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1	Identification of Features Associated with Plant Cell Wall Recalcitrance to Pretreatment by
2	Alkaline Hydrogen Peroxide in Diverse Bioenergy Feedstocks Using Glycome Profiling
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20	ABSTRACT
21	A woody dicot (hybrid poplar), an herbaceous dicot (goldenrod), and a graminaceous
22	monocot (corn stover) were subjected to alkaline hydrogen peroxide (AHP) pretreatment and
23	subsequent enzymatic hydrolysis in order to assess how taxonomically and structurally diverse
24	biomass feedstocks respond to a mild alkaline oxidative pretreatment and how differing features
25	of the cell wall matrix contribute to its recalcitrance. Using glycome profiling, we determined
26	changes in the extractability of non-cellulosic glycans following pretreatment and screened a
27	panel of 155 cell wall glycan-specific monoclonal antibodies against these extracts to determine
28	differences in the abundance and distribution of non-cellulosic glycan epitopes in these extracts
29	and assess pretreatment-induced changes in the structural integrity of the cell wall. Two
30	taxonomically-dependent outcomes of pretreatment were identified that both improved the
31	subsequent enzymatic hydrolysis yields but differed in their impacts on cell wall structural

integrity. Specifically, it was revealed that the goldenrod exhibited decreases in all classes of alkali-extractable glycans indicating their solubilization during pretreatment, which was accompanied by an improvement in the subsequent extractability of the remaining cell wall glycans. The corn stover did not show the same decreases in glycan abundance in extracts following pretreatment, but rather mild increases in all classes of cell wall glycans, indicating

overall weaker associations between cell wall polymers and improved extractability. The hybrid
poplar was relatively unaffected by pretreatment in terms of composition, enzymatic hydrolysis,
and the extractability of cell wall glycans due presumably to its higher lignin content and denser
vascular structure.

Keywords: glycome profiling, cell wall recalcitrance, bioenergy, AHP pretreatment, plant cellwalls

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44 BROAD IMPACTS AND HIGHLIGHTS

The cell wall matrix of plants provides an effective barrier to the chemical and/or 45 biological deconstruction to its monosaccharide components for use in biofuels applications. 46 The properties of the cell wall matrix impacting its recalcitrance set by many features of cell wall 47 macromolecules across a wide range of length scales. Changes in the physical and chemical 48 properties of cell wall polymers and the overall cell wall environment are important outcomes of 49 pretreatments. Quantitatively identifying how properties of structurally and taxonomically 50 51 diverse cell walls are impacted by pretreatments and how this correlates to reduced recalcitrance is an important goal with implications for both designing plants with properties suited for 52 deconstruction^{1, 2} and designing effective deconstruction strategies. This work applied glycome 53 profiling to identify how a mild alkaline oxidative pretreatment impacts the composition and 54 55 structural organization of the cell wall and identified two distinct mechanisms by which this pretreatment overcomes cell wall recalcitrance in either herbaceous dicots or graminaceous 56 monocots. These results highlight both the taxonomic differences in cell wall organization and 57 the differences in their response to pretreatment. 58

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60 **INTRODUCTION**

The lignified cell walls of vascular plants are the result of nearly half a billion years of evolution to resist biological degradation while serving the structural and physiological needs of

the plant.^{3, 4} It is believed that the majority of terrestrial carbon in the biosphere is sequestered within plant cell walls,⁵ yet this vast resource of reduced carbon is used primarily by humans for its existing structural value (as fiber/paper and as a structural/building material), as a fuel for combustion, or as ruminant forage rather than for the value contained in its existing chemical constituents. This is due to the recalcitrance of the cell wall to deconstruction by chemical and biological treatments and is set by features that are both structural and chemical that cut across the length scales at the molecular, macromolecular, and cellular levels.^{6, 7}

The plant cell wall matrix is a complex network of cellulose and other matrix 70 polysaccharides including hemicelluloses and pectin, lignin, and structural proteins and presents 71 an obvious barrier to water and cellulolytic enzyme penetration, in particular due to lignin's 72 capacity to set cell wall hydrophobicity and porosity. However, underpinning this lignin barrier 73 is a network of non-cellulose, cell wall matrix polysaccharides providing structure and 74 organization. The major classes of matrix polysaccharides include the xylans (GAXs and GXs), 75 glucomannans (GMs), xyloglucans (XyGs), mixed-linkage glucans (MLGs), pectins, and cell 76 wall protein glycosylations.⁸⁻¹⁰ The abundance, composition, and substitution patterns of these 77 glycans vary temporally during plant growth and cell wall expansion, spatially within cell walls 78 and between plant tissues, and taxonomically across diversity plant. In dicots, XyGs and GAXs 79 are the predominant non-cellulosic glycans in the primary cell walls with GXs the predominant 80 hemicellulose in dicot the secondary cell walls.¹¹ In grasses, GAXs have been found to occupy a 81 significant fraction of the interstitial space between cellulose microfibrils in the primary cell 82 walls in addition to mixed-linkage MLGs and grass-specific XyGs and GMs.^{2, 12} Pectic 83 polysaccharides are complex and can comprise up to 30% of the primary cell walls of dicots and 84 significantly less in grasses¹³ where some of this function is thought to be performed by other 85 glycans including MLGs and GAXs.² Structural proteins may comprise up to 10% of the cell 86 wall in some plant tissues and can be significantly glycosylated, for example with 87 arabinogalactans (AGs).⁸ 88

Matrix polysaccharide content, diversity, interactions, and distribution play a role in recalcitrance by setting the accessibility of cellulose to celluloytic enzymes and defining the porosity of the cell wall.¹³⁻¹⁵ This network of macromolecules is built up from a combination of physical entanglement of structural polymers as well as non-covalent and covalent cross-links between macromolecules.¹⁶ These non-covalent interactions are important and form the

principle mechanism of association between cell wall glycans.¹⁵ Taken together, this complex 94

composite structure presents a challenge for characterization and a number of approaches have 95

been developed recently with a focus on relating structural features of the cell wall to its 96

recalcitrance as reviewed recently by Foston et al.¹⁷ 97

Immunological methods using glycan directed mAbs are widely used tools to investigate 98 plant cell wall structure.^{18, 19} Besides the cell wall composition and structure, mAbs can be used 99 100 for qualitative and quantitative detection of carbohydrate epitopes in plant sequential extracts. This has been performed in order to characterize pretreatments using one approach that plotted 101 polysaccharides from three increasingly severe cell wall extracts (CDTA, NaOH, cadoxen) onto 102 a microarray, which was then probed with mAbs and CBMs²⁰ in order to identify changes in 103 content and extractability of xylans, XyGs, and MLGs epitopes in hydrothermal pretreated wheat 104 straw.²¹ Recently, Pattathil *et al.* assembled a collection of glycan-specific antibodies²² shown in 105 Supplemental Table S1 and an ELISA-based screen was used to categorize these mAbs with 106 respect to binding affinity for structurally diverse plant cell wall glycans.^{22, 23} This technique has 107 been applied to evaluate the structural, accessibility, and extractability changes of cell wall 108 glycans in hybrid poplar during dilute acid pretreatments²⁴ and to compare differences in the 109 structural features underpinning cellulose digestibility in switchgrass and hybrid poplar.²⁵ 110

Chemical pretreatments are one route to overcoming cell wall recalcitrance²⁶ and can be 111 coupled to an enzymatic depolymerization of cell wall polysaccharides whereby the sugars 112 113 monomers generated can be subsequently biologically converted to fuels and chemicals, providing a path forward for developing a bio-based fuels and chemicals industry that is 114 renewable and petroleum-displacing. Ultimately this recalcitrance to cell wall deconstruction 115 lies in the challenge for cellulolytic enzyme infiltration into the cell wall which can be seen as a 116 117 combination of cell wall porosity, hydrophobicity (or water penetration/swelling), and glycan accessibility. Alkaline hydrogen peroxide (AHP) pretreatment is comparable to alkaline 118 hydrogen peroxide pulp bleaching stages in the paper industry with the difference that higher pH 119 and higher hydrogen peroxide loadings are employed to affect a mild delignification rather than 120 brightening as the outcome.^{27, 28} AHP pretreatment has several potential advantages compared to 121 other pretreatment processes including minimal loss of polysaccharides,²⁹ operation at low 122 temperature and pressure, and minimal formation of inhibitors for fermentation,³⁰ and potentially 123 high enzymatic digestibilities in grasses. In particular, we have recently been able to achieve 124

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ethanol titers greater than 50 g/L from undetoxified hydrolysates of 12.5% (w/w) H₂O₂ loading
 AHP-pretreated corn stover and switchgrass including complete glucose and xylose utilization
 using *Saccharomyces cerevisiae* strains metabolically engineered and evolved for xylose
 fermentation.³⁰

It has been relatively well-established that hydrothermal pretreatments such as dilute 129 acid³¹ or liquid hot water³² overcome recalcitrance through thermal effects by melting and 130 redistributing lignin and catalyzing xylan hydrolysis and solubilization. This effect on xylan 131 removal has been validated using similar characterization approaches for quantifying glycan 132 extractability and epitope abundance.^{21, 24} As AHP pretreatment is mechanistically different from 133 acidic hydrothermal pretreatments and targets lignin solubilization at low temperature while 134 preserving carbohydrates considerably, glycome profiling should be able to provide new insight 135 into matrix polysaccharide-specific contributions to cell wall recalcitrance. 136

Specifically, in this study we investigate the response of cell wall glycans of diverse 137 plants including hybrid poplar (woody dicot), goldenrod (herbaceous dicot), and corn stover 138 (monocot grass) to AHP pretreatment with increasing H₂O₂ loadings. The cell wall response to 139 pretreatment was characterized by compositional changes and overall mass loss of the cell wall, 140 glucan and xylan enzymatic yields using a commercial enzyme cocktail, and glycome profiling 141 of the sequential glycan extracts of the untreated and AHP-pretreated cell walls at 12.5% (w/w) 142 H_2O_2 loading on biomass. Using this information, we draw conclusions about the structural 143 144 changes associated with AHP pretreatment and additionally are able to gain insights into the role that differences in plant cell wall architecture have on cell wall recalcitrance. 145

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147 **EXPERIMENTAL**

148 **1. Pretreatment**

Biomass consisted of a commercial hybrid corn stover (Pioneer Hi-Bred 36H56)
provided through the Great Lakes Bioenergy Research Center (GLBRC), debarked hybrid poplar
(*Populus nigra* var. *charkoviensis* x *caudina* cv. NE-19) grown at the University of Wisconsin
Arlington Agricultural Research Station and provided through the GLBRC, and goldenrod
(*Solidago canadensis*) collected locally in East Lansing, MI and obtained from Dr. Jonathan
Walton (MSU, Plant Biology). Biomass was initially milled with a Wiley MiniMill (Thomas
Scientific) to pass a 2 mm screen and air-dried to ~5% moisture. The milled biomass was

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subjected to alkaline hydrogen peroxide (AHP) pretreatment using H₂O₂ loadings of either 156 12.5%, 25%, 50% (g H_2O_2 per g biomass). These were performed in shake flasks with a 100 g 157 total mass and 2% (w/v) solids concentration for 24 h at 30°C with orbital shaking at 170 rpm 158 and periodic pH adjustment to 11.5. After pretreatments, the liquid in the samples were removed 159 by vacuum filtration (#113 Whatman filter paper) and the slurry was washed several times with 160 deionized water to remove solubles and air-dried at room temperature. The mass yield during 161 pretreatment was quantified as the ratio between the mass (dry basis) after pretreatment and the 162 initial mass before pretreatment. 163

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2. Composition analysis, enzymatic hydrolysis and digestibility determination

The pretreated biomass was subjected to a two-stage acid hydrolysis according to NREL 166 composition analysis³³ to determine neutral polysaccharide content, Klason lignin, and ash with 167 the difference that Aminex HPX-87H (Bio-Rad, Hercules, CA) column was used to quantify the 168 glucan, xylan+galactan+mannan, arabinan, as well as acetate content. The total uronic acids were 169 assayed enzymatically (K-Uronic, Megazyme, Wicklow, Ireland). The extractives content was 170 determined by a sequential 3-step extraction including 70% ethanol, followed by 1:1 (v/v)171 methanol and chloroform mixture, then acetone. Three extraction cycles for each solvent were 172 performed and followed by centrifugation at 10500 x g for 10 minutes for each cycle. Before the 173 enzymatic hydrolysis, the pretreated biomass was ball-milled for 3 cycles using a QIAGEN 174 175 TissueLyser II equipped with 25 mL Teflon jars and 20 mm diameter Teflon balls at 30 Hz for 2 minutes with liquid nitrogen cooling. The ball-milled samples were incubated with Cellic CTec2 176 (Novozymes, Bagsværd, Denmark) at a loading of 30 mg protein/g glucan at 50°C, 10% solid 177 loading and 5 mL total volume in 0.05 M Na-citrate buffer pH 4.8, for 24 hours or 72 hours. The 178 glucan and xylan yields (based on only glucan and xylan remaining after pretreatment) were 179 determined by the HPLC analyzable glucose and xylose concentrations after hydrolysis divided 180 by the original glucan and xylan contents in the pretreated samples. 181

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183 **3.** Sequential extraction, glycome profiling

Sequential cell wall extractions and glycome profiling were carried out as described
 previously.^{22, 24, 34} The six extractions included (in order of extraction) oxalate to remove "loose"
 pectins, carbonate to remove "more tightly" bound pectins, 1M KOH to remove "loose"

187 hemicelluloses along with tightly bound pectins, 4M KOH to remove "tightly" bound hemicelluloses along with tightly bound pectins, acid chlorite to oxidize and solubilize lignin and 188 release lignin-embedded hemicelluloses, and a 4M KOH post-chlorite treatment to remove 189 additional lignin-bound polysaccharides. Plant glycan-directed monoclonal antibodies were from 190 laboratory stocks (CCRC, JIM and MAC series) at the Complex Carbohydrate Research Center 191 (available through CarboSource Services; http://www.carbosource.net) or were obtained from 192 BioSupplies (Australia) (BG1, LAMP). A description of the mAbs used in this study can be 193 found in the Supporting Information, Table S1, which includes links to a web database, 194 WallMAbDB (http://www.wallmabdb.net) that provides detailed information about each 195 antibody. 196

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198 RESULTS AND DISCUSSION

199 1. Changes in composition and mass loss

Three types of biomass representing three classes of plants that may offer promise as 200 feedstocks for cellulosic biofuels were tested in this study. Corn stover represents the 201 agricultural residue with the highest production and availability for bioenergy applications in the 202 U.S.,³⁵ while short-rotation hybrid poplar has agronomic, logistical, and environmental 203 advantages as a feedstock.³⁶ "Low-input high-diversity" bioenergy landscapes have many 204 attractive sustainability attributes³⁷ and comprise mixed communities of plants on marginal or 205 206 degraded lands, and in this study we use goldenrod (Solidago canadensis) as a representative herbaceous dicot that may be present in these landscapes. The composition of the untreated 207 biomass is presented in Table 1. Notable differences include the low content of pectic 208 polysaccharides (as uronic acids) in the corn stover, which is 5-fold lower than the goldenrod and 209 210 2-fold lower than the hybrid poplar. The goldenrod has a substantially higher extractives content (23.5%) relative to the other two biomass types. Additionally, lignin content of the corn stover is 211 nearly half that of the goldenrod and poplar. 212

AHP pretreatment was performed at increasing H_2O_2 loadings (12.5, 25, and 50% w/w on biomass) which would be significantly higher than would be economically practicable industrially. The reason for these high loadings was to compare and analyze how the cell walls from phylo-genetically diverse differ in their susceptibility to low temperature mild oxidative delignification and hemicellulose extraction and as a screen for overall differences in enzymatic

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218 hydrolyzability. The total cell wall mass loss (excluding extractives), xylan loss, and lignin loss for the three biomass types following pretreatment with increasing H₂O₂ loadings show distinct 219 220 responses between the biomass types (Figure 1). For poplar, minimal material was solubilized with pretreatment (<1% by mass), while up to 20% and 25% of the mass the cell walls of the 221 goldenrod and corn stover, respectively, was solubilized by pretreatment at the higher H_2O_2 222 loadings. The mass of corn stover decreased continuously with increasing H₂O₂ loading, while 223 the sample mass of poplar and goldenrod decreased abruptly with the mildest treatment (12.5%) 224 H₂O₂). For individual cell wall components, AHP pretreatments resulted in minimal changes in 225 glucan for all biomass types representing preservation of cellulose (data not shown) which is 226 consistent with our previous findings.^{27, 29} while the xylan content decreased only for the corn 227 stover (Figure 1B). For goldenrod, the Klason lignin content was only slightly reduced by AHP 228 pretreatment and did not change significantly by increasing H₂O₂ loading. The simultaneous 229 removal of xylan and Klason lignin in corn stover was increased by increasing H₂O₂ loading. 230 This is a well-known property of grass cell walls and is due to the higher solubility of grass 231 lignins³⁸ and alkali-only extraction of lignin and xylans in grasses is known to be significantly 232 higher in grasses than in woody dicots.^{25, 39} Besides alkali solubility, cleavage of ester and ether 233 cross-links between xylan and lignin or lignin and lignin mediated by ferulate^{12, 40} in grasses are 234 thought to be an important target of AHP pretreatment²⁹ and likely contribute to these outcomes. 235 236

237 2. Enzymatic hydrolysis yields of poplar, goldenrod, and corn stover

Figure 2 shows the enzymatic hydrolysis yields of glucose for poplar, goldenrod, and 238 239 corn stover subjected to increasing H_2O_2 loadings for 24 and 72 hr hydrolysis times using a commercial cellulase (Cellic CTec2) with no xylanase supplementation. These results represent 240 241 screening only and were not focused on optimizing enzyme cocktail or loading. As such, improved glucose (and xylose) yields at lower enzyme loadings would be observed if xylanase 242 and pectinase were used. These results show that the glucose yields were increased with 243 increasing H₂O₂ loading for the three types of biomass. Similar to the results for compositional 244 245 changes (Figure 1), the yield changes with pretreatment are significantly different between the three classes of plants tested. For poplar, the glucose yields were significantly lower than 246 goldenrod and corn stover with the highest glucose yields approaching 40%. For goldenrod, the 247 glucose yields "saturate" at approximately 70% at the mildest pretreatment condition. For corn 248

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249 stover, the glucose yield approaches 100% with increased H₂O₂ loading and it is known that corn stover as well as many other cereal stovers are often considerably more digestible than many 250 undomesticated grasses.^{29, 40} Gould⁴¹ determined that diverse graminaceous monocots responded 251 considerably better to AHP pretreatment at 100% (w/w) H₂O₂ loadings than herbaceous 252 dicots/forbs (including goldenrod) and that the 7 grasses tested had on average more than double 253 the improvement in digestibility of the 11 forbs tested following AHP pretreatment, although the 254 255 goldenrod showed the largest improvement in digestibility of the dicots tested and was among the highest in terms of absolute glucose release per gram of biomass. 256

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258 3. Glycome profiling of poplar, goldenrod, and corn stover

Glycome profiling (GP) provides quantitative information on both how pretreatment 259 impacts the strength of association between cell wall glycans and other cell wall matrix polymers 260 and how pretreatment impacts cell wall glycan composition. For GP, the samples were subjected 261 to increasingly severe extractions to sequentially remove the cell wall polymers, followed by 262 quantification of recovered materials in each extraction step, and probing of the binding strength 263 for the diverse array of antibodies covering a range of non-cellulose cell wall polysaccharide 264 epitopes.^{22, 24, 34} The categories of glycan-specific mAbs were determined previously based on 265 hierarchical clustering of the antibodies against a panel of 54 known plant polysaccharides,²² 266 with complete data on the binding specificity and cross-reactivity for each mAb on the web 267 268 database, WallMAbDB (http://www.wallmabdb.net). It should be noted that the GP approach does not yield information on small glycan molecules (e.g. oligomeric glycans and 269 monosaccharides) as only larger cell wall glycans are able to effectively adsorb to the ELISA 270 plates.²² For the sequential extractions, the strength of association between individual glycans 271 272 and other cell wall matrix polymers may be hypothesized to be mechanistically due to differences in: (1) the strength of non-covalent association or physical entanglement between 273 polymers or "encrustation" within lignin, (2) location within the cell wall (surface vs. interior), 274 and (3) tissue type (e.g. lignified parenchyma and sclerenchyma versus low lignin pith tissues). 275 276 The role of lignin in setting the alkali-extractability of cell wall glycans was recently demonstrated by Pattathil et al.,⁴² whereby it was identified that alfalfa lines with disrupted 277 monolignol synthesis resulting in a low-lignin phenotype contained considerably more alkali-278 extractable glycans than the control line where much more of the cell wall glycans were 279

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280 extractable only after chlorite delignification.

The GP results from this work are presented in Figure 3 for the three biomass categories 281 for either no pretreatment or AHP pretreatment at 12.5% (w/w) H₂O₂ loading on the biomass. 282 Substantial differences can be observed between biomass types and for biomass subjected to 283 pretreatment as quantified for both mass partitioning of extracted glycans (top panel of Figure 3) 284 and differences in the abundance of glycan epitopes in these extracts (heat map in the lower part 285 of Figure 3). For the extract mass partitioning of glycans, it is clear that the two dicots have 286 similar profiles for the four most severe extracts, *i.e.* the 1 M KOH, 4 M KOH, chlorite, and 4 M 287 KOH PC. The goldenrod shows a very high content of the oxalate- and carbonate-extractable 288 polysaccharides relative to the other two types of biomass. This may be a consequence of the 289 goldenrod having a higher proportion of pectic polysaccharide-rich leaves compared to the 290 poplar which consists of only stem heartwood, and the corn stover, which as a graminaceous 291 monocot is known to have low pectic polysaccharide content.⁴³ 292

A number of noteworthy differences are apparent in comparing the glycan epitope 293 abundances within the six extracts for the three types of untreated biomass. One difference is 294 that the XyG epitopes show significantly different partitioning between the three biomass types 295 and the goldenrod is the only biomass exhibiting abundant XyG epitopes in the 1 M KOH 296 extract. The xylan epitopes are more abundant in the corn stover extracts and more abundantly 297 distributed into the two most severe extracts (chlorite and 4 M KOH PC) that might correspond 298 299 to "lignin-bound" xylan. Pectic polysaccharides and AG domains show different partitioning behavior between the biomass types as well, with the cell wall extracts from goldenrod 300 exhibiting the most abundant content of these classes of epitopes. Two intense MLG epitopes 301 are present in all the corn stover cell wall extracts corresponding to antibodies LAMP2H12H7 302 303 and BG1 and the abundance of these epitopes increase in extracts (particularly in the 2 mildest and the chlorite delignification) following AHP pretreatment. Weak epitope binding for both of 304 these antibodies is present in some of the poplar extracts and goldenrod extracts. These 305 observations are consistent with the primary cell wall models proposed by Carpita,¹² where for 306 307 grasses (Type II cell wall), the MLGs and GXs have a more important structural role and may act in the capacity that XyGs and pectic polysaccharides in dicots (Type I cell wall). 308 Pretreatment can conceivably alter the binding of mAbs to their glycan epitopes from the 309

309 Fredeathent can concervably after the binding of mAbs to then grycan epitopes from the 310 same extraction condition in three ways by: (1) altering the cell wall structural integrity to shift

the glycan epitope into a more easily (or more difficult) extractable category, (2) solubilizing the 311 glycan epitope during pretreatment, and (3) structurally altering the glycan epitope, for example, 312 through alkali-induced deacetylation or demethylesterification. These three modes of action are 313 used to interpret the changes associated with pretreatment. To better visualize the effects of 314 pretreatment on glycan extractability, the GP data are replotted in Figures 4 and 5 after 315 normalizing to epitope abundance per mass of original cell wall. In this representation, glycan 316 epitopes that are increased in their abundance in individual extracts after pretreatment will 317 appear to the left of the x-y line, while epitopes that are decreased will appear to the right. It can 318 be observed that for the poplar, pretreatment has very little effect on the total extractable glycans 319 in most of the six fractions. The apparent increase in the xylan epitopes in oxalate and carbonate 320 extracts suggest that the extractability of xylan by mild solvents may be enhanced by 321 pretreatment (Figure 4, subplot B). However, considering that the total content of carbohydrates 322 in these extracts are unchanged and that the xylan-specific antibodies were developed for 323 deacetylated, alkali-extracted xylans, this result likely indicates that easily extractable xylans 324 were deacetylated by pretreatment and that the abundance of deacetylated xylan epitopes 325 increase as a consequence. Another possibility is that a small fraction of the total xylan becomes 326 more easily extractable following pretreatment. The slight differences in other epitopes in the 4 327 harshest extracts (Figure 5, subplots A-C) suggests that AHP pretreatment results in only minor 328 alterations in the extractability of other major cell wall glycans indicating minimal impact on the 329 330 structural and compositional organization of the cell wall in agreement with the results in Figures 1 and 2. An exception is the xyloglucan epitopes in the 1M KOH extract (Figure 5A), which are 331 slightly improved in their extractability by pretreatment. 332

For goldenrod, the GP results show that xylan epitopes and XyG epitopes in the oxalate 333 334 extract increased considerably (Figure 4, subplots D-F); potentially as a consequence of pretreatment-induced deacetylation or by pretreatment increasing the extractability in agreement 335 with the increase in total glycan mass in the oxalate extract with pretreatment (Figure 3). Unlike 336 the poplar, the epitopes for HG backbones, RG-I/AG, and AG are decreased in both the oxalate 337 338 and carbonate extracts, indicating that these pectic polysaccharides in goldenrod are likely solubilized during AHP pretreatment. In the four most severe extracts, a number of important 339 trends are apparent (Figure 5, subplots D-F). The first is that virtually all epitopes are decreased 340 as a consequence of pretreatment in the 1M KOH extract (corresponding to alkali-soluble 341

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glycans not closely associated with lignin), while slight increases in the abundance of epitopes
are observed after pretreatment in the 4M KOH, chlorite and 4M KOH post-chlorite extracts
(corresponding to lignin-embedded glycans). These results indicate that the likely target of AHP
pretreatment for improving enzymatic hydrolysis in goldenrod is glycan (XyG and xylan)
solubilization to improve cell wall accessibility to glycolytic enzymes and minor delignification
which slightly improves the extractability of lignin-embedded polysaccharides.

The corn stover results show considerable differences in both their glycan extraction plots 348 and antibody binding profiles relative to the hybrid poplar and goldenrod (Figure 3). From the 349 glycan mass extraction profile at the top of the panel, it can be observed that, unlike the 350 goldenrod, the amounts of extractable glycans in the three most severe extracts were significantly 351 altered by pretreatment. One noticeable alteration is that the glycan epitopes in the 4M KOH 352 post-chlorite extract were shifted into the chlorite extracts after AHP pretreatment. This 353 phenomenon of the pretreatment changing the glycan extractability profile was not shown for 354 dilute acid pretreated hardwood.²⁴ This result supports the identified changes in composition 355 shown in Figure 1, and together with Figure 2, make clear that alteration of the non-cellulosic 356 glycan extractability directly impacts the glucose hydrolysis yield in the subsequent enzymatic 357 treatment. In the two mildest extracts from corn stover, the XyG and xylan epitopes were 358 increased, which could be a consequence of improving extractability or likely due to 359 deacetylation of these glycans by pretreatment (Figure 4, subplots G and H). Unlike goldenrod 360 361 and poplar, the epitopes for pectic polysaccharides were increased in the two mildest extracts, possibly indicating differences in their structural role between monocots and dicots in pectic 362 polysaccharides.¹² Additionally, the results for changes in epitope abundance as a result of 363 pretreatment for the four harshest extracts were considerably different for the corn stover than for 364 365 the goldenrod. Compared to goldenrod, where all glycan epitopes were decreased by pretreatment in the 1M KOH extract but slightly increased in the lignin-associated extracts 366 (Figure 5, subplots D-F), the corn stover glycan epitopes showed minimal change or slight 367 increases across all four extracts except in the 4M KOH post-chlorite extract, in which the lignin-368 369 embedded glycan epitopes are decreased by pretreatment (Figure 5, subplots G-I). These reductions of corn stover glycans in the harshest extracts are likely a consequence of these 370 epitopes being already removed by less harsh extractions. Increases in the 2 MLG epitopes were 371 observed with pretreatment for corn stover (Figure 3) for all extracts except the 4M KOH post 372

373 chlorite treatment.

These differing responses to AHP pretreatment between monocots and dicots have 374 important implications for structural features of the cell wall contributing to recalcitrance as well 375 as the mechanism or target of pretreatment. The composition and structure of the cell wall are 376 obviously important and many properties of the cell wall impacting recalcitrance have been 377 described in the literature including cell wall hydrophobicity,⁴⁴ porosity,^{45, 46} xylan content,⁴⁷ 378 lignin content, cross-linking, and higher order structure.⁴⁸ Jung et al.¹ noted that lignified 379 secondary cell walls were the primary obstacle hindering ruminant digestibility in dicots with 380 stem secondary xylem (*i.e.* woody biomass) the most recalcitrant, while increasing lignification 381 in grasses hinders, but does not completely inhibit digestion. The current work identified that 382 AHP pretreatment has a relatively minor impact on the hybrid poplar composition, hydrolysis 383 yields, and glycan extractability profiles. 384

Substantial work has been devoted to understanding the cell wall properties contributing 385 to ruminant digestibility of grasses and properties including ferulate content, total lignin content, 386 syringyl:guaicyl ratio of lignin monomers, and degree of arabinosylation of xylans have all been 387 linked to differences in hydrolysis yields.^{1, 48-50} Pretreatments may impact any of these afore-388 mentioned cell wall properties to improve cell wall digestibility. DeMartini et al.²⁵ found that 389 treatment of switchgrass with alkali alone to solubilize xylan (and lignin) from the cell wall was 390 sufficient to result in glucan enzymatic hydrolysis yields approaching the theoretical maximum, 391 while for hybrid poplar, chlorite delignification was necessary to improve enzymatic hydrolysis 392 significantly past alkali-only treatment. This is consistent with models for grass cell walls that 393 include alkali-labile ferulate ester cross-links between cell wall polymers as an important 394 structural feature controlling lignin integration into cell walls.⁴⁰ Our previous work identified 395 396 that lignin and ferulate removal by AHP pretreatment are important predictors of digestibility in diverse grasses.²⁹ We have previously shown that AHP pretreatment results in the destruction of 397 β -O-4 bonds in grasses²⁹ and, for example, the content of free phenolics in grass ligning may 398 enable improved alkali solubilization or potentially participate in the initiation β -O-4 scission 399 400 reactions.

Re-engineering plant cell walls for improved bioconversion outcomes is currently the
 subject of substantial research interest,⁵¹ and the findings of this work and the literature suggest
 strategies for tailoring bioenergy feedstock phenotypes to an alkaline-oxidative pretreatment

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404 process. Specifically, low initial lignin content and/or the capacity of the pretreatment to 405 effectively remove lignin are recognized as important contributors to high enzymatic hydrolysis 406 yields. As such, engineered plant phenotypes that would optimally couple to an alkaline-407 oxidative pretreatment might include decreased lignin content (without impacting plant fitness) 408 and increasing alkali-labile bonds in lignin for example through the introduction of ester cross-409 links⁵² or increasing the β-O-4 content through increasing the S/G ratio.⁵³

410

411 CONCLUSIONS

Untreated and AHP-pretreated biomass from phylogenetically diverse plants were 412 compared to understand fundamental features impacting cell wall recalcitrance. We found that 413 enzymatic hydrolysis yields, cell wall biopolymer and total mass solubilization, cell wall glycan 414 extractabilities, and glycan epitope abundances in these extracts differed significantly in their 415 response to AHP pretreatment for a woody dicot (hybrid poplar), an herbaceous dicot 416 (goldenrod), and an graminaceous monocot (corn stover). Using glycome profiling, we 417 identified different mechanisms for how AHP pretreatment overcomes cell wall recalcitrance in 418 the goldenrod versus the corn stover, while it was relatively ineffective on the poplar. For the 419 corn stover, mild alkaline-oxidative pretreatment resulted in slight delignification and 420 presumably disruption of cell wall polymer cross-linking. This had the consequence of 421 disrupting the structural integrity of the cell wall which was manifested through improved 422 423 extractability of important structural glycans including xylans, MLGs, and XyGs and presumably allowed for improved accessibility for glycolytic enzymes into the cell wall during hydrolysis. 424 425 Goldenrod was found to respond differently in the extractability profiles where all classes of glycan epitopes exhibited considerable decreases in the 1M KOH extracts following pretreatment 426 427 rather than an increase for the case of corn stover. Besides these differences, it was revealed that the pectic polysaccharides (HG, RG-I, and AG) were not only significantly more abundant in the 428 goldenrod than in the corn stover, but were solubilized by pretreatment as indicated by their 429 decrease following pretreatment in the three mildest extracts for goldenrod. For corn stover, the 430 431 pectic polysaccharides as well as the significantly more abundant MLGs showed mild increases in extractability following pretreatment indicating "loosening" from the cell wall rather than 432 solubilization. These results call attention to the important role that differences in cell wall 433 structure (e.g. MLGs in graminaceous monocots and pectic polysaccharides in herbaceous 434

dicots) and organization (*e.g.* ester cross-linking) play in setting the cell wall recalcitrance to

- 436 deconstruction by pretreatment and enzymatic hydrolysis. Thus, the pretreatment conditions that
- 437 are feedstock-specific are likely to be more effective than general approaches, and the future
- 438 work in breeding and engineering plants with cell walls designed for specific deconstruction
- approaches should make use of positive synergistic interactions between specific pretreatments
- 440 and particular cell wall features.
- 441

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- 450

451 ABBREVIATIONS

- 452 AG: arabinogalactan; AHP: alkaline hydrogen peroxide; AIR: alcohol insoluble residue; AX:
- 453 arabinoxylan; CBM: cellulose binding module; CCRC: Complex Carbohydrate Research Center;
- 454 CDTA: (1,2-cyclohexylenedinitrilo)tetraacetic acid; CS: corn stover; ELISA: enzyme-linked
- immunosorbent assay; GAX: glucuronoarabinoxylan; GM: glucomannan; GP: glycome
- 456 profiling; GX: glucuronoxylan; HGA: homogalacturonic acid; HPLC: high pressure liquid
- 457 chromatography; mAb: monoclonal antibody; MLG: mixed-linkage glucan; RG-I:
- 458 rhamnoglucuronan-I; SG: switchgrass; XyG: xyloglucan.
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460 **REFERENCES**

- 461 1. H.-J. G. Jung, D. A. Samac and G. Sarath, *Plant Sci.*, 2012, **185–186**, 65-77.
- 462 2. M. C. McCann and N. C. Carpita, *Curr. Opin. Plant Biol.*, 2008, **11**, 314-320.
- 463 3. P. Sarkar, E. Bosneaga and M. Auer, J. Exp. Bot., 2009, 60, 3615-3635.
- 464 4. Z. A. Popper, G. Michel, C. Hervé, D. S. Domozych, W. G. T. Willats, M. G. Tuohy, B.
- 465 Kloareg and D. B. Stengel, *Annu. Rev. Plant Biol.*, 2011, **62**, 567-590.

466 5. H. J. Gilbert, Plant Physiol., 2010. 6. T. Jeoh, C. I. Ishizawa, M. F. Davis, M. E. Himmel, W. S. Adney and D. K. Johnson, 467 Biotechnol. Bioeng., 2007, 98, 112-122. 468 S.-Y. Ding, Y.-S. Liu, Y. Zeng, M. E. Himmel, J. O. Baker and E. A. Bayer, Science, 7. 469 2012, 338, 1055-1060. 470 M. Josè-Estanvol and P. Puigdomènech, *Plant Physiol. Biochem.*, 2000, **38**, 97-108. 471 8. 472 9. A. Voragen, G.-J. Coenen, R. Verhoef and H. Schols, Struct. Chem., 2009, 20, 263-275. 10. A. Ebringerová, Macromol. Symp., 2005, 232, 1-12. 473 11. P. J. Harris and B. A. Stone, in Biomass Recalcitrance, Blackwell Publishing Ltd., 2009, 474 pp. 61-93. 475 12. N. C. Carpita, Annu. Rev. Plant Physiol. Plant Molec. Biol., 1996, 47, 445-476. 476 13. W. G. T. Willats, L. McCartney, W. Mackie and J. P. Knox, Plant Mol. Biol., 2001, 47, 9-477 27. 478 14. C. Somerville, S. Bauer, G. Brininstool, M. Facette, T. Hamann, J. Milne, E. Osborne, A. 479 Paredez, S. Persson, T. Raab, S. Vorwerk and H. Youngs, Science, 2004, 306, 2206-2211. 480 15. D. J. Cosgrove, Nat Rev Mol Cell Biol, 2005, 6, 850-861. 481 16. D. J. Cosgrove, Plant Physiol. Biochem., 2000, 38, 109-124. 482 17. M. Foston and A. Ragauskas, Ind. Biotechnol., 2012, 8, 191-208. 483 W. G. T. Willats, C. G. Steele-King, L. McCartney, C. Orfila, S. E. Marcus and J. P. 18. 484 485 Knox, Plant Physiol. Biochem., 2000, 38, 27-36. 19. J. P. Knox, in International Review of Cytology, ed. W. J. Kwang, Academic Press, 1997, 486 vol. 171, pp. 79-120. 487 20. I. Moller, I. Sørensen, A. J. Bernal, C. Blaukopf, K. Lee, J. Øbro, F. Pettolino, A. Roberts, 488 489 J. D. Mikkelsen, J. P. Knox, A. Bacic and W. G. T. Willats, *Plant J.*, 2007, 50, 1118-1128. 21. A. Alonso-Simón, J. B. Kristensen, J. Øbro, C. Felby, W. G. T. Willats and H. Jørgensen, 490 Biotechnol. Bioeng., 2010, 105, 509-514. 491 22. S. Pattathil, U. Avci, D. Baldwin, A. G. Swennes, J. A. McGill, Z. Popper, T. Bootten, A. 492 493 Albert, R. H. Davis, C. Chennareddy, R. Dong, B. O'Shea, R. Rossi, C. Leoff, G. Freshour, R. Narra, M. O'Neil, W. S. York and M. G. Hahn, Plant Physiol., 2010, 153, 494 514-525. 495 S. Pattathil, U. Avci, J. S. Miller and M. G. Hahn, in *T Biomass Conversion*, 2012, vol. 496 23.

497		908, pp. 61-72.
498	24.	J. D. DeMartini, S. Pattathil, U. Avci, K. Szekalski, K. Mazumder, M. G. Hahn and C. E.
499		Wyman, Energy Environ. Sci., 2011, 4, 4332-4339.
500	25.	J. DeMartini, S. Pattathil, J. S. Miller, H. Li, M. G. Hahn and C. E. Wyman, Energy
501		Environ. Sci., 2013.
502	26.	N. Mosier, R. Hendrickson, N. Ho, M. Sedlak and M. R. Ladisch, Bioresour. Technol.,
503		2005, 96 , 1986-1993.
504	27.	G. Banerjee, S. Car, T. Liu, D. L. Williams, S. L. Meza, J. D. Walton and D. B. Hodge,
505		Biotechnol. Bioeng, 2012, 109, 922-931.
506	28.	G. Banerjee, S. Car, J. Scott-Craig, D. Hodge and J. Walton, Biotechnol. Biofuels, 2011,
507		4 , 16.
508	29.	M. Li, C. Foster, S. Kelkar, Y. Pu, D. Holmes, A. Ragauskas, C. Saffron and D. Hodge,
509		Biotechnol. Biofuels, 2012, 5, 38.
510	30.	T.K. Sato, T. Liu, L.S. Parreiras, D. L. Williams, D. J. Wohlbach, B. D. Bice, I. M. Ong,
511		R. J. Breuer, L. Qin, D. Busalacchi, S. Deshpande, C. Daum, A. P. Gasch, D. B. Hodge,
512		Appl. Environ. Microbiol., 2014. 80, 540-554.
513	31.	B. S. Donohoe, S. R. Decker, M. P. Tucker, M. E. Himmel and T. B. Vinzant, Biotechnol.
514		Bioeng., 2008, 101, 913-925.
515	32.	M. A. T. Hansen, J. B. Kristensen, C. Felby and H. Jørgensen, Bioresour. Technol., 2011,
516		102 , 2804-2811.
517	33.	A. Sluiter, B. Hames, R. Ruiz, C. Scarlata, J. Sluiter and D. Templeton, NREL Analytical
518		Procedure, 2004, National Renewable Energy Laboratory, Golden, CO.
519	34.	S. Pattathil, U. Avci, J. S. Miller and M. G. Hahn, in Biomass Conversion: Methods and
520		Protocols, ed. M. E. Himmel, Humana Press, 2012, vol. 908, pp. 61-72.
521	35.	U.S. Department of Energy, U.S. Billion-Ton Update: Biomass Supply for a Bioenergy
522		and Bioproducts Industry., 2011. R.D. Perlack and B.J. Stokes (Leads), ORNL/TM-
523		2011/224. Oak Ridge National Laboratory, Oak Ridge, TN. 227p.
524	36.	V. H. Dale, K. L. Kline, L. L. Wright, R. D. Perlack, M. Downing and R. L. Graham,
525		<i>Ecol. Appl.</i> , 2011, 21 , 1039-1054.
526	37.	D. Tilman, J. Hill and C. Lehman, Science, 2006, 314, 1598-1600.
527	38.	C. Lapierre, D. Jouin and B. Monties, Phytochemistry, 1989, 28, 1401-1403.

528	39.	R. J. Stoklosa and D. B. Hodge, Ind. Eng. Chem. Res., 2012, 51, 11045-11053.
529	40.	J. Ralph, S. Guillaumie, J. H. Grabber, C. Lapierre and Y. Barrière, C. R. Biol., 2004, 327,
530		467-479.
531	41.	J. M. Gould, Biotechnol. Bioeng., 1985, 27, 893-896.
532	42.	S. Pattathil, T. Saffold, L. Gallego-Giraldo, M. O'Neill, W. S. York, R. A. Dixon and M.
533		G. Hahn, Ind. Biotechnol., 2012, 8, 217-221.
534	43.	M. C. Jarvis, W. Forsyth and H. J. Duncan, Plant Physiol., 1988, 88, 309-314.
535	44.	S. Nakagame, R. P. Chandra, J. F. Kadla and J. N. Saddler, Biotechnol. Bioeng., 2011,
536		108 , 538-548.
537	45.	W. R. Grous, A. O. Converse and H. E. Grethlein, Enzyme Microb. Technol., 1986, 8,
538		274-280.
539	46.	D. N. Thompson, HC. Chen and H. E. Grethlein, Bioresour: Technol., 1992, 39, 155-
540		163.
541	47.	J. D. McMillan, in Enzymatic Conversion of Biomass for Fuels Production, American
542		Chemical Society, 1994, vol. 566, ch. 15, pp. 292-324.
543	48.	Y. Zhang, T. Culhaoglu, B. Pollet, C. Melin, D. Denoue, Y. Barrière, S. p. Baumberger
544		and V. R. Méchin, J. Agric. Food. Chem., 2011, 59, 10129-10135.
545	49.	M. A. Bernard Vailhé, G. J. Provan, L. Scobbie, A. Chesson, M. P. Maillot, A. Cornu and
546		J. M. Besle, J Agric. Food. Chem., 2000, 48, 618-623.
547	50.	A. Mohagheghi, K. Evans, YC. Chou and M. Zhang, Appl. Biochem. Biotechnol., 2002,
548		98-100 , 885-898.
549	51.	R. G. Ong, S. P. Chundawat, D. B. Hodge,, S. Keskar, and B. E. Dale, in Plants and
550		BioEnergy, Springer, 2014, ch.14, pp. 231-253.
551	52.	J. H. Grabber, J. Ralph, and R. D. Hatfield RD, J. Agric. Food Chem. 1998, 46, 2609-
552		2614.
553	53.	J. J. Stewart, T. Akiyama, C. Chapple, J. Ralph, S. D. Mansfield, Plant Physiol. 2009,
554		150 , 621-635
555		
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Table 1: Composition of untreated biomass. Neutral polysaccharides, lignin, and ash were determined according to NREL Laboratory Analytical Protocol NREL/TP-510-42619. Acetyl content was determined by HPLC following two-stage acid hydrolysis. Uronic acids were assayed enzymatically (K-Uronic, Megazyme). Extractives were determined gravimetrically following sequential extraction with ethanol, methanol-chloroform, and acetone. Compositions are reported on a whole sample basis rather than an extractives-free basis. Errors represent data range of duplicate measurements.

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Figure 1. Compositional changes (extractives-free basis) associated with AHP pretreatment with increasing H_2O_2 loading showing solubilization of total cell wall mass (A), xylan (B), and lignin (C). The results indicate substantially different responses to AHP pretreatment by the three biomass types in all three categories. The basis for these differences arising from the structural organization of cell wall glycans are further explored by glycome profiling.

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Figure 2. Enzymatic hydrolysis yields of cell wall glucan to glucose in untreated and AHP pretreated biomass with increasing H_2O_2 loading for poplar, goldenrod, and corn stover with hydrolysis (Cellic CTec2) for 24 h and 72 h.

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576 **Figure 3.** Glycome profiling of hybrid poplar, goldenrod and corn stover biomass samples before and after AHP pretreatment (12.5% H₂O₂ loading). Sequential cell wall extracts were made from 577 578 untreated and pretreated biomass samples using increasingly harsh reagents as explained in materials and methods. Various extracts obtained were ELISA screened using 155 mAbs 579 580 directed against most major plant cell wall glycans (See Supplemental Table S1). The resulting binding response data are represented as heatmaps with yellow-red-black scale indicating the 581 strength of the ELISA signal (yellow, red and dark-blue colors depict strong, medium, and no 582 binding, respectively). The mAbs are grouped based on the cell wall glycans they recognize as 583 584 depicted in the panel at right hand side of the figure. The actual amounts of materials extracted out at each extraction condition are depicted as bar graph at the top of heatmaps with color codes 585 for reagents used for extraction. 586

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588 Figure 4: Comparison of the changes in the relative abundance of three glycan epitope categories due to AHP pretreatment from the oxalate (blue dots) and carbonate (red dots) extracts 589 in glycome profiling, for poplar (A, B and C, for xyloglucans, xylans and pectic 590 polysaccharides), goldenrod (D, E and F for xyloglucans, xylans and pectic polysaccharides) and 591 corn stover (G, H and I for xyloglucans, xylans and pectic polysaccharides). Data are replotted 592 from Figure 3, but are normalized to represent mAb binding strength per mass of original AIR to 593 594 make absolute values comparable between conditions and samples. The red dash line represents x = y and denotes the case where the abundance of these glycan epitopes was unchanged after 595 AHP pretreatment in these extracts. Data points above and below the dash lines represent 596 increased or decreased abundance of the glycan epitopes appearing in each extract, respectively. 597 598 **Figure 5:** Comparison of the changes in the relative abundance of three glycan epitope 599 categories due to AHP pretreatment from each of the four harshest extracts including 1M KOH 600 (green triangles), 4M KOH (blue diamonds), chlorite (red squares), and 4M KOH PC (purple 601

crosses) in glycome profiling. Data are normalized to represent relative abundance per massoriginal AIR.

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606	Table 1.
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	Hybrid Poplar (wt/wt %)	Goldenrod (wt/wt %)	Corn Stover (wt/wt %)
Glc	48.3±2.0	27.1±0.8	38.4±0.3
Xyl+Man+Gal	16.9±0.6	13.6±0.8	25.2±0.2
Ara	1.33±0.07	2.96±0.03	3.92 ± 0.02
Acetyl	4.14±0.09	2.59±0.06	3.20±0.10
Total Uronic Acids	1.88 ± 0.10	4.75±0.03	0.88 ± 0.02
Lignin (Klason)	20.85±1.1	19.92±0.7	12.57±1.2
Extractives	5.79±0.22	23.5±0.23	12.0±0.74
Ash	1.95 ± 0.48	6.08±1.02	3.03±0.29

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Figure 1: Compositional changes (extractives-free basis) associated with AHP pretreatment with increasing H₂O₂ loading showing solubilization of total cell wall mass (A), xylan (B), and lignin (C). The results indicate substantially different responses to AHP pretreatment by the three biomass types in all three categories. The basis for these differences arising from the structural organization of cell wall glycans are further explored by glycome profiling.
 789x230mm (96 x 96 DPI)







Figure 3: Glycome profiling of hybrid poplar, goldenrod and corn stover biomass samples before and after AHP pretreatment (12.5% H₂O₂ loading). Sequential cell wall extracts were made from untreated and pretreated biomass samples using increasingly harsh reagents as explained in materials and methods. Various extracts obtained were ELISA screened using 155 mAbs directed against most major plant cell wall glycans (See supplemental Table S2). The resulting binding response data are represented as heatmaps with yellow-red-black scale indicating the strength of the ELISA signal (yellow, red and dark-blue colors depict strong, medium, and no binding, respectively). The mAbs are grouped based on the cell wall glycans they recognize as depicted in the panel at right hand side of the figure. The actual amounts of materials extracted out at each extraction condition are depicted as bar graph at the top of heatmaps with color codes for reagents used for extraction.

544x806mm (96 x 96 DPI)



Comparison of the changes in the relative abundance of three glycan epitope categories due to AHP pretreatment from the oxalate (blue dots) and carbonate (red dots) extracts in glycome profiling, for poplar (A, B and C, for xyloglucans, xylans and pectic polysaccharides), goldenrod (D, E and F for xyloglucans, xylans and pectic polysaccharides) and corn stover (G, H and I for xyloglucans, xylans and pectic polysaccharides). Data are replotted from Figure 3, but are normalized to represent mAb binding strength per mass of original AIR to make absolute values comparable between conditions and samples. The red dash line represents x = y and denotes the case where the abundance of these glycan epitopes was unchanged after AHP pretreatment in these extracts. Data points above and below the dash lines represent increased or decreased abundance of the glycan epitopes appearing in each extract, respectively. 438x365mm (96 x 96 DPI)



Comparison of the changes in the relative abundance of three glycan epitope categories due to AHP pretreatment from each of the four harshest extracts including 1M KOH (green triangles), 4M KOH (blue diamonds), chlorite (red squares), and 4M KOH PC (purple crosses) in glycome profiling. Data are normalized to represent relative abundance per mass original AIR. 410x337mm (96 x 96 DPI)