes.<sup>76</sup>

**Table 3** Complete mass balances in terms of total solid yields (as % of initial dry weight wb), total AX yields (as % of the initial AX content in wb) and as total FA yields (as % of the initial FA content in wb)

Fraction	Total yields (%)		AX yields (%)		FA yields (%)	
	NE	HE-160-7	NE	HE-160-7	NE	HE-160-7
H	24.8	14.7	68.6	42.9	2.3	32.7
L	2.9	7.3	4.1	12.2	0.3	7.3
ZS + ZE	n.d.	9.2	n.d.	17.3	n.d.	15.8
<b>Total</b>	<b>27.6</b>	<b>31.2</b>	<b>72.6</b>	<b>72.3</b>	<b>2.5</b>	<b>55.9</b>

NE: alkaline Na(OH) extraction; HE-160-7: optimized SWE at 160 °C and pH 7; H: high molar mass fraction; L: low molar mass fraction; EZ + SWE: fraction after enzymatic treatment (FE + X) and subsequent SWE at 160 °C pH 7.



## Conclusions

We have designed a cascade bioprocess using subcritical water extraction, membrane filtration, and enzymatic treatments for the isolation of functional biomolecules from cereal byproducts. With this approach, we do not only fractionate hemicelluloses with high molecular weight and intact phenolic functionalities, but we also produce valuable oligosaccharides from the non-extractable residues. The use of both subcritical water and xylanolytic enzymes offers evident environmental benefits as green solvents and biocatalysts avoiding the use of hazardous substances. The purity, substitution pattern, and molecular weight of the polymeric extracts can be modulated by controlling the temperature and pH conditions during subcritical water extraction. The SW extracts show significant radical scavenging activity compared to the alkaline extracts, which can be attributed to the presence of ferulic acid attached to the carbohydrate backbones. Moreover, xylanolytic enzymes release functional oligosaccharides from the unextractable residue, increasing the total AX and phenolic yields. This study opens up the possibility of exploring the use of different families of xylan-degrading enzymes to disrupt the recalcitrant crosslinked network, in order to maximize and tune the yields, molecular structure and properties of the released oligosaccharides. These polysaccharide and oligosaccharide fractions show great potential for their utilization in advanced nutritional (prebiotic) and material applications in the food and biomedical sectors, combining the macroscopic properties of the carbohydrates with the antioxidant activity of the ferulic moieties. This sequential procedure can be extended to basically any cereal side stream and opens the possibility to an improved utilization of plant biomass.

## Experimental

### Materials

Fine granulometry defatted wheat bran (wb) was provided by Lantmännen (Stockholm, Sweden). All chemicals, analytical standards and reagents were from Sigma-Aldrich (Stockholm Sweden). Folin Ciocalteu Phenol reagent was from Merck (Solna, Sweden). Spectra/Por 1 and 3 Dialysis Membrane, 6–8 kDa and 3.5 kDa MWCO, respectively and Biotech CE dialysis tubing, 0.1–0.5 kDa and 20 kDa MWCO were purchased from SpectrumLabs (Breda, The Netherlands). Centramate cassette 30 kDa MWCO was purchased from VWR. Feruloyl esterase 1A (*Ruminococcus albus*) was purchased from NZytech (Lisbon, Portugal). *endo*-1,4- $\beta$ -Xylanase (*Cellvibrio mixtus*), cellulase (*Talaromyces emersonii*) and lichenase (*Bacillus subtilis*), and wheat endosperm high viscosity arabinoxylan were purchased from Megazyme (Wicklow, Ireland).

### Bioprocess design

**Destarching.** The bran was enzymatically destarched with  $\alpha$ -amylase. In brief, the bran was suspended in 0.01 M phosphate buffer with 100 mM KCl (pH 7.5) in a ratio of 1 : 10 (w/v)

and boiled for 5 min to gelatinize the starch granules. The samples were equilibrated at 40 °C and incubated with  $\alpha$ -amylase (2 U mg<sup>-1</sup> of carbohydrate) for 1 h at 40 °C; a second incubation was performed for further 30 min. The polysaccharides were precipitated by addition of four volumes of cold absolute ethanol at –8 °C overnight and centrifuged (1500g, 5 min). The precipitate was washed with cold absolute ethanol ( $\times 3$ ) and water, centrifuging between washes, prior to freeze-drying. Starch removal was verified under the microscope by iodine staining. The total starch content was determined gravimetrically and enzymatically before and after the  $\alpha$ -amylase treatment.

**Alkaline extraction.** Destarched wheat bran (Dwb) was submitted to alkaline extraction by 0.5 M NaOH or Ca(OH)<sub>2</sub> solutions at 80 °C for 16 h using a ratio of 1 : 8 (w/v). The recovered supernatants (NE and CE) and the residues (NR and CR) were neutralized with acetic acid, dialyzed for 48 h using a 3.5 kDa MWCO dialysis membrane and freeze-dried.

**Subcritical water extraction.** Subcritical water extraction (SWE) of Dwb was performed by pressurized hot water using a laboratory accelerated solvent extraction Dionex™ ASE™ 350. Four different temperatures (100, 120, 140 and 160 °C) were tested at pH 5.0, whereas two additional pH (pH 7.0 and 9.0) were also tested at 160 °C. The extraction aqueous solvents were buffered with 0.1 M sodium formate. The pH was adjusted either by addition of formic acid or NaOH and again measured in the final extracts to check for changes. The extractions were performed using 1 g of Dwb under a static cycle of 15 min, resulting in a hydrothermal extract (HE) and a hydrothermal residue (HR) for each one of the tested conditions. The extracts and residues were dialyzed with a 3.5 kDa MWCO membrane for 72 h and freeze-dried.

**Fractionation of high and low molar mass fractions.** The different extracts after SWE were resuspended in water and submitted to closed dialysis using a 20 kDa MWCO membrane for 48 h. The collected eluents and retained fractions were concentrated and freeze-dried. Membrane filtration was upscaled for HE-160-7 in a continuous flow (initial concentration 10 mg mL<sup>-1</sup>), using a 30 kDa cassette membrane ultrafiltration device. The retentate (high molar mass; H) and the eluent (low molar mass; L), were collected separately, concentrated and freeze-dried.

**Enzymatic treatments of the residues.** Two enzymatic approaches were tested in order to further valorize the residue. In both cases, the hydrothermal residues were incubated with the enzymes, centrifuged, freeze-dried and further submitted to a new SWE at 160 °C pH 7.0 for 15 min. The first approach involved the use of auxiliary enzymes (lichenase and/or cellulase) to assist the relaxation of bran residue material. Briefly, 1 g of the hydrothermal residue HR-160-7 was treated with lichenase (30  $\mu$ L, 1000 U mL<sup>-1</sup>; 40 °C, 16 h), cellulase (30  $\mu$ L, 700 U mL<sup>-1</sup>; 40 °C, 16 h) or a combination of both enzymes, in 20 mL of the appropriate buffer (sodium phosphate pH 6.5, 4.5 or 5.0, respectively). The second approach involved the use of feruloyl esterase 1A (FE) alone or in synergy with an *endo*-1,4- $\beta$ -D-xylanase (X). The incubations were done in water



(20 mL) at 37 °C for 16 h. The supernatants generated after the enzymatic incubations were used for the quantification of the reducing sugars (DNS assay).

### Characterization of the extracted fractions

**Fluorescence microscopy.** The samples were visualized under a fluorescence microscope (Axio Imager M2; Carl Zeiss, Oberkochen, Germany) using the reflective index of differential interference contrast (DIC). Phenolic acid auto-fluorescence signals were observed under the DAPI filter set at 365 nm excitation and 445/50 nm emission wavelengths. All images were captured using an AxioCam HRc camera (Carl Zeiss, Oberkochen, Germany).

**Monosaccharide composition.** The monosaccharide composition was analyzed by acid hydrolysis followed by high-pH anion exchange chromatography with pulsed amperometric detection (HPAEC-PAD) and GC-MS analysis (triplicate runs). Two-step sulfuric acid hydrolysis (72% H<sub>2</sub>SO<sub>4</sub>, 3 h, R.T., followed by 1 M H<sub>2</sub>SO<sub>4</sub>, 100 °C, 3 h) was performed for the original bran (wb) and the residues (NR and HR) using the conditions originally reported by Saeman.<sup>77</sup> TFA hydrolysis (2 M TFA at 121 °C for 3 h) was performed on the extracts using 1–5 mg of freeze-dried material. The hydrolyzed monosaccharides were analyzed by HPAEC-PAD on an ICS3000 system (Dionex, Sunnyvale, CA) using a Dionex CarboPac PA1 column at 30 °C at a flow rate of 1 mL min<sup>-1</sup>. Different gradients were employed for the quantification of neutral monosaccharides and uronic acids.<sup>28</sup> For the GC-MS analyses, the remaining hydrolyzed samples were dried until complete acid removal, reduced with sodium borohydride (NaBH<sub>4</sub>) at 60 °C for 1 h and acetylated with pyridine and acetic anhydride (1 : 1 v/v, 100 μL) at 100 °C for 30 min.<sup>78</sup> The alditol acetates were extracted with ethyl acetate and analyzed by GC-MS using a HP-6890 gas chromatographer and an HP-5973 electron-impact mass spectrometer (Agilent Technologies, Santa Clara, CA).

**Protein content and composition.** The total protein content was determined using the Dumas method.<sup>39</sup> In parallel, the amino acid composition was analysed by GC-MS after acid hydrolysis (6 M HCl with 1% phenol (v/v), 100 °C, 20 h) and derivatization (pyridine–methanol–acetic anhydride (1 : 1 : 4 v/v), 100 °C, 60 min) into amino acid methyl esters (aAMAs).<sup>79</sup>

**Klason lignin** was determined gravimetrically from the residues after two-step total sulfuric acid hydrolysis (72 wt% H<sub>2</sub>SO<sub>4</sub>, 3 h, R.T.; 1 M H<sub>2</sub>SO<sub>4</sub>, 100 °C, 3 h).<sup>80</sup>

**Hydroxycinnamic acid quantification.** The total hydroxycinnamic acid profile was determined as described by Comino *et al.*,<sup>33</sup> with minor modifications. In brief, 10 mg of dry sample (in duplicates) were saponified with 500 μL of 2 M NaOH overnight at room temperature under stirring in the dark and acidified with 12 M HCl (to pH 3.0) with an internal standard (3,4-dimethoxycinnamic acid – 5 μg). The samples were then extracted three times with 1 mL ethyl acetate, dried, and silylated with 50 μL N-O-bis(trimethylsilyl)acetamide) at 100 °C for 5 min. The samples were resuspended in acetone and injected onto a GC-MS fitted with a CP Sil 5CB column (Agilent Technologies, Santa Clara, CA).

**Acetyl content and degree of acetylation.** 10 g of wheat bran (wb and Dwb) were hydrolyzed with NaOH at 70 °C overnight. The acetyl content was determined from the extracts using high-performance liquid chromatography (HPLC) with UV detection. The degree of acetylation (DA) was determined according to Xu *et al.*<sup>81</sup>

**Phytic acid content.** The phytic acid content was determined using the phytic acid (total phosphorous) assay kit from Megazyme, according to manufacturers instructions.

**Glycosidic linkage analysis.** Per-*O*-methylation was carried out as described by Pettolino *et al.*,<sup>82</sup> with some modifications. The samples (1 mg) were dissolved in dimethyl sulfoxide (400 μL) with NaOH (20 mg). Methyl iodide was added (5 × 50 μL) under continuous stirring. The methylated polysaccharides were recovered in the organic phase after partition (×3) with H<sub>2</sub>O and dichloromethane (DCM) and dried. The samples were hydrolyzed with 2 M TFA at 121 °C for 3 h, dried, reduced with NaBD<sub>4</sub> at 60 °C for 1 h, and acetylated with pyridine and acetic anhydride (1 : 1 v/v, 200 μL) at 100 °C for 30 min. The per-*O*-methylated alditol acetates were extracted with ethyl acetate and quantified in triplicate by GC-MS, by comparison with typical polysaccharide standards.

**Molar mass distributions by SEC.** The molar mass distributions of the extracted fractions were analyzed by size-exclusion chromatography (SECcurity 1260, Polymer Standard Services, Mainz, Germany) coupled to a refractive index detector (SECcurity 1260, Polymer Standard Services, Mainz, Germany) at 45 °C. The extracted fractions were dissolved directly in the SEC eluent consisting of dimethyl sulfoxide (DMSO, HPLC grade, Scharlab, Sweden) with 0.5% w/w LiBr (ReagentPlus) at 60 °C. SEC analyses were performed with a flow rate of 0.5 mL min<sup>-1</sup> at 60 °C using a column set consisting of a GRAM PreColumn, 30 and 10 000 analytical columns (PSS, Mainz, Germany). Calibration of the SEC separation was performed using pullulan standards provided by Polymer Standards Services (PSS, Mainz, Germany).

**Dinitrosalicylic acid (DNS) method – reducing sugar determination.** 1 mL of sample was mixed with 1 mL of DNS reagent, vortexed and incubated in a boiling water bath for 15 min.<sup>83</sup> After cooling, the sample was diluted to 10 ml and the absorbance measured using a spectrophotometer at 540 nm against a reagent blank. Calibration was performed using glucose standard solutions.

**Electrospray ionization mass spectrometry (ESI-MS).** Positive-ion ESI-MS was carried out on a Q-TOF<sup>2</sup> ESI mass spectrometer (Micromass, UK). Samples were dissolved in acetonitrile (50%) containing 0.1% formic acid and infused directly into the Q-TOF through the capillary liquid chromatography (CapLC) module at a rate of 8 μL min<sup>-1</sup>. The ESI source was operated at 3.3 kV with a desolvation temperature of 140 °C and a cone voltage of 70–80 V. A collision energy of 40–60 eV was applied for the fragmentation of the selected oligosaccharides (ESI-MS/MS) using argon as collision gas. The oligosaccharides were detected as [M + Na]<sup>+</sup> adducts.

**DPPH scavenging activity.** The scavenging activity of the wheat bran extracts on DPPH radicals was measured in tripli-



cate according to the method of Brand-Williams *et al.*,<sup>84</sup> with minor modifications. Ascorbic acid and ferulic acid were used as reference materials. Briefly, an aliquot of 1.5 mL of sample aqueous solution at different concentrations (10, 30, 50 and 100  $\mu\text{g mL}^{-1}$ ) was mixed with 0.5 mL of 100  $\mu\text{M L}^{-1}$  methanolic solution of DPPH $\cdot$ . The blank control contained all the reaction reagents and methanol instead of the wheat bran extracts or ascorbic and ferulic acid. The reaction mixture was incubated for 30 min in the dark at room temperature, and the absorbance of the resulting solution was measured at 517 nm. The radical scavenging activity of the tested samples was measured as a decrease in the absorbance of DPPH $\cdot$ .

**ABTS<sup>+</sup> scavenging activity.** The scavenging activity of the wheat bran extracts on ABTS radical cation (ABTS<sup>+</sup>) was measured in triplicate according to the method described by Re *et al.*,<sup>85</sup> with minor modifications. Ascorbic acid and ferulic acid were used as reference materials. Briefly, an aliquot of 10  $\mu\text{L}$  of sample aqueous solution at different concentrations (10, 30, 50 and 100  $\mu\text{g mL}^{-1}$ ) was mixed with 1.0 mL of a dilution of ABTS<sup>+</sup> in ethanol to an absorbance of  $0.70 \pm 0.02$  nm at 734 nm. The blank control contained all the reaction reagents and ethanol instead of the wheat bran extracts or ascorbic and ferulic acid. The reaction mixture was incubated for 5 min in the dark at room temperature, and the absorbance of the resulting solution was measured at 734 nm. The radical scavenging activity of the tested samples was measured as a decrease in the absorbance of ABTS<sup>+</sup>.

## Acknowledgements

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## References

- O. O. Onipe, A. I. O. Jideani and D. Beswa, *Int. J. Food Sci. Technol.*, 2015, **50**, 2509–2518.
- L. Stevenson, F. Phillips, K. O'Sullivan and J. Walton, *Int. J. Food Sci. Nutr.*, 2012, **63**, 1001–1013.
- S. Apprich, Ö. Tirpanalan, J. Hell, M. Reisinger, S. Böhrmdorfer, S. Siebenhandl-Ehn, S. Novalin and W. Kneifel, *LWT – Food Sci. Technol.*, 2014, **56**, 222–231.
- J. Hell, L. Donaldson, H. Michlmayr, M. Kraler, W. Kneifel, A. Potthast, T. Rosenau and S. Böhrmdorfer, *Carbohydr. Polym.*, 2015, **121**, 18–26.
- M. S. Izydorczyk and C. G. Biliaderis, *Carbohydr. Polym.*, 1995, **28**, 33–48.
- L. Saulnier, F. Guillon, P.-E. Sado and X. Rouau, in *Comprehensive Glycoscience*, ed. H. Kamerling, Elsevier, France, 2007, pp. 653–689, DOI: 10.1016/B978-044451967-2/00147-1.
- A. Ebringerova, Z. Hromadkova and T. Heinze, in *Polysaccharides I*, ed. T. Heinze, 2005, ch. 136816, vol. 186, pp. 1–67.
- M. Aguedo, C. Fougnyes, M. Dermience and A. Richel, *Carbohydr. Polym.*, 2014, **105**, 317–324.
- I. Egüés, A. M. Stepan, A. Eceiza, G. Toriz, P. Gatenholm and J. Labidi, *Carbohydr. Polym.*, 2014, **102**, 12–20.
- Z. Zhang, C. Smith and W. Li, *Food Res. Int.*, 2014, **65**, 423–436.
- A. Höije, M. Gröndahl, K. Tommeraas and P. Gatenholm, *Carbohydr. Polym.*, 2005, **61**, 266–275.
- D. J. Rose and G. E. Inglett, *Food Chem.*, 2010, **119**, 1613–1618.
- A. Ebringerová, Z. Hromádková, J. Alfödi and V. Hribalová, *Carbohydr. Polym.*, 1998, **37**, 231–239.
- Z. Hromádková, J. Kováčiková and A. Ebringerová, *Ind. Crops Prod.*, 1999, **9**, 101–109.
- S. G. Allen, D. Schulman, J. Lichwa, M. J. J. Antal, M. Laser and L. R. Lynd, *Ind. Eng. Chem. Res.*, 2001, **40**, 2934–2941.
- Ó. Benito-Róman, E. Alonso and M. J. Cocero, presented in part at the III Iberoamerican Conference on Supercritical Fluids, Cartagena de Indias, 2013.
- Ó. Benito-Román, E. Alonso, M. J. Cocero and M. Goto, *Food Bioprod. Process.*, 2016, **98**, 21–28.
- A. Mustafa and C. Turner, *Anal. Chim. Acta*, 2011, **703**, 8–18.
- C. C. Teo, S. N. Tan, J. W. H. Yong, C. S. Hew and E. S. Ong, *J. Chromatogr., A*, 2010, **1217**, 2484–2494.
- O. Hosseinaei, S. Wang, T. G. Rials, C. Xing and Y. Zhang, *Cellulose*, 2011, **18**, 841–850.
- J. V. Rissanen, H. Grénaman, C. Xu, S. Willför, D. Y. Murzin and T. Salmi, *ChemSusChem*, 2014, **7**, 2947–2953.
- T. Song, A. Pranovich and B. Holmbom, *Holzforschung*, 2011, **65**, 35–42.
- P. O. Kilpeläinen, S. S. Hautala, O. O. Byman, L. J. Tanner, R. I. Korpinen, M. K. J. Lillandt, A. V. Pranovich, V. H. Kitunen, S. M. Willför and H. S. Ilvesniemi, *Green Chem.*, 2014, **16**, 3186–3194.
- A. Martínez-Abad, A. C. Ruthes and F. Vilaplana, *J. Appl. Polym. Sci.*, 2016, **133**, 42523.
- R. Deutschmann and R. F. H. Dekker, *Biotechnol. Adv.*, 2012, **30**, 1627–1640.
- R. Faryar, J. A. Linares-Pastén, P. Immerzeel, G. Mamo, M. Andersson, H. Stålbrand, B. Mattiasson and E. N. Karlsson, *Food Bioprod. Process.*, 2015, **93**, 1–10.
- I. Finore, A. Poli, P. Di Donato, L. Lama, A. Trincone, M. Fagnano, M. Mori, B. Nicolaus and A. Tramice, *Green Chem.*, 2016, **18**, 2460–2472.
- L. S. McKee, H. Sunner, G. E. Anasontzis, G. Toriz, P. Gatenholm, V. Bulone, F. Vilaplana and L. Olsson, *Biotechnol. Biofuels*, 2016, **9**, 1–13.
- E. Dornez, K. Gerbruers, S. Wiame, J. A. Delcour and C. M. Courtin, *J. Agric. Food Chem.*, 2006, **54**, 8521–8529.
- Y. Hemery, X. Rouau, V. Lullien-Pellerin, C. Barron and J. Abecassis, *J. Cereal Sci.*, 2007, **46**, 327–347.
- G. Di Lena, V. Vivanti and G. B. Quaglia, *Food*, 1997, **41**, 285–288.
- M. Reisinger, Ö. Tirpanalan, M. Prückler, F. Huber, W. Kneifel and S. Novalin, *Bioresour. Technol.*, 2013, **144**, 179–185.
- P. Comino, H. Collins, J. Lahnstein, C. Beahan and M. J. Gidley, *Food Hydrocolloids*, 2014, **41**, 219–226.



- 34 S. Zhou, X. Liu, Y. Guo, Q. Wang, D. Peng and L. Cao, *Carbohydr. Polym.*, 2010, **81**, 784–789.
- 35 A. Fadet, *Nutr. Res. Rev.*, 2010, **23**, 65–134.
- 36 C. Barron, A. Surget and X. Rouau, *J. Cereal Sci.*, 2007, **45**, 88–96.
- 37 K. Gebruers, E. Dornez, D. Boros, A. Fras, W. Dynkowska, Z. Bedő, M. Rakszegi, J. A. Delcour and C. M. Courtin, *J. Agric. Food Chem.*, 2008, **56**, 9740–9749.
- 38 M. Dubois, K. A. Gilles, J. K. Hamilton, P. A. Rebers and F. Smith, *Anal. Chem.*, 1956, **28**, 350–356.
- 39 J. B. A. Dumas, *Ann. Chim. Phys.*, 1831, **2**, 198–213.
- 40 M. E. F. Schooneveld-Bergmans, A. M. C. P. Hopman, G. Beldman and A. G. J. Voragen, *Carbohydr. Polym.*, 1998, **35**, 39–47.
- 41 G. B. Fincher, *J. Inst. Brew.*, 1976, **82**, 347–349.
- 42 H. K. Lichtenthaler and J. Schweiger, *J. Plant Physiol.*, 1998, **152**, 272–282.
- 43 V. Pussayanawin, D. L. Wetzell and R. G. Fulcher, *J. Agric. Food Chem.*, 1988, **36**, 515–520.
- 44 B. Albinsson, S. Li, K. Lundqvist and R. Stomberg, *J. Mol. Struct.*, 1999, **508**, 1–3.
- 45 L. Donaldson, *IAWA J.*, 2013, **34**, 3–19.
- 46 B. Laddomada, S. Caretto and G. Mita, *Molecules*, 2015, **20**, 15666–15685.
- 47 X. Yuan, J. Wang and H. Yao, *Food Chem.*, 2005, **90**, 759–764.
- 48 Y. Sun, S. W. Cui, X. Gu and J. Zhang, *Carbohydr. Polym.*, 2011, **85**, 615–621.
- 49 B. Domon and C. E. Costello, *Glycoconjugate J.*, 1988, **5**, 397–409.
- 50 D. M. de Oliveira, A. Finger-Teixeira, T. R. Mota, V. H. Salvador, F. C. Moreira-Vilar, H. B. C. Molinari, R. A. C. Mitchell, R. Mearchiosi, O. Ferrarese-Filho and W. D. dos Santos, *Plant Biotechnol. J.*, 2015, **13**, 1224–1232.
- 51 V. F. Crepin, C. B. Faulds and I. F. Connerton, *Appl. Microbiol. Biotechnol.*, 2004, **63**, 647–652.
- 52 I. Benoit, E. G. J. Danchin, R.-J. Bleichrodt and R. P. de Vries, *Biotechnol. Lett.*, 2008, **30**, 387–396.
- 53 D. B. R. K. G. Udatha, I. Kouskoumvekaki, L. Olsson and G. Panagiotou, *Biotechnol. Adv.*, 2011, **29**, 94–110.
- 54 C. B. Faulds, D. Zanichelli, V. F. Crepin, I. F. Connerton, N. Juge, M. K. Bhat and K. W. Waldron, *J. Cereal Sci.*, 2003, **38**, 281–288.
- 55 S. Hu, J. Yin, S. Nie, J. Wang, G. O. Phillips, M. Xie and S. W. Cui, *Bioact. Carbohydr. Diet. Fibre*, 2016, **7**, 19–27.
- 56 H. Kikuzaki, M. Hisamoto, K. Hirose, K. Akiyama and H. Taniguchi, *J. Agric. Food Chem.*, 2002, **50**, 2161–2168.
- 57 K. Masisi, T. Beta and M. H. Moghadasian, *Food Chem.*, 2016, **196**, 90–97.
- 58 M. Mendis and S. Simsek, *Food Hydrocolloids*, 2014, **42**, 239–243.
- 59 M. Srinivasan, A. R. Sudheer and V. P. Menon, *J. Clin. Biochem. Nutr.*, 2007, **40**, 92–100.
- 60 G. G. Duthie, S. J. Duthie and J. A. M. Kyle, *Nutr. Res. Rev.*, 2000, **13**, 79–106.
- 61 J. Wang, X. Yuan, B. Sun, Y. Tian and Y. Cao, *Food Technol. Biotechnol.*, 2009, **47**, 39–46.
- 62 S. Itagaki, T. Kurokawa, C. Nakata, Y. Saito, S. Oikawa, M. Kobayashi, T. Hirano and K. Iseki, *Food Chem.*, 2009, **114**, 466–471.
- 63 P. Katapodis, M. Vardakou, E. Kalogeris, D. Kekos, B. J. Macris and P. Christakopoulos, *Eur. J. Nutr.*, 2003, **42**, 55–60.
- 64 A. Mpofo, H. D. Spirstein and T. Beta, *J. Agric. Food Chem.*, 2006, **54**, 1265–1270.
- 65 D. Povilaitis, V. Sulniute, P. R. Venskutonis and V. Kraujaliene, *J. Cereal Sci.*, 2015, **62**, 117–123.
- 66 B. R. Veenashri and G. Muralikrishna, *Food Chem.*, 2011, **126**, 1475–1481.
- 67 V. Bijalwan, U. Ali, A. K. Kesarwani, K. Yadav and K. Mazunder, *Int. J. Biol. Macromol.*, 2016, **88**, 296–305.
- 68 A. Rogowski, J. A. Briggs, J. C. Mortimer, T. Tryfona, N. Terrapon, E. C. Lowe, A. Basle, C. Morland, A. M. Day, H. Zheng, T. E. Rogers, P. Thompson, A. R. Hawkins, M. P. Yadav, B. Henrissat, E. C. Martens, P. Dupree, H. J. Gilbert and D. N. Bolam, *Nat. Commun.*, 2015, **6**, 7481.
- 69 E. C. Martens, A. G. Kelly, A. S. Tauzin and H. Brumer, *J. Mol. Biol.*, 2014, **426**, 3851–3865.
- 70 B. R. Hamaker and Y. E. Tuncil, *J. Mol. Biol.*, 2014, **426**, 3838–3850.
- 71 G.-G. Chen, X.-M. Qi, Y. Guan, F. Peng, C.-L. Yao and R.-C. Sun, *ACS Sustainable Chem. Eng.*, 2016, **4**, 1985–1993.
- 72 M. J. Costa, M. A. Cerqueira, H. A. Ruiz, C. Fougnyes, A. Richel, A. A. Vicente, J. A. Teixeira and M. Aguedo, *Ind. Crops Prod.*, 2015, **66**, 305–311.
- 73 N. Kumar and V. Pruthi, *Biotechnol. Rep.*, 2014, **4**, 86–93.
- 74 G. Oudgenoeg, R. Hilhorst, S. R. Piersma, C. G. Boeriu, H. Gruppen, M. Helsing, A. G. J. Voragen and C. Laane, *J. Agric. Food Chem.*, 2001, **49**, 2503–2510.
- 75 E. Carvajal-Millan, V. Landillon, M.-H. Morel, X. Rouau, J.-L. Doublier and V. Micard, *Biomacromolecules*, 2005, **6**, 309–317.
- 76 Y.-H. Cheng, S.-H. Yang and F.-H. Lin, *Biomaterials*, 2011, **32**, 6953–6961.
- 77 J. F. Saeman, W. E. Moore, R. L. Mitchell and M. A. Millett, *Tappi J.*, 1954, **37**, 336–343.
- 78 A. B. Blakeney, P. J. Harris, R. J. Henry and B. A. Stone, *Carbohydr. Res.*, 1983, **113**, 291–299.
- 79 G. L. Sasaki, L. M. Souza, R. V. Serrato, T. R. Cipriani, P. A. J. Gorin and M. Iacomini, *J. Chromatogr., A*, 2008, **1208**, 215–222.
- 80 J. B. Sluiter, R. O. Ruiz, C. J. Scarlata, A. D. Sluiter and D. W. Templeton, *J. Agric. Food Chem.*, 2010, **58**, 9043–9053.
- 81 C. Xu, A. S. Leppänen, P. Eklund, P. Holmlund, R. Sjöholm, K. Sundberg and S. Willför, *Carbohydr. Res.*, 2010, **345**, 810–816.
- 82 F. A. Pettolino, C. Walsh, G. B. Fincher and A. Basic, *Nat. Protoc.*, 2012, **7**, 1590–1606.
- 83 G. L. Miller, *Anal. Chem.*, 1959, **31**, 426–428.
- 84 W. Brand-Williams, M. E. Cuvelier and C. Berset, *LWT – Food Sci. Technol.*, 1995, **28**, 25–30.
- 85 R. Re, N. Pellegrini, A. Proteggente, A. Pannala, M. Yang and C. Rice-Evans, *Free Radical Biol. Med.*, 1999, **26**, 1231–1237.

