

1 **Changes in relative molecular weight distribution of soluble barley beta-glucan during**  
2 **passage through the small intestine of pigs**

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17

18 **Abstract**

19 The relative molecular weight distribution of soluble barley beta-glucans (SBB) was  
20 monitored through the small intestine in pigs by analyzing water extracts of duodenal- and  
21 ileal digesta with HPLC-SEC. Variations among four diets, based on four different barley  
22 varieties, were documented as well as variations between animals fed the same diet. The  
23 results showed depolymerisation of the SBB throughout the whole small intestine  
24 independent of diet. The average molecular weight of the SBB was reduced to approximately  
25 50% in duodenum in all the experimental animals.

26

27

28 **Key words:** beta-glucan; depolymerisation; relative molecular weight distribution; pigs;  
29 small intestine

30

31

## 32        **1. Introduction**

33        Dietary fiber will affect digestive physiology in pigs and influence digesta flow, voluntary  
34        feed intake and thus nutritional absorption and feed digestibility (Bach Knudsen, Hedemann  
35        et al. 2012), in addition to manure odor and ammonia emissions (O'Shea, Gahan et al. 2010).  
36        Thus, different factors such as grain type and their chemical composition as well as cereal  
37        derived endogenous enzyme activities will affect gastrointestinal function, bacteria population  
38        and microbial metabolites in the gut (Högberg and Lindberg 2004; Högberg, Lindberg et al.  
39        2004; Bindelle J., Leterme P. et al. 2008; Pieper, Jha et al. 2008). These effects will further  
40        depend on the size, solubility and molecular structure of the dietary fiber (Bach Knudsen,  
41        Jensen et al. 1993; Glitsø, Brunsgaard et al. 1998; Bach Knudsen, Hedemann et al. 2012).

42                Dietary fiber, here/often referred to as non-starch polysaccharides (NSP), is  
43        depolymerized in the gastrointestinal (GI) tract in different biological systems (Bach Knudsen  
44        and Canibe 2000; Coles, Moughan et al. 2005). It is evident that cereal beta-glucans are  
45        digested in the upper GI tract of pigs at various degrees, and especially in the distal part of the  
46        small intestine (ileum). Digestibility of the cereal beta-glucans will depend on different  
47        factors; not only particle size or the feed matrix is important, but also source of beta-glucan  
48        and diet composition. Also different grain types and varieties with parallel variation in the  
49        fiber content, as well as different biological systems and individual biological differences  
50        between subjects will influence the monitored experimental results. However, not only  
51        digestibility is important, but physiological properties of beta-glucans are also significant for  
52        both animal nutrition and health. Despite different reports on digestion of cereal beta-glucans  
53        based on quantitative recovery (Fadel, Newman et al. 1988; Bach Knudsen, Jensen et al.  
54        1993), there is less information on quantitative changes in their molecular weights (Mw).  
55        There is a few studies showing changes in the molecular size of oat beta-glucans and of wheat  
56        and rye arabinoxylans during digestion in the upper GI tract (Johansen, Wood et al. 1993;

57 Johansen, Bach Knudsen et al. 1997; Le Gall, Eybye et al. 2010). However, there is scarce  
58 information in the literature regarding specific information on the Mw changes of soluble  
59 barley beta-glucans during passage in the GI tract and possible variations among/with  
60 different barley varieties. This is important since changes in Mw will affect the physico-  
61 chemical properties of the beta-glucans significant for their possible influence on gut health in  
62 both human and animals.

63 The main objectives of the present experiment were to measure and document the  
64 degree of depolymerization (changes in Mw) of soluble barley beta-glucans in the small  
65 intestine of pigs, and study possible differences between different dietary treatments using  
66 four barley varieties.

67

## 68 **2. Material and methods**

### 69 *2.1 Dietary treatments*

70 Four pelleted diets were produced at the Centre for Feed Technology, Ås, Norway. These  
71 were based on four Norwegian barley varieties: Olve (normal starch), Marigold (normal  
72 starch), Karmosè (high amylose starch) and Magdalena (waxy starch). The barley varieties  
73 were grown at the same location (Landvik, Norway) under the same growth conditions in  
74 2010. The diets were formulated to meet the requirements for all nutrients (Subcommittee on  
75 Swine Nutrition, Committee on Animal Nutrition et al. 1998). The composition of the diets is  
76 given in Table 1.

77

### 78 *2.2 Experimental animals*

79 The feeding experiment was performed at the Experimental Farm, Department of Animal and  
80 Aquacultural Sciences, Norwegian University of Life Sciences, Ås, Norway. All pigs were  
81 cared for according to laws and regulations controlling experiments with live animals in  
82 Norway (Animal Protection Act of December 20, 1974, and the Animal Protection Ordinance  
83 concerning experiments with animals of January 15, 1996).

84 A total of 16 female pigs ((Norwegian Landrace x Yorkshire) X (Norwegian Landrace  
85 x Duroc)) from 4 litters were used in the experiment with an average initial weight at 29.8 kg  
86 and an average final weight at 37.6 kg. They were blocked by litter and by live weight, and  
87 groups of four animals were fed each experimental diet.

88

### 89 *2.3 Experimental procedure*

90 The total experimental period lasted for 14 days; a 5-day adaptation period followed by a 9-  
91 day experimental period with collection of faeces the last four days. The pigs were given feed  
92 twice daily according to a restricted Norwegian feeding scale (Øverland, Granli et al. 2000).  
93 The experimental animals were fed in pens designed for individual feeding in a room with an  
94 average temperature of 20.4°C, and had free access to water.

95

### 96 *2.4 Sample collection*

97 The pigs were slaughtered at a commercial slaughter house three hours after the last meal.  
98 The digestive tract was separated from the animal at the slaughter line, and the collection of  
99 digesta from duodenum and ileum was performed immediately. The duodenal samples were  
100 collected from the pyloric ring and 64 cm distally, and the ileal samples from the ileacaecal  
101 opening and 64 cm proximally. The samples were put in closed boxes and kept on ice until

102 being frozen at -20°C. The samples were freeze dried and ground homogenously before being  
103 analysed.

104

## 105 *2.5 Analytical methods*

106 The four diets were analyzed for yttrium by inductively coupled plasma mass spectrometry  
107 (ICP-AES analysis, Perkin-Elmer Optia 3000DV; Perkin-Elmer, Wellesley, MA, USA) at 371  
108 nm, after mineralization and solubilization in acid of the pooled sample.

109

### 110 *2.5.1 Extraction of soluble barley beta-glucans for molecular weight determination*

111  $\beta$ -Glucans were extracted as described by Rieder et al. (Rieder, Holtekjølen et al. 2012). The  
112 initial step involved adding 10 mL of 50% ethanol to a 200 mg sample of the ground diets and  
113 of freeze dried duodenal and ileal samples. The mixture was boiled for 15 min., cooled and  
114 centrifuged (2000 g, 15 min; Heraeus Multifuge 4 KR). The supernatant was discarded before  
115 20 mL 2.5 mM CaCl<sub>2</sub> and 50  $\mu$ L thermostable  $\alpha$ -amylase (Termamyl, Novozymes A/S,  
116 Denmark) was added to each sample. The samples were boiled for 90 min. with mixing every  
117 15 min. After cooling, samples were centrifuged (2500 g, 15 min; Heraeus Multifuge 4 KR)  
118 and the supernatants collected. Another 10 mL of 2.5 mM CaCl<sub>2</sub> was added and the procedure  
119 repeated with boiling for 60 min. The supernatants were combined with the previously  
120 obtained supernatants and stored frozen before molecular weight analysis.

121 Content of soluble beta-glucan was calculated as the difference between total beta-  
122 glucan and insoluble beta-glucan determined by a mixed-linkage beta-glucan assay kit  
123 (Megazyme International Ltd., Wicklow, Ireland). Insoluble beta-glucan was determined in  
124 aliquot samples after removal of soluble beta-glucan by extraction.

125

126 *2.5.2 Relative estimation of molecular weight distribution of soluble barley beta-glucans*

127 (*M<sub>w</sub>-SBB*)

128 The apparent molecular weights of soluble barley beta-glucans (hereafter referred to as *M<sub>w</sub>-*  
129 *SBB*) were determined by HPLC-SEC equipped with a post column addition of calcofluor  
130 combined with fluorescence detection. The HPLC system consisted of a dual pump system  
131 (DIONEX P680) one pump delivering the eluent (50 mM Na<sub>2</sub>SO<sub>4</sub>) at a flow rate of 0.5  
132 mL/min and the other delivering calcofluor (Megazyme International Ltd.) solution (25 mg/L  
133 in 0.1 M tris(hydroxymethyl)aminomethane, Sigma, Schnellendorf, Germany) at a flow rate of  
134 0.25 mL/min. A Spectraphysics AS3500 auto injector was coupled to two serially connected  
135 columns (Tosho; TSK G6000PWXL + G5000PWXL (7.8mm ID x 30.0cm) in series equipped  
136 with a TSK Gel PWXL (6.0mm ID x 4.0cm) guard column).

137 A T-valve placed in the oven containing the columns (40°C) delivered the calcofluor  
138 post column. Injection volume was 20mL and a fluorescent detector (Shimadzu RF-6A,  
139 Shimadzu Europa, Duisburg, Germany) was used with 415nm excitation and 445nm emission  
140 for detection. The HPLC system was controlled with Chromeleon 6.80 (DIONEX, Sunnyvale,  
141 CA, USA).

142 Beta-glucan *M<sub>w</sub>* standards with average given *M<sub>w</sub>* values of 35600, 70600, 229000,  
143 26500, 391000 and 650000 were obtained from Megazyme. The standards were solubilised in  
144 the eluent (50mM Na<sub>2</sub>SO<sub>4</sub>) added 0.02% NaN<sub>3</sub>) by boiling for 5 min. and filtered through a  
145 Millex-AA filter, syringe-driven filter, 33mm, 0.8µm (Merck Millipore Ltd, Ireland). The  
146 standards were then diluted with eluent to give a final concentration of 300µg/mL. A  
147 calibration curve based on the *M<sub>p</sub>* (peak molecular weight) of the *M<sub>w</sub>* standards versus their  
148 elution volume) was established based on the classical principle of narrow molecular weight

149 standards. Weight average Mw distributions of the samples were calculated from this using  
150 WINGPC-6.2 (PSS) offline using a polynomial fitted standard curve. The classification of the  
151 molecular weight distribution into high and low molecular weights (HMw and LMw) was  
152 based on dividing the chromatogram in two regions (by elution time); high (20-30 min.) and  
153 low (30-42 min.). This cutting point corresponded to ca. 250 kDa in the standard curve.

154 The calculated weight average Mw's (Mwcalc) only include  $\beta$ -glucan molecules large  
155 enough to interact with calcofluor and hence be detected by the resulting fluorescence signal  
156 (Rieder, Knutsen et al. 2012). From in-house experiments this cut-off value is approximately  
157 30.000-40.000, but this value is so far not been exactly determined. The reported values  
158 therefore do not represent the exact weight average Mw of the samples, but rather the  
159 calcofluor based average Mwcalc. Furthermore, since high molecular weight standards (Mw >  
160 650.000) are not available, there is no accurate determination of the molecular weight in the  
161 upper range Mw > 650.000). However, for comparative purposes and assessing relative  
162 changes in Mw, the methodology was considered appropriate. In fact the unique specificity of  
163 the system does not display any or very little interference with starch and other soluble  
164 polysaccharides such as arabinoxylan in the system. Cellulose is not soluble and hence not  
165 detected.

166 The SBB were solubilized in water as described by Rieder et al. (Rieder, Holtekjølen  
167 et al. 2012) and for the analysis of the actual samples 1.0mL of each water extract was filtered  
168 as above and diluted 1:1 with 0.04% NaN<sub>3</sub> before injecting into the system. The results of the  
169 duodenal and ileal samples are an average of 4 biological replicas. The variation between the  
170 technical parallels was less than 10% with a few exceptions.

171

172 *2.6 Data analysis*

173 Analysis of variance and significant differences among means were tested by one-way  
174 ANOVA, using Minitab (version 16; Minitab Inc., State College, PA). Significant differences  
175 were declared at  $P < 0.05$ .

176

### 177 **3. Results and discussions**

#### 178 *3.1 Molecular weight distribution of soluble barley beta-glucan (Mw-SBB) in the diets*

179 The SBB in the four experimental diets exhibited similar monomodal size distribution as seen  
180 in Figure 1. The Mw-SBB of the four diets however varied and the diet including the barley  
181 variety *Magdalena* (hereafter referred to as Diet-Mag) had a significantly higher average Mw-  
182 SBB than the rest of the diets. The diet including the barley variety *Karmosè* (hereafter  
183 referred to as Diet-Kar) had the lowest Mw-SBB of the four diets (Fig. 1).

184

#### 185 *3.2 Effect of digestion on molecular weight distribution of soluble barley beta-glucans (Mw- 186 SBB)*

##### 187 *3.2.1 Duodenum – beginning of the small intestine*

188 The results show a significant depolymerisation of the SBB already at the beginning of the  
189 small intestine (duodenum) (Fig. 2). The average molecular weight (average of all diets and  
190 all pigs) (AMw-SSB) decreased from approximately 1050 kDa in the diets to ca. 460 kDa in  
191 the duodenal samples, a reduction of 55%. There was also a shift in retention time and a  
192 broadening of the peak into a bimodal size distribution in the duodenal samples independent  
193 of diet (Fig. 2). This showed that the Mw-SBB was depolymerized and that the reduction  
194 resulted in two significantly different populations; one population of high molecular weight  
195 SBB (HMw-SBB) and one of low molecular weight (LMw-SBB). In the literature there are

196 many studies on fermentation pattern and degradation rate of barley beta-glucans in pigs.  
197 However, there is scarce information regarding changes in molecular weight of barley beta-  
198 glucans. For oat beta-glucans similar depolymerisation pattern has been observed (Johansen,  
199 Wood et al. 1993; Johansen, Bach Knudsen et al. 1997).

200 The average HMw-SBB size distribution (as average of all diets and all pigs) was 940  
201 kDa and it accounted for ca. 45% of the molecular size distribution in the duodenal samples,  
202 while the average LMw-SBB was 105 kDa with a 55% share. Also oat beta-glucans showed  
203 depolymerisation in the upper small intestine of pigs (up to 55%) (Johansen, Bach Knudsen et  
204 al. 1997).

205 All diets showed the same change into a bimodal size distribution. Still, some  
206 significant differences were seen depending on the diet. Overall, diet-Mag had the  
207 significantly highest average Mw-SBB, followed by diet-Kar and diet-Olv, with diet-Mar  
208 having the lowest. Also, the portion of high molecular weight SBB (HMw-SBB) differed and  
209 the largest part of HMw-SBB was found in the Diet-Mag (51%), while diet-Mar had the  
210 lowest (32%).

211

### 212 *3.2.2 Ileum – end of the small intestine*

213 The average molecular weight of SBB in ileum showed a significant decrease  
214 compared with the duodenal samples, from 460 kDa to 250 kDa respectively ( $P < 0.05$ ). The  
215 corresponding decrease in AMw-SBB compared to the original diets was 75%.

216 The results showed that the SBB was depolymerized throughout the small intestine  
217 with a shift towards a higher portion of LMw-SBB in the ileal samples (Fig. 3) compared with  
218 the duodenal samples. Thus, the low molecular weight portion increased moving through the  
219 small intestine from the duodenum to the ileum. The share of HMw-SBB decreased equally,  
220 and again, diet-Mag had the highest Mw-SBB and the largest portion of HMw-SBB (only

221 28%) in the ileal samples, with diet-Mar the lowest (15%) (Fig. 3). This is consistent with  
222 findings for oat. Johansen et al. (1997) showed an increased depolymerisation for oat beta-  
223 glucans going from the proximal to the distal small intestine in pigs. Thus, the oat beta-  
224 glucans in the distal small intestine after 3h post-prandial showed higher depolymerisation,  
225 decreasing the share of high Mw oat beta-glucan.

226

### 227 *3.3 Variations among pigs in distribution of molecular weight distribution of soluble barley* 228 *beta-glucans (Mw-SBB) in the duodenal and ileal samples*

229 Some variations were seen among the experimental animals fed the same diet (see figure 4  
230 and 5). Figure 4 shows the variations found in the duodenal samples within pigs fed Diet-  
231 Mag, while Figure 5 shows the variation among the ileal samples of the pigs fed the Diet-  
232 Mag. The observed variations among pigs fed the same diet might relate to differences in the  
233 microorganisms present in their gastrointestinal tract. It could also be associated with  
234 variation in the matrix of the pellets after chewing as well as different drinking pattern. The  
235 variations between the biological parallels make it important to include a sufficient number of  
236 biological parallels to obtain reliable data as well as to verify the results when working with  
237 animals and animal trials. Still, despite some variation among pigs fed the same diet, the  
238 effect on SBB is evident. The molecular weight of the SBB is reduced and the  
239 depolymerisation starts at duodenum and continues all the way through the small intestine. At  
240 ileum the Mw-SBB is reduced up to 80% compared to the original diet.

241

## 242 **4. Conclusion**

243 Soluble barley beta-glucan (SBB) is depolymerized during digestion in pigs and there is a  
244 significant depolymerisation of SBB naturally occurring already in the upper GI tract, in the

245 small intestine. Our results show that depending on variety, the SBB is depolymerized up to  
246 60% in the duodenum and 80% in the ileum. Thus, before the SBB has reached the hindgut  
247 and is fermented, its Mw has already been significantly reduced into a larger share of low  
248 molecular weight SBB (ca. 100 kDa). The depolymerisation of the beta-glucan might be due  
249 to hydrolytic enzymes excreted by microbiota in the upper digestive tract of the  
250 animal. However, retained endogenous hydrolase activities in the barley material may be  
251 present despite barley processing and transit through the upper GI-tract.

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### **Conflict of interest statement**

The authors confirm no conflict of interest with this article.

## Tables

**Table 1. Composition of the four diets and their amount of soluble beta-glucan (%)**

	Diet 1	Diet 2	Diet 3	Diet 4
Barley <i>Marigold</i>	83.47			
Barley <i>Magdalena</i>		83.47		
Barley <i>Karmosè</i>			83.47	
Barley <i>Olve</i>				83.47
Soybean meal (HiPro)	15.0	15.0	15.0	15.0
Limestone meal (CaCO <sub>3</sub> )	1.3	1.3	1.3	1.3
Mineral premix	0.16	0.16	0.16	0.16
Vitamin premix	0.06	0.06	0.06	0.06
Y <sub>2</sub> O <sub>3</sub> *	0.01	0.01	0.01	0.01
Soluble beta-glucan	1.6	3.0	2.6	2.6

\*Yttrium oxide was used as the indigestible dietary marker.

## Figure legends

**Figure 1:** Chromatogram showing the relative molecular weight profile of the soluble barley beta-glucans (Mw-SBB) in the diets based on the different barley varieties including their calculated average Mw-SBB as bar graphs. The error bars represent the standard deviations (two technical parallels).

**Figure 2:** Chromatogram showing the relative molecular weight profile of the soluble barley beta-glucans (Mw-SBB) in the different duodenal samples including their calculated average Mw-SBB as bar graphs. The degree of depolymerisation compared to the Mw-SBB in the corresponding diets are given (in %) above the bars. The error bars represent the standard deviations (four biological replicas (pigs)).

**Figure 3:** Chromatogram showing the relative molecular weight profile of the soluble barley beta-glucans (Mw-SBB) in the different ileal samples including their calculated average Mw-SBB as bar graphs. The degree of depolymerisation compared to the Mw-SBB in the corresponding diets are given (in %) above the bars. The error bars represent the standard deviations (four biological replicas (pigs)).

**Figure 4:** Example of the variation found in the relative molecular weight profile of duodenal samples among the four pigs (1-4) fed the same diet (Magdalena). The two overlapping chromatograms represent the two technical parallels.

**Figure 5:** Example of the variation found in the relative molecular weight profile of ileal samples among the four pigs (1-4) fed the same diet (Magdalena). The two overlapping chromatograms represent the two technical parallels.

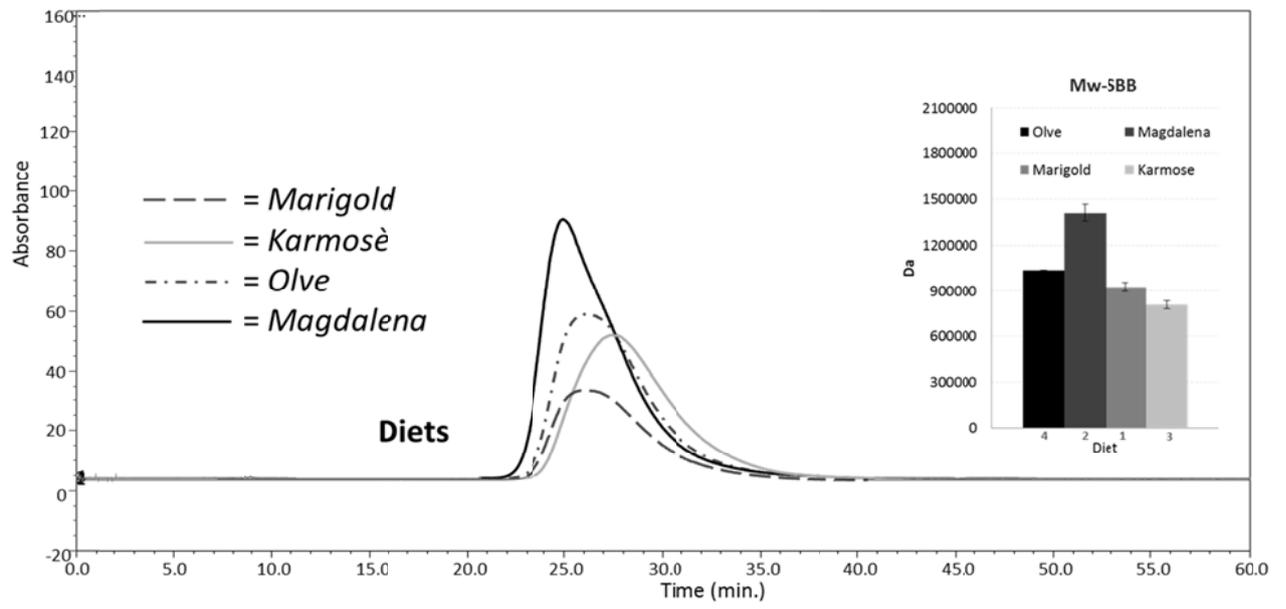
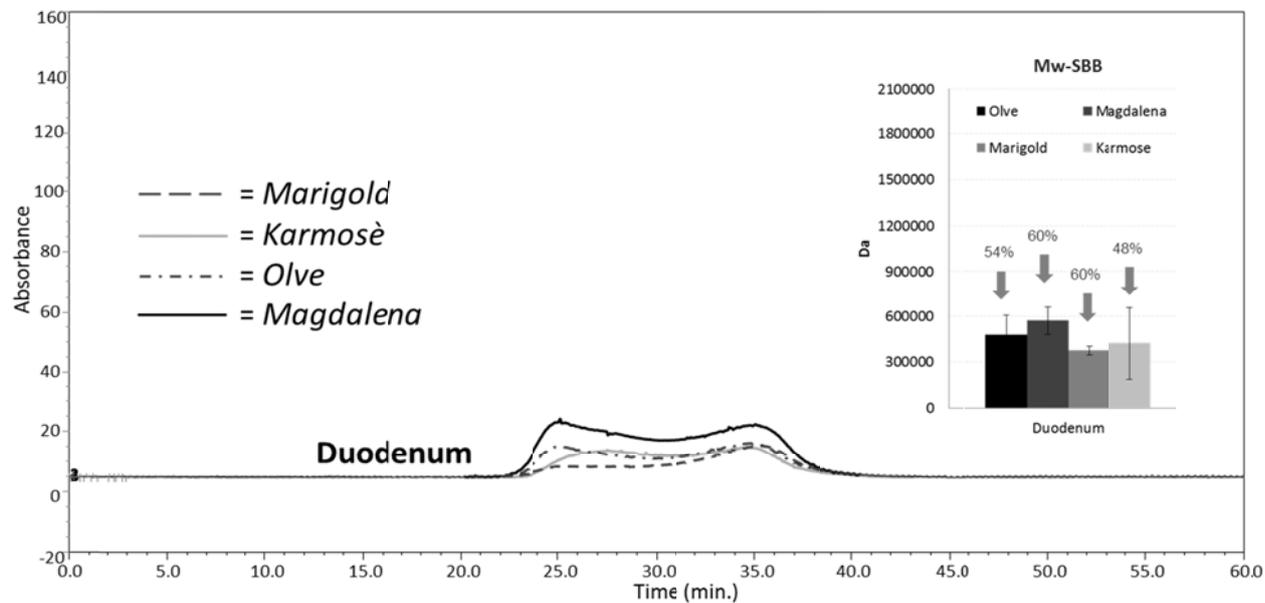
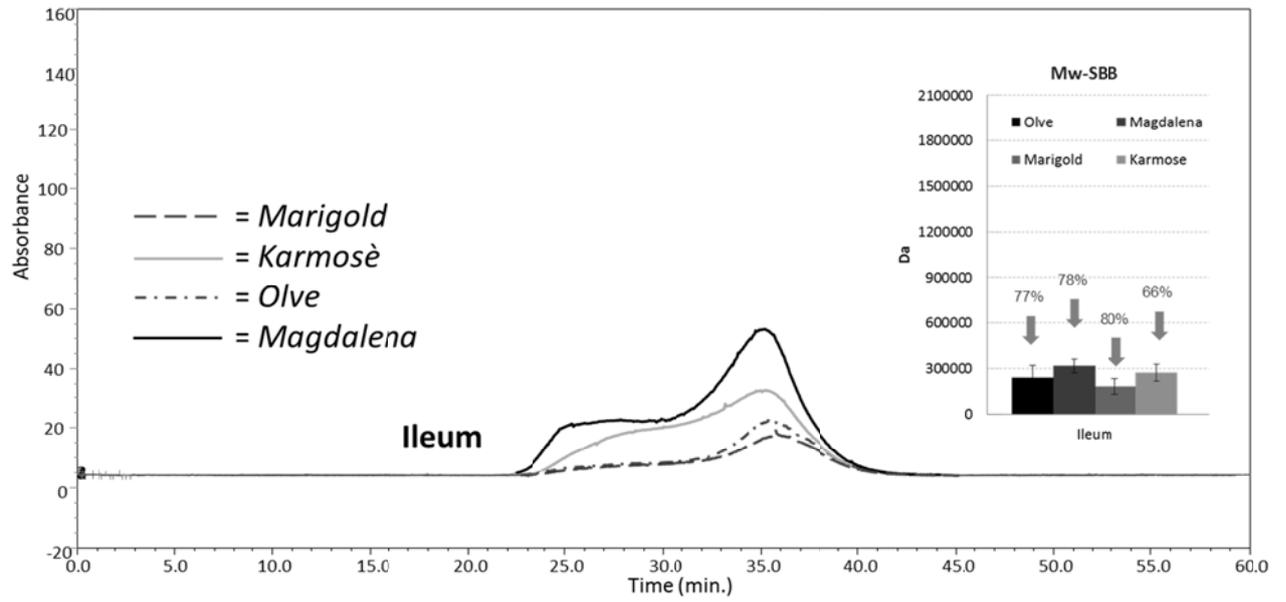


Figure 1



**Figure 2**



**Figure 3**

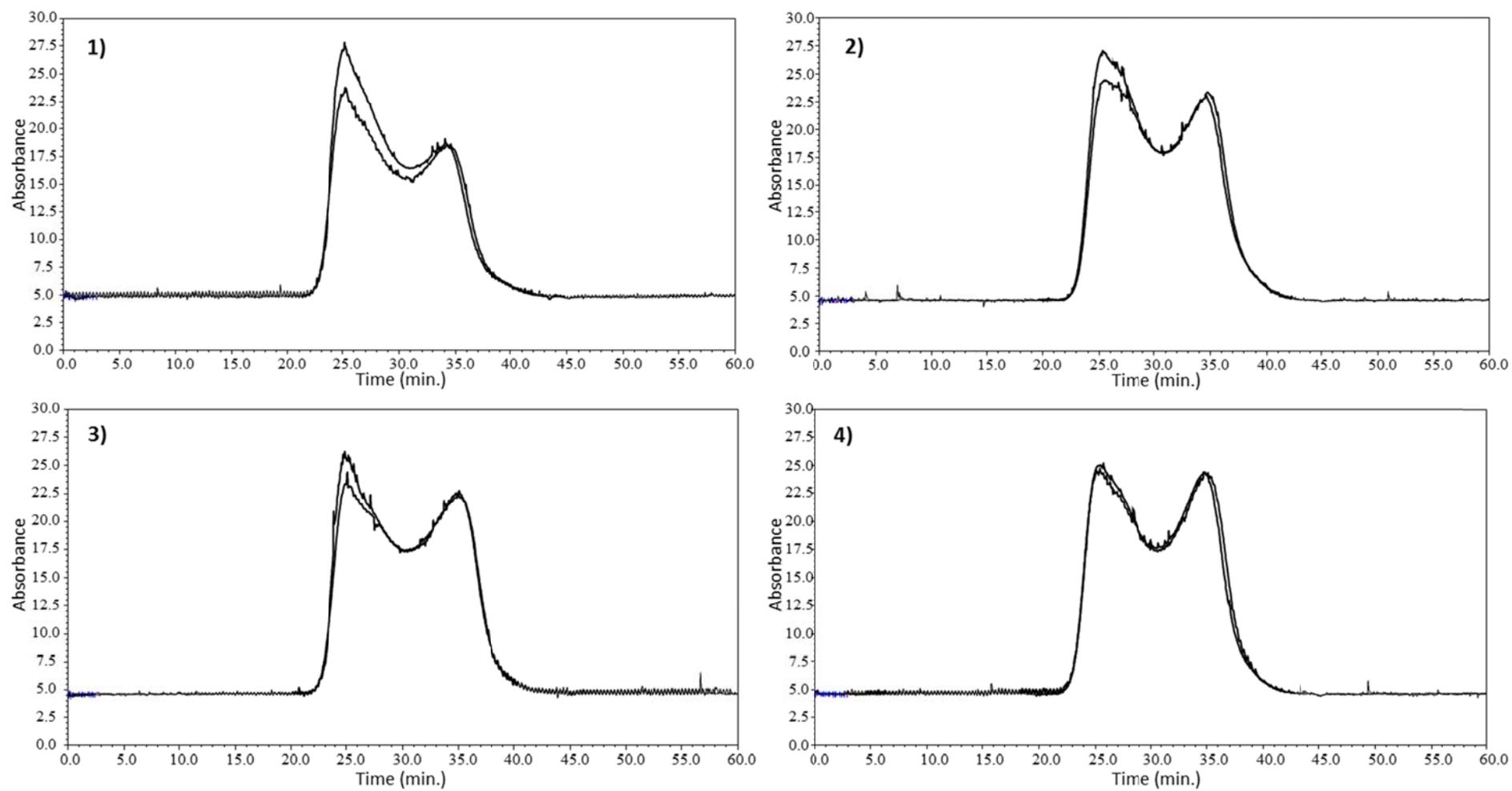


Figure 4

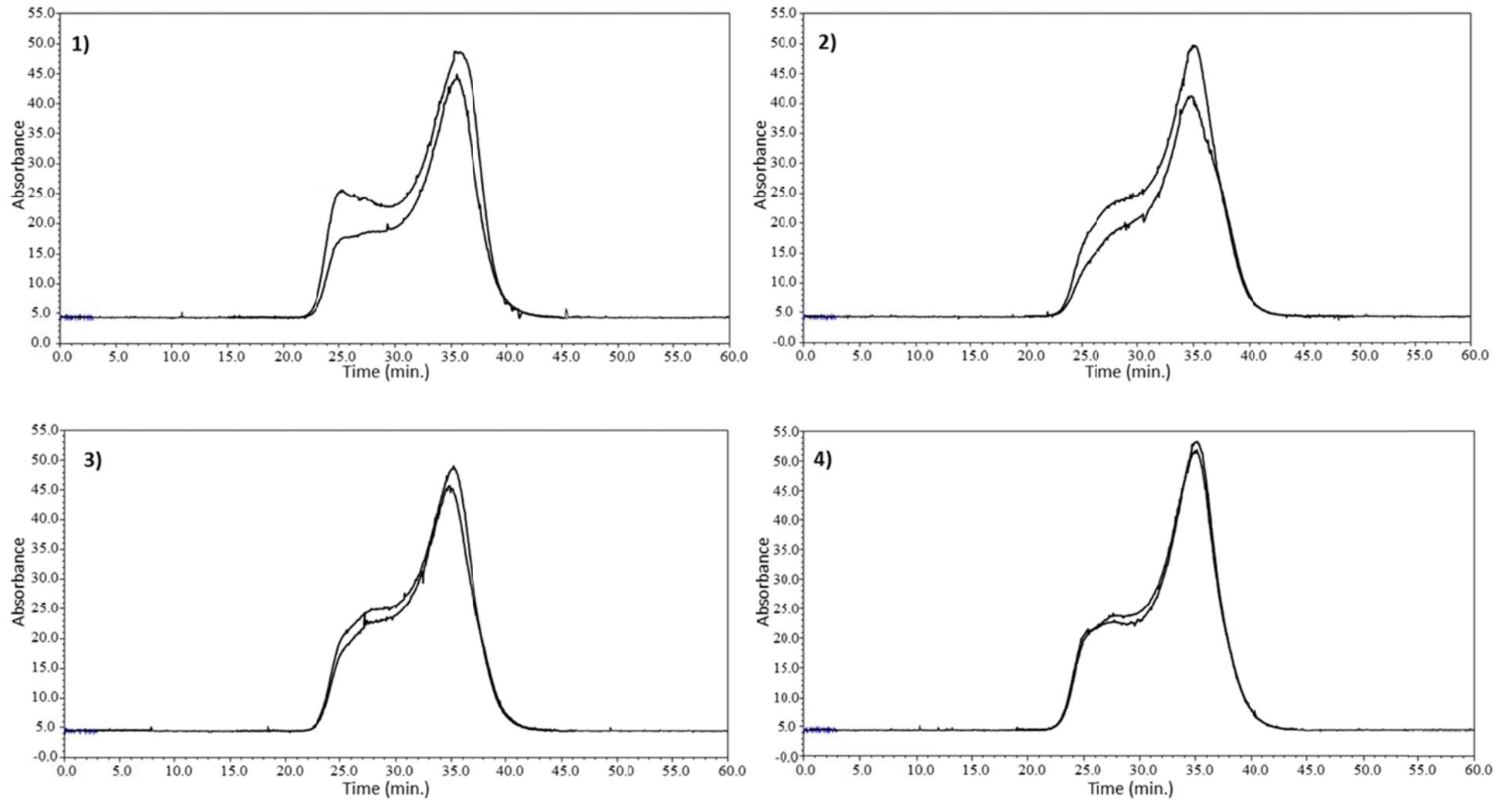


Figure 5

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