

1 **Structural imaging of native cryo-preserved secondary cell walls**
2 **reveals the presence of microfibrils and their formation requires**
3 **normal cellulose, lignin and xylan biosynthesis.**

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30 1 Abstract

31 The woody secondary cell walls of plants are the largest repository of renewable carbon biopolymers
32 on the planet. These walls are made principally from cellulose and hemicelluloses and are impregnated
33 with lignin. Despite their importance as the main load bearing structure for plant growth, as well as
34 their industrial importance as both a material and energy source, the precise arrangement of these
35 constituents within the cell wall is not yet fully understood. We have adapted low temperature scanning
36 electron microscopy (cryo-SEM) for imaging the nanoscale architecture of angiosperm and
37 gymnosperm cell walls in their native hydrated state. Our work confirms that cell wall macrofibrils,
38 cylindrical structures with a diameter exceeding 10 nm, are a common feature of the native hardwood
39 and softwood samples. We have observed these same structures in *Arabidopsis thaliana* secondary cell
40 walls, enabling macrofibrils to be compared between mutant lines that are perturbed in cellulose,
41 hemicellulose and lignin formation. Our analysis indicates that the macrofibrils in Arabidopsis cell
42 walls are dependent upon the proper biosynthesis, or composed, of cellulose, xylan and lignin. This
43 study establishes that cryo-SEM is a useful additional approach for investigating the native nanoscale
44 architecture and composition of hardwood and softwood secondary cell walls and demonstrates the
45 applicability of Arabidopsis genetic resources to relate fibril structure with wall composition and
46 biosynthesis.

47 2 Introduction

48 The majority of carbon in terrestrial biomass is stored in forests as wood (Ramage et al., 2017, Pan et
49 al., 2011). The current classification system distinguishes two types of timber. Wood from Angiosperm
50 trees is known as hardwood and the wood made by Gymnosperm species is described as softwood
51 (Ramage et al., 2017). Despite significant differences in tissue organisation and chemical composition,
52 both these types of timber are almost entirely formed from plant secondary cell walls – an extracellular
53 matrix made primarily from cellulose, lignin and hemicelluloses (Schweingruber, 2007). Considering
54 the ecological and industrial importance of wood and other cell wall materials, our knowledge of the
55 exact arrangement of these polymers in the cell wall remains poor. A better understanding of the
56 molecular architecture and ultrastructure of cell walls is needed to describe the complex spatio-
57 temporal deposition pattern of the cell wall polymers. This may contribute to the development of more
58 efficient biofuel feedstocks (Loque et al., 2015), to the improvement in our understanding of novel
59 biomaterials such as nanocellulose (Jarvis, 2018), and to applications such as advanced approaches for
60 the use of timber in the construction industry (Ramage et al., 2017)

61 Cellulose is the main constituent of plant cell walls (Pauly and Keegstra, 2008). At the molecular level,
62 cellulose has a simple repeating structure of β -1,4-linked glucopyranosyl residues. These glucan chains
63 coalesce to form a crystalline cellulose microfibril. The exact structure of the microfibril is unknown,
64 however, it has been suggested the elementary microfibril consists of 18 or 24 individual glucan chains
65 (Gonneau et al., 2014, Hill et al., 2014, Turner and Kumar, 2017). Individual cellulose microfibrils
66 associate to form larger order structures known as macrofibrils (Niklas, 2004). In plant primary cell
67 walls this close-contact association may be limited to selected parts of the microfibril which is proposed
68 to lead to formation of so-called biomechanical hotspots (Cosgrove, 2014). A range of imaging and
69 spectroscopic techniques has been used to investigate cellulose macrofibrils in secondary cell walls, as
70 reviewed by (Purbasha et al., 2009), but due to technical challenges the precise structure in native,
71 unprocessed, hydrated secondary cell walls remains poorly described. Lignin is the main non-
72 polysaccharide component of both hardwood and softwood and is made by coupling of monolignol
73 radicals in secondary cell walls. Three main monolignols exist in plants, which, once turned into
74 chemical radicals by the activity of laccases and peroxidases, can couple in a random manner to form

75 a lignin polymer made from guaiacyl (G), syringyl (S), and p-hydroxyphenyl (H) units (Ralph et al.,
76 2004). The monolignol composition of hardwood and softwood differs, with the former consisting of
77 predominantly S and G units and the latter being made almost solely from G units (Vanholme et al.,
78 2010). The process of lignification is important for wood mechanical properties. Arabidopsis mutant
79 plants with reduced lignin content or altered monolignol composition often have collapsed xylem
80 vessels and can be severely dwarfed (Bonawitz and Chapple, 2010). Lignin is proposed to associate
81 with cell wall polysaccharides to form the recalcitrant matrix (Terrett and Dupree, 2019).

82 Xylan and galactoglucomannan are the principal hemicelluloses in hardwood and softwood. Xylan is
83 a polymer of β -1,4-linked xylopyranosyl residues and is the main hemicellulose in hardwood but is
84 also present in softwood (Scheller and Ulvskov, 2010). Hardwood and softwood xylans carry α -1–2
85 linked glucuronic acid (GlcA) branches which can be methylated on carbon 4 leading to formation of
86 4-O-Methyl-glucuronic acid (MeGlcA) (Scheller and Ulvskov, 2010). In addition to GlcA and
87 MeGlcA (together, [Me]GlcA) decorations, hardwood xylan hydroxyls are acetylated on carbon 2,
88 carbon 3 or both carbons of the monomer. The softwood xylan, in addition to the MeGlcA branches,
89 carries α -1,3-linked arabinofuranosyl decorations (Scheller and Ulvskov, 2010, Busse-Wicher et al.,
90 2016b). The presence of [Me]GlcA branches on xylan is important for the maintenance of biomass
91 recalcitrance (Lyczakowski et al., 2017) and, together with acetylation in hardwood and arabinose
92 decorations in softwood, these substitutions are mostly distributed with an even pattern on xylosyl units
93 (Bromley et al., 2013, Busse-Wicher et al., 2014, Busse-Wicher et al., 2016b, Martinez-Abad et al.,
94 2017). This so-called ‘compatible’ patterning of xylan substitutions is thought to allow the hydrogen
95 bonding between xylan, in a two-fold screw conformation, and the hydrophilic surface of the cellulose
96 microfibril (Busse-Wicher et al., 2016a, Simmons et al., 2016, Grantham et al., 2017).
97 Galactoglucomannan (GGM) is the main hemicellulose in softwood (Scheller and Ulvskov, 2010) but
98 is also present in hardwood xylem. GGM has a backbone formed from both β -1,4-linked mannosyl and
99 glucosyl residues with some mannosyl residues substituted by an α -1,6-linked galactosyl branch. The
100 GGM backbone can also be acetylated. The arrangement of mannose and glucose units in softwood
101 GGM is thought to be random, but a recently described regular structure GGM found in Arabidopsis
102 mucilage was proposed to bind to both the hydrophilic and hydrophobic surface of the cellulose
103 microfibril (Yu et al., 2018). *In vitro* studies using TEM and 1D ^{13}C NMR indicate that a range of
104 branched and unbranched mannan and glucomannan structures can interact with bacterial cellulose
105 (Whitney et al., 1998). Softwood GGM is also proposed to interact with the cellulose microfibril
106 (Terashima et al., 2009) and recent evidence demonstrates that it can form covalent linkages with lignin
107 (Nishimura et al., 2018).

108 Although we now have a better understanding of secondary cell wall composition and the nature of the
109 interactions between its main constituents, a picture of the ultrastructural assembly of wall polymers
110 into a secondary cell wall matrix is not yet complete. Solid state NMR (ssNMR) analysis has been
111 applied extensively to the study of polymer interactions in both primary and secondary walls. This, for
112 example, provided evidence that in dried primary wall samples from Arabidopsis, pectin and
113 xyloglucan may be interacting with the cellulose microfibril (Dick-Perez et al., 2011). Analysis of
114 hydrated secondary cell wall of Arabidopsis with solid state NMR indicated that xylan is likely to
115 interact with the hydrophilic surface of the cellulose microfibril as a two-fold screw (Simmons et al.,
116 2016, Grantham et al., 2017). Recent ssNMR analysis indicates that in dried cell walls of grasses, xylan
117 is likely to interact with lignin (Kang et al., 2019). Despite providing excellent insights into the
118 proximity of different cell wall components ssNMR cannot provide information about the assembly of
119 these constituents into higher order structures. Some insights into this process have been achieved with
120 other techniques. This includes application of vibrational microspectroscopy techniques such as FT-IR
121 and Raman to study the orientation of cellulose and other cell wall components in the matrix, as

Cell wall macrofibrils are composed of cellulose, xylan and lignin

122 reviewed by (Gierlinger, 2018). AFM has been applied to the study of cell wall matrix assembly, but
123 the work has been focused on primary cell walls (Cosgrove, 2014) and only recent advances allowed
124 nanoscale resolution imaging of dried spruce secondary cell walls (Casdorff et al., 2017). Moreover,
125 insights into the assembly of cellulose microfibrils in wood walls of conifers (Fernandes et al., 2011)
126 and dicots (Thomas et al., 2014) have been obtained using wide-angle X-ray scattering (WAXS) and
127 small-angle neutron scattering (SANS).

128 In addition to these various approaches, other studies have attempted to use scanning electron
129 microscopy (SEM) to study the structure of plant cell walls. Low temperature SEM (cryo-SEM), in
130 which the sample is rapidly frozen and then maintained cold during imaging, has been used to study
131 the collapse of pine needle tracheid cell walls upon prior dehydration (Cochard et al., 2004) and to
132 visualise the bulging of root hairs in the *kojak* (cellulose synthase-like) mutant (Favery et al., 2001).
133 Additionally, higher magnification cryo-SEM has been used to visualise cell walls of wheat awns
134 (Elbaum et al., 2008). Some awn cell walls exhibit structural differences that are dependent upon the
135 level of hydration and cryo-SEM revealed extensive layering within the wall, however, the technique
136 was not further optimised to investigate individual fibrils. Field emission (FE) SEM techniques were
137 effectively used to study the alignment of cellulose microfibrils in *Arabidopsis* hypocotyls (Refregier
138 et al., 2004), roots (Himmelspach et al., 2003) and stems (Fujita et al., 2013). FE-SEM has also been
139 applied to investigate wood structure, including observations of microfibril alignment in fixed cell
140 walls of fir tracheids (Abe et al., 1997) and lignin distribution in spruce tracheids (Fromm et al., 2003).
141 Importantly, FE-SEM analysis of dehydrated pine and poplar wood suggests that secondary cell walls
142 of these species contain macrofibrils – cylindrical fibrillar structures with a diameter of up to 60 nm,
143 which presumably comprise of bundles of elementary cellulose microfibrils (Donaldson, 2007).
144 Moreover, the diameter of these macrofibrils was observed to increase with increasing lignification,
145 suggesting that the macrofibrils may be formed from association of lignin and cell wall
146 polysaccharides. This analysis was extended further to wood from Ginkgo where the FE-SEM was
147 combined with density analysis to propose a model of macrofibril formation based on cellulose, GGM,
148 xylan and lignin interaction (Terashima et al., 2009).

149 It has been suggested that some of the treatments used in preparation of the FE-SEM cell wall samples
150 have little impact on the microfibril arrangement and that the technique may provide a true
151 representation of native (unprocessed) cell wall features (Marga et al., 2005). The FE-SEM techniques
152 applied to secondary cell wall samples, however, included additional steps such as (i) fixation and
153 exposure to organic solvents (ii) a thermal treatment that may result in some degree of wall degradation
154 (Fromm et al., 2003) and (iii) a thick coating of heavy metal which may impact upon the resolution
155 (Donaldson, 2007), raising questions about the effect these may have on interpretation of the wall
156 structure. Visualisation of native, hydrated, secondary cell walls with environmental FE-SEM has been
157 challenging and the resolution of obtained images has been low (Donaldson, 2007). We present here a
158 technique for the analysis of native, fully-hydrated, secondary cell wall material from angiosperm and
159 gymnosperm plant species using cryo-SEM. The use of an ultrathin 3 nm platinum film, together with
160 cryo-preservation at high vacuum, enabled us to demonstrate that cell wall macrofibrils are a common
161 feature in all types of native secondary cell wall material analysed. Importantly, we were able to detect
162 the presence of macrofibrils in *Arabidopsis thaliana* vessel secondary cell walls. This allowed us to
163 make use of the readily available cell wall-related genetic resources, revealing *Arabidopsis* microfibril
164 diameter to be dependent upon cellulose, xylan and lignin.

165

166 3 Materials and Methods

167 3.1 Plant material

168 *Picea abies*, (spruce) one-year old branch was acquired from 30-50cm tall potted plants grown
169 outdoors purchased from Scotsdale (Great Shelford, Cambridgeshire, UK). *Ginkgo biloba*, (Ginkgo)
170 material, consisting of the narrow ends of branches of diameter approximately 3-5 mm was obtained
171 from 15 year old trees trees grown at the Cambridge University Botanic Garden. For both spruce and
172 Ginkgo, samples from two individuals were analysed.

173 Hybrid aspen (*Populus tremula* x *Populus tremuloides*, clone T89), referred to as poplar in the text,
174 was grown *in vitro* (20°C, with a 16-h light, 8-h dark photoperiod, with illumination at 85
175 microeinsteins.m⁻².s⁻¹) during 76 to 80 days after micro-propagation on 1/2MS media with vitamins
176 (Duchefa M0222), 1% sucrose, 0.7% Agar. Samples from three individuals were analysed. For field
177 grown poplar (*Populus tremula*), material was obtained from one year old branches of two
178 approximately 30 year old individuals grown at the Cambridge University Botanic Garden.

179 *Arabidopsis thaliana* (*Arabidopsis*) Columbia-0 ecotype plants were grown in a cabinet maintained at
180 21 °C, with a 16-h light, 8-h dark photoperiod. Stem material was collected from 7-week-old plants.
181 Mutant insertion lines described in published work were used in this study. Specifically, Col-0 ecotype
182 *irx3-7* plants (Simmons et al., 2016, Kumar and Turner, 2015), representing a mutant allele of CESA7,
183 *irx9-1* (Brown et al., 2005), *irx10-1* (Brown et al., 2005), *esk1-5* (Lefebvre et al., 2011, Grantham et
184 al., 2017), *4cl1-1* (Vanholme et al., 2012), *lac4-2* (Berthet et al., 2011) and *cs1a2-1cs1a3-2cs1a9-1*
185 (Goubet et al., 2009) were studied. Mutants of *IRX1* and *IRX5* gene were in Ler ecotype (Taylor et al.,
186 2003). Plants were analysed alongside the Col-0 or Ler wild type (WT) material. For each genotype
187 three individuals were analysed.

188 3.2 Cryo-SEM sample preparation and imaging

189 Fresh stems of 7 week old *Arabidopsis* plants were prepared for imaging as outlined in Supplementary
190 Material Figure S1. Firstly, 1 cm length sections were cut from the bottom part of the stems and
191 mounted vertically in recessed stubs containing a cryo glue preparation consisting of a 3:1 mixture of
192 Tissue-Tec (Scigen Scientific, USA) and Aquadog colloidal graphite (Agar Scientific, Stansted, UK)
193 (see steps 1 to 4 on Figure S1). Stem sections were immediately (within 5 minutes of harvest) plunge
194 frozen in liquid nitrogen slush (step 5 on Figure S1), transferred under vacuum, fractured and then
195 coated with 3 nm of platinum (step 6 on Figure S1) using a PT3010T cryo-apparatus fitted with a film
196 thickness monitor (Quorum Technologies, Lewes, UK). The short time between freezing and
197 harvesting serves to prevent drying of the sample where only the exposed surface, not the fractured
198 face, is expected to exhibit some water loss during the short time it is exposed to air. Finally, fractured
199 stems were imaged using a Zeiss EVO HD15 Scanning Electron Microscope (step 7 on Figure S1) and
200 maintained at -145 °C using a Quorum cryo-stage assembly. The electron source is a Lanthanum
201 Hexaboride HD filament. Images were acquired using a secondary electron detector and an accelerating
202 voltage of between 5 and 8 kV with a working distance between 4 and 6 mm. Quantification of the
203 width of cell wall macrofibrils was performed using ImageJ software (Schneider et al., 2012). For the
204 measurements of macrofibril width between 25 and 50 macrofibrils were selected at random on each
205 image analysed (Figure 1a). To quantify the width a line was drawn parallel to the fibril axis. The
206 length of a second line, perpendicular to the fibril axis line and across the width of the macrofibril, was
207 quantified as the macrofibril width (Figure 1b). Each fibril width measurement was standardised for
208 the platinum layer applied during the coating process by subtracting the width of the standardised coat
209 from the original measurement. Imaging without the cryo-preservation was performed by visualising
210 hand sectioned platinum coated specimens with the stage maintained at room temperature. For
211 preparation of these samples all freezing steps were omitted.

212 **3.3 Sampling and statistical analysis**

213 For spruce, Ginkgo and field grown poplar, stem sections were taken from two individual trees and
214 150 macrofibrils were measured from three tracheids/vessels that had each been coated with platinum
215 separately. Imaging of poplar was performed in technical triplicate from three *in vitro* grown trees and
216 150 poplar macrofibrils were measured from three separately coated vessels as for the gymnosperm
217 samples. For Arabidopsis, cryo-SEM imaging of vessels was carried out on three biological replicates,
218 each from separate individuals. 150 macrofibril diameters were measured across the three individuals.

219 Statistical analysis was performed using packages available with R software (Team, 2014). Statistical
220 tests, either Student's T test or ANOVA, used to compare average measurements for samples are
221 defined in Figure legends. The variance between each pairwise combination was estimated to be similar
222 with Levene's test.

223 **4 Results**

224 **4.1 Softwood and hardwood secondary cell walls contain macrofibrils**

225 In order to investigate and compare the nanoscale architecture of gymnosperm and angiosperm cell
226 walls we analysed stem sections taken from spruce, Ginkgo and poplar using cryo-SEM. Stems were
227 placed in the SEM specimen stub and immediately frozen in nitrogen slush, fractured and then coated
228 with platinum, before being passed in to the SEM chamber for imaging. Nitrogen slush is a suspension
229 of solid nitrogen that enables high freezing rates, greatly reducing the Leidenfrost effect during plunge
230 freezing and thus minimising structural damage (Sansinena et al., 2012). The fine grain size attributed
231 to platinum sputtering allows small and densely packed objects to be resolved. This rapid sample
232 preparation protocol serves to better maintain sample hydration levels and native structures for optimal
233 EM imaging in a high vacuum environment.

234 We first investigated whether our cryo-SEM protocol gave comparable results to the previous FE-SEM
235 analysis of both softwood and hardwood secondary cell walls (Donaldson, 2007). To examine if
236 macrofibrils are found in natively hydrated, non-pretreated cell walls, cryo-SEM imaging was
237 performed on unprocessed, frozen softwood and hardwood samples. For observing gymnosperm cell
238 wall architecture, we first prepared softwood samples from spruce and used a low magnification to see
239 an overview of stem cross-section (Figure 2a) and tracheid structure (Figure 2b). The inner part of the
240 stem cross section was composed of densely packed xylem tracheids, each surrounded by cell walls.
241 To investigate the appearance of the secondary cell walls, higher magnification images of these parts
242 of tracheid cells were acquired. This enabled us to observe that the tracheid cell walls contain fibrous
243 structures which frequently assembled into larger aggregates (Figure 2c and 2d, red arrows). After a
244 further increase in magnification, individual fibrils became resolvable (Figure 2e and 2f) and their
245 diameter was found to exceed the 3 nm diameter calculated for a single softwood elementary
246 microfibril (Fernandes et al., 2011). Therefore the observed fibrils, if composed of cellulose, represent
247 a higher order structure that fits the description of a "macrofibril" (Niklas, 2004, Donaldson, 2007).
248 Similarly to spruce stem, sections from another gymnosperm, the Ginkgo, were also observed to
249 contain macrofibrils (Figure S2). These data show that, in line with previously reported SEM imaging
250 of dried, processed plant material (Donaldson, 2007, Terashima et al., 2009), the native, hydrated cell
251 walls of spruce and Ginkgo also contain macrofibrils. Therefore, these structures may contribute to the
252 higher order assembly of native gymnosperm cell walls.

253 We extended the analysis to the model hardwood species, poplar. Vessels, a distinct cell type of
254 hardwood xylem, were clearly visible using low magnification (Figure 3a and 3b). In addition to the

255 vessels, xylem fibre cells were also observed (Figure 3b; red and yellow arrows for vessels and fibre
256 cells respectively). For some cells we were able to observe spiral thickenings which were preserved
257 during sample preparation and extended above the surface of the fracture plane (Figure 3b). We focused
258 upon the vessel cell walls which showed clearly visible fibrous structures at a vessel-to-vessel boundary
259 (Figure 3c). Analysis of vessel cell walls at a higher magnification revealed a clear presence of
260 macrofibril structures, similar to those observed in spruce, in the poplar samples (Figure 3d and 3e).
261 To investigate the dimensions of the macrofibrils we measured their diameter in poplar and spruce
262 (Figure 3f). Our measurements are broadly similar to those reported in a previous study (Donaldson,
263 2007). We carried out comparative analysis of macrofibril diameter between hardwood and softwood
264 by measuring 150 individual macrofibrils in poplar, spruce and Ginkgo. While the diameter of spruce
265 and Ginkgo macrofibrils was not significantly different (Figure S3), the diameter of macrofibrils in
266 poplar secondary cell walls was significantly smaller than that of spruce macrofibrils (Figure 3f).
267 Spruce and Ginkgo were grown in the field while poplar samples were obtained from *in vitro* grown
268 plants. To control for this difference in growth conditions we also analysed samples from field grown
269 poplar trees. There was no statistically significant difference in the macrofibril diameter between the
270 two poplar samples (Figure S4). For both hardwood and softwood we observed variation in the
271 macrofibril diameter. This may reflect biological differences or may be a result of technical challenges
272 associated with macrofibril width measurement.

273 **4.2 Arabidopsis secondary cell walls macrofibrils contain a cellulose scaffold**

274 To further evaluate the nanoscale architecture of plant cell walls and identify possible constituents of
275 the cell wall macrofibrils, the high magnification cryo-SEM imaging was used to analyse wild type
276 (WT) Arabidopsis secondary cell walls (Figure 4). The initial analysis investigated the structure of WT
277 xylem vessels (Figure 4a and 4b). Sets of vessel bundles were detected and, using higher magnification,
278 fibrous structures similar to those observed in spruce and poplar were also visible in the fractured
279 Arabidopsis material. The width of WT Arabidopsis macrofibrils was comparable to that of poplar
280 macrofibrils but not spruce and suggests Arabidopsis macrofibrils could be used as a good structural
281 model for hardwoods (Figure S3, S4). Despite the use of ultra-thin platinum coating, the use of SEM
282 without the cryo-preservation steps did not allow us to observe the Arabidopsis macrofibrils with good
283 resolution (Figure S5) highlighting the critical importance of sample cryo-preservation to resolve a
284 native cell wall ultrastructure.

285 Based on the data available in the literature, we hypothesized that the macrofibrils may be mostly
286 composed of cellulose (Fahlen and Salmen, 2002, Donaldson, 2007). To investigate this, and to
287 understand the nature of these macrofibrils further, we performed a comparative analysis between WT
288 vessel cell walls (Figure 4c) and a commercially available fibrous cellulose standard (Figure 4d)
289 extracted from cotton linters and consisting of 99% pure cellulose (Sczostak, 2009). In this experiment,
290 clear individual fibrils with distinct bright termini were observed in both samples indicating that the
291 vessel wall macrofibrils have a similar appearance to the cellulose fibrils present in this polysaccharide
292 standard. To determine whether these macrofibrils are dependent upon the proper production of
293 cellulose, the morphology of WT Arabidopsis vessel cell walls (Figure 4e and 4g) was compared to
294 that of the *irx3* mutant (Figure 4f and 4h). IRX3 is one of three CESA proteins that make up the
295 secondary wall cellulose synthase complex and *irx3* plants are almost completely devoid of cellulose
296 in their secondary cell walls, but not primary cell walls (Taylor et al., 1999). As previously reported,
297 *irx3* plants had collapsed vessels (Figure S6), since secondary cell wall cellulose contributes to vessel
298 wall strength (Turner & Somerville, 1997). Interestingly, the *irx3* stems lacked the fibrous structures
299 in their vessel secondary cell walls and, in contrast to WT, the *irx3* cell walls were formed from a
300 largely amorphous matrix (Figure 4f). It is likely that this matrix is composed of xylan and lignin,

Cell wall macrofibrils are composed of cellulose, xylan and lignin

301 which can still be deposited in the secondary cell wall in the absence of IRX3 activity (Takenaka et al.,
302 2018). Some structures which may resemble cellulose fibrils were present in the primary cell walls of
303 *irx3* plants (Figure S6). To further support these observations we analysed the cell walls of plants
304 mutated in *IRX1* and *IRX5*, encoding other members of the secondary cell wall cellulose complex
305 (Figure S7). Similar to *irx3*, the *irx1* and *irx5* plants lacked fibril-type structures in their cell walls.
306 Taken together, the data show that macrofibril formation is dependent upon cellulose production.

307 **4.3 Reduction in cell wall xylan and lignin, but not in galactoglucomannan content decreases** 308 **the dimensions of Arabidopsis macrofibrils**

309 To investigate the role of xylan in macrofibril formation, cryo-SEM was used to visualise the secondary
310 walls from *irx9*, *irx10* and *eskl* Arabidopsis plants (Figure 5a and S8, 5b and S9, 5c and S10). IRX9
311 and IRX10 are required for proper xylan synthesis and mutations in the corresponding genes lead to
312 cell wall weakening and collapse of xylem vessels in the Arabidopsis model (Brown et al., 2007, Bauer
313 et al., 2006, Brown et al., 2005). The *irx9* plants have impaired xylan synthesis resulting in a decrease
314 of xylan by more than 50% compared to WT (Brown et al., 2007). In *irx10* plants the reduction in
315 xylan content is smaller and does not exceed 20% (Brown et al., 2009). Macrofibrils are clearly
316 observed in *irx9* and *irx10* Arabidopsis (Figure 5a and 5b). However, the median macrofibril diameter
317 between WT and *irx9* cell wall fibres showed a ~30% reduction in the xylan synthesis mutant (Figure
318 5g). The median macrofibril diameter of *irx10* plants was ~10% smaller than that of WT Arabidopsis
319 (Figure 5g). Although there was a wide variation in macrofibril diameter within each genotype, the
320 difference between the WT macrofibril diameter and the one quantified for the two mutants is
321 statistically significant, suggesting that xylan is incorporated along with cellulose to generate the
322 normal macrofibril size. To investigate the role of xylan-cellulose interaction in the macrofibril
323 formation we assessed the macrofibril size in the *eskl* Arabidopsis mutant (Figure 5c). Mutation in the
324 *ESK1* gene results in reduction of xylan acetylation, but not in a decrease in xylan quantity (Xiong et
325 al., 2013), which leads to changes in xylan [Me]GlcA patterning and loss of interaction between xylan
326 and the hydrophilic surface of the cellulose microfibril (Grantham et al., 2017). In line with the results
327 observed for *irx9* and *irx10* plants the loss of xylan-cellulose interaction caused a reduction in the
328 macrofibril diameter (Figure 5g).

329 Previous work in softwood suggested that lignin (Donaldson, 2007) and galactoglucomannan (GGM)
330 (Terashima et al., 2009) may be involved in macrofibril formation. To investigate the role of these two
331 cell wall components in the maintenance of macrofibril structure we performed imaging of *4cl1* (Figure
332 5d and S11), *lac4* (Figure 5e and S12) and *csla2/3/9* (Figure 5f and S13) mutant Arabidopsis cell walls.
333 Both 4CL1 and LAC4 are involved in lignin biosynthesis and plants mutated in genes encoding these
334 enzymes have a 30% and 15% reduction in lignin content respectively (Li et al., 2015, Berthet et al.,
335 2011). The median macrofibril diameter for both *4cl1* and *lac4* was significantly smaller than that
336 calculated for WT (Figure 5g). Importantly, the extent of the reduction in macrofibril diameter was in
337 line with the decrease in the lignin content observed for the two mutants, with *4cl1* macrofibrils being
338 ~15% smaller than the WT ones and *lac4* macrofibrils having ~7% reduction in the median diameter.
339 Proteins from the CSLA family are involved in the biosynthesis of a hemicellulose
340 galactoglucomannan and mutations in *csla2/3/9* leads to nearly complete loss of stem GGM in the
341 Arabidopsis model (Goubet et al., 2009). Our quantitative analysis indicates that the diameter of
342 macrofibrils of *csla2/3/9* Arabidopsis was not significantly different to that of the WT plants (Figure
343 5g).

344

345 **5 Discussion**

346 The native nanoscale architecture of woody plant secondary cell walls remains poorly understood due
347 to the challenges of keeping the sample hydrated, which is incompatible with some types of techniques.
348 Studies that analyse dehydrated and fixed plant cell wall samples with FE-SEM (Donaldson, 2007),
349 together with other work which includes SANS experiments investigating spruce (Fernandes et al.,
350 2011) and bamboo samples (Thomas et al., 2015), suggest there is a higher order arrangement of
351 cellulose microfibrils in plant secondary cell walls. Our work reports the application of a cryo-SEM
352 based analysis technique which, using exclusively samples that have not been dried, heated or
353 chemically processed, indicates that secondary cell wall cellulose microfibrils are likely to come
354 together to form larger macrofibril structures. Our study strongly suggests that these structures, at least
355 in the model plant species *Arabidopsis thaliana*, are sensitive to changes in xylan and lignin.

356 Previous studies investigated the presence and diameter of macrofibrils in dehydrated softwood
357 samples (Donaldson, 2007). In line with results presented in our work, Donaldson did observe
358 macrofibrils in cell walls of pine tracheids. Moreover, also in agreement with the results presented here
359 (Figure S4), these softwood macrofibrils were larger than those seen in hardwoods. In softwood, in
360 addition to various patterned types of xylan (Busse-Wicher et al., 2016b, Martinez-Abad et al., 2017),
361 most of which are likely to be compatible with binding to the hydrophilic surface of the cellulose fibril,
362 the cell walls contain large quantities of acetylated GGM (Scheller and Ulvskov, 2010) which may
363 contribute to macrofibril width. Indeed, gymnosperm GGM was proposed to interact with the cellulose
364 microfibril in cell walls of Ginkgo (Terashima et al., 2009). Therefore, the significant difference in
365 macrofibril diameter observed between hardwood and softwood samples may be due to the differences
366 in the cell wall composition. Consequently, we hypothesize that in gymnosperms, GGM, along with
367 xylan, may contribute to the macrofibril size in a way similar to what we observed for xylan in
368 *Arabidopsis* macrofibrils. With an average diameter ranging between 20 and 34 nm, the size of pine
369 macrofibrils measured by Donaldson was somewhat smaller than that measured in spruce wood in the
370 current work. However, these observations are not necessarily inconsistent. Donaldson dehydrated the
371 wood samples prior to the SEM imaging. As the spacing between bundled softwood cellulose
372 microfibrils, estimated to be equal to 3 nm by small angle neutron scattering, is sensitive to wood
373 hydration levels (Fernandes et al., 2011), at least part of the difference in the macrofibril diameter
374 might be due to the changes in the water content within the sample analysed with SEM. Interestingly,
375 Donaldson reported that macrofibrils in dried poplar wood, depending on their position in cell wall,
376 have an average diameter ranging from 14 to 18 nm, which is similar to what was measured for both
377 poplar and *Arabidopsis* as a part of our study. This observation suggests that the softwood macrofibril
378 size may be more sensitive to drying than the hardwood one. This in turn suggests that, in addition to
379 compositional disparities, hydration could contribute to the differences in softwood and hardwood
380 macrofibril characteristics. In addition to providing scientific insight, this result highlights that imaging
381 of the cryo-preserved secondary cell walls offers significant advance over the previously used
382 techniques.

383 Interestingly, similar to a previous report (Donaldson, 2007), we observed that macrofibrils in both
384 hardwood and softwood have a range of diameters. The reasons for this variation in size are not clear.
385 It is possible that the number of individual cellulose microfibrils that come together to form the
386 macrofibril structure in both hardwood and softwood is not constant. This may be regulated by
387 coordinated movement of CesA complexes or their density during cell wall synthesis (Li et al., 2016).
388 It was proposed that the macrofibril diameter is proportional to the degree of cell wall lignification
389 (Donaldson, 2007), which may also vary between the structures. This hypothesis is supported by our
390 results which indicate that the cell wall lignin content influences macrofibril diameter in *Arabidopsis*.

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391 Variations may also originate from environmental conditions. For example, it was shown that wood
392 density may vary correlatively with climate change (Bouriaud et al., 2005). Although much of this
393 effect is likely to be due to cell size and wall thickness, it can be hypothesized that change in wood
394 density may also originate from compositional changes that impact macrofibril assembly and
395 ultrastructure. It would therefore be relevant to assess macrofibrils of perennial trees with samples
396 spanning several years of growth. We cannot rule out that the width variance may originate from the
397 technical limitations of resolving the macrofibrils by SEM. It will be interesting to see if the emerging
398 He-ion technologies, with an increase in resolution and less dependence upon metal coating, reduce
399 this variance (Joens et al., 2013). The cryo-SEM techniques developed as part of our study offer a
400 significant advantage over the previous investigation (Donaldson, 2007) which applied a thicker coat
401 of chromium (mostly 12 nm) that yield films with coarser grains than the thinner (3 nm) platinum films
402 used in our work. Thus, taking the results described by Donaldson and our technological improvements
403 into consideration, we believe that the variance in the macrofibril width observed in both studies is
404 likely to reflect natural material variation.

405 The prominence of macrofibril structures in Arabidopsis cell walls is a surprising discovery of this
406 study. Previously published results using AFM analysis indicate the presence of some bundled
407 microfibrils in primary cell walls of Arabidopsis but the extent of this bundling is lower than what was
408 observed in primary cell wall samples from other species (Zhang et al., 2016). AFM is not yet
409 technically feasible for analysis of bundling of hydrated secondary cell walls although recent technical
410 advances allowed visualisation of dried spruce wood at a nanometer resolution (Casdorff et al., 2017).
411 The observation of the macrofibrils by cryo-SEM in Arabidopsis allowed us to determine the
412 contribution of cellulose, xylan, lignin and galactoglucomannan to macrofibril formation, thanks to the
413 availability of secondary cell wall related mutants in this model. Macrofibrils were completely absent
414 in vessel cell walls of *irx1*, *irx3* and *irx5* plants, which lack secondary cell wall cellulose, indicating
415 that proper cellulose biosynthesis is required for formation and assembly of secondary cell walls
416 polymers into macrofibrils. In addition, we observed that vessel macrofibril diameter is significantly
417 decreased in *irx9*, *irx10* and *esk1* plants, suggesting that xylan may also participate in the correct
418 assembly of such structures. While in *irx9* and *irx10* reduction in macrofibril diameter may be
419 associated with decrease in the xylan content the ~25% reduction in the median macrofibril diameter
420 observed for *esk1* Arabidopsis is harder to explain. Hardwood xylan is proposed to interact with the
421 hydrophilic surface of the cellulose microfibril as a two-fold screw (Simmons et al., 2016, Busse-
422 Wicher et al., 2016a), and this interaction is facilitated by the even pattern of the [Me]GlcA and acetyl
423 branches on the xylan backbone which is lost in *esk1* plants (Grantham et al., 2017). Therefore, the
424 decrease in macrofibril diameter observed in *esk1* Arabidopsis indicates that xylan-cellulose interaction
425 may have a role in spacing or proper coalescence of microfibrils to form the elementary macrofibril. It
426 is unclear why the macrofibril diameter is reduced in *esk1*, but perhaps fewer elementary fibrils are
427 incorporated into each macrofibril when xylan is not interacting with the hydrophilic surface of the
428 cellulose fibril. This may be different to the effect observed in flax where the absence of xylan may
429 lead to aggregation of glucan chains into larger fibres (Thomas et al., 2013). Such difference may be
430 associated with variations in the stoichiometry of the cellulose synthase complex which were recently
431 reported for angiosperms (Zhang et al., 2018).

432 In addition to implicating xylan in the process of macrofibril formation our results indicate that lignin
433 may contribute to assembly of the structures. As such, our results use genetic assignment to extend
434 previous work which has correlated macrofibril diameter with the degree of wall lignification
435 (Donaldson, 2007). Interestingly, we observed that the macrofibril diameter does not correlate with the
436 cell wall GGM content. This may be associated with low abundance of GGM in angiosperms where
437 the polysaccharide accounts for only up to 5% of the cell wall material (Scheller and Ulvskov, 2010).

438 Alternatively, this result may indicate that in Arabidopsis GGM might be not involved in microfibril
439 formation. GGM may play a more significant role in the microfibril assembly in gymnosperms where
440 it accounts for up to 30% of the cell wall material. Importantly, all our conclusions are based on the
441 analysis of native, hydrated, cell wall samples. The assignment of cell wall microfibril composition,
442 in their native state, would be impossible using techniques such as immunogold due to the pre-
443 treatment steps needed before the antibody labelling.

444 In conclusion, our analysis indicates that Arabidopsis vessel cell walls contain fibrous structures
445 composed of cellulose and likely contain xylan and lignin. These structures are present in both
446 hardwood and softwood and have a diameter larger than a single cellulose microfibril. Therefore, these
447 structures can be described as cell wall microfibrils. The reduction in microfibril diameter observed
448 in *eskl* Arabidopsis suggests that the interaction between xylan and the hydrophilic surface of the
449 cellulose microfibril may be involved in the assembly of these structures. Therefore, this xylan-
450 cellulose interaction may be important for the maintenance of plant cell wall ultrastructure and
451 mechanical properties (Simmons et al., 2016). The techniques developed here and the discovery of the
452 ubiquitous presence of microfibrils in hardwood and softwood in their native state will contribute to a
453 better understanding of cell wall assembly processes. Furthermore, the ability to resolve microfibrils
454 in Arabidopsis, along with the availability of genetic resources in this model, will offer the community
455 a valuable tool to further study the complex deposition of secondary cell walls polymers and their role
456 in defining the cell wall ultrastructure. The assembly of cell wall microfibrils is likely to influence the
457 properties of wood, such as density, which may vary due to different stimuli such as tree fertilisation
458 (Makinen et al., 2002) or environmental changes (Bouriaud et al., 2005). Therefore, we expect that the
459 methodology described here will enable to correlate the native nanoscale features of the cell walls,
460 such as the microfibril diameter, or a specific microfibril patterning within the cell wall, with wood
461 properties. Consequently, our approach may be useful to assess this aspect of wood quality at a new
462 level and could benefit numerous industries ranging from building construction, paper manufacturing
463 and biofuel production to generation of novel biomaterials such as nanocrystalline cellulose.

464

465 **6 Conflict of Interest**

466 The authors declare that the research was conducted in the absence of any commercial or financial
467 relationships that could be construed as a potential conflict of interest.

468 **7 Author Contributions**

469 JJL designed the study, conducted the experiments, analysed the data and wrote the paper. MB
470 performed poplar imaging experiments, analysed the data and wrote the paper. OMT analysed the data
471 and wrote the paper. YH contributed to data analysis and manuscript preparation, RW designed the
472 study, conducted experiments, analysed the data and wrote the paper. PD designed the study and
473 contributed to data analysis and manuscript preparation.

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496 **10 List of abbreviations**

497 1D – one dimensional

498 AFM – atomic force microscopy

499 CesA – Cellulose synthase

500 cryo-SEM – low temperature scanning electron microscopy

501 FE-SEM – field emission scanning electron microscopy

502 FT-IR - Fourier-transform infrared spectroscopy

503 GGM – galactoglucomannan

504 He-ion – Helium ion

505 IRX – irregular xylem

506 [Me]GlcA – methylated and unmethylated form of glucuronic acid

507 NMR – nuclear magnetic resonance

508 SANS – small angle neutron scattering

509 TEM – transmission electron microscopy

510 WAXS – wide angle x-ray scattering

511 **11 Data Availability Statement**

512 All quantitative datasets generated and analysed for this study are presented on graphs included in the
513 manuscript and the supplementary files.

514 12 References

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732

733 **13 Short legends for supporting material figures:**

734 Figure S1: Overview of the cryo-SEM procedure.

735 Figure S2: Cryo-SEM analysis of Ginkgo cell walls.

736 Figure S3: Comparison of macrofibril diameter in Arabidopsis, poplar, spruce and Ginkgo.

737 Figure S4: Imaging of macrofibrils in field grown poplar.

738 Figure S5: Analysis of native Arabidopsis samples without the cryo-preservation protocol.

739 Figure S6: Cryo-SEM analysis of vessel collapse and primary cell wall cellulose in *irx3* Arabidopsis
 740 plants.

741 Figure S7: Analysis of *irx1* and *irx5* cell walls.

742 Figure S8: Further images of *irx9* plants.

Cell wall macrofibrils are composed of cellulose, xylan and lignin

743 Figure S9: Further images of *irx10* plants.

744 Figure S10: Further images of *esk1* plants.

745 Figure S11: Further images of *4c11* plants.

746 Figure S12: Further images of *lac4* plants.

747 Figure S13: Further images of *csla2/3/9* plants.

748

749 **14 Main text figure legends**

750 **Figure 1. Measurement of cell wall macrofibrils.** (a) Example of macrofibrils which would be
751 considered for measurement. Only macrofibrils that were resolvable from their neighbours were
752 analysed. The diameter was measured at a site along the length of the macrofibril and not at the
753 fractured ends. Measurement (b) was carried out by placing one line in parallel to the macrofibril and
754 measuring the length of a line perpendicular to it and spanning the width of the structure to be analysed.

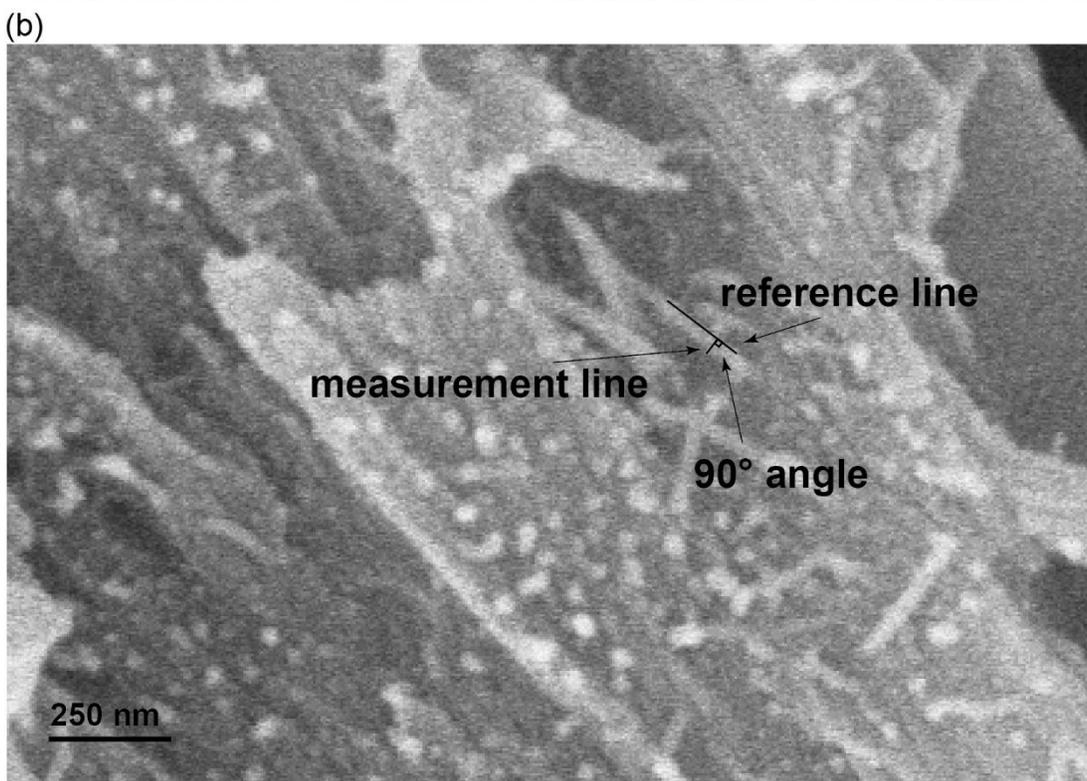
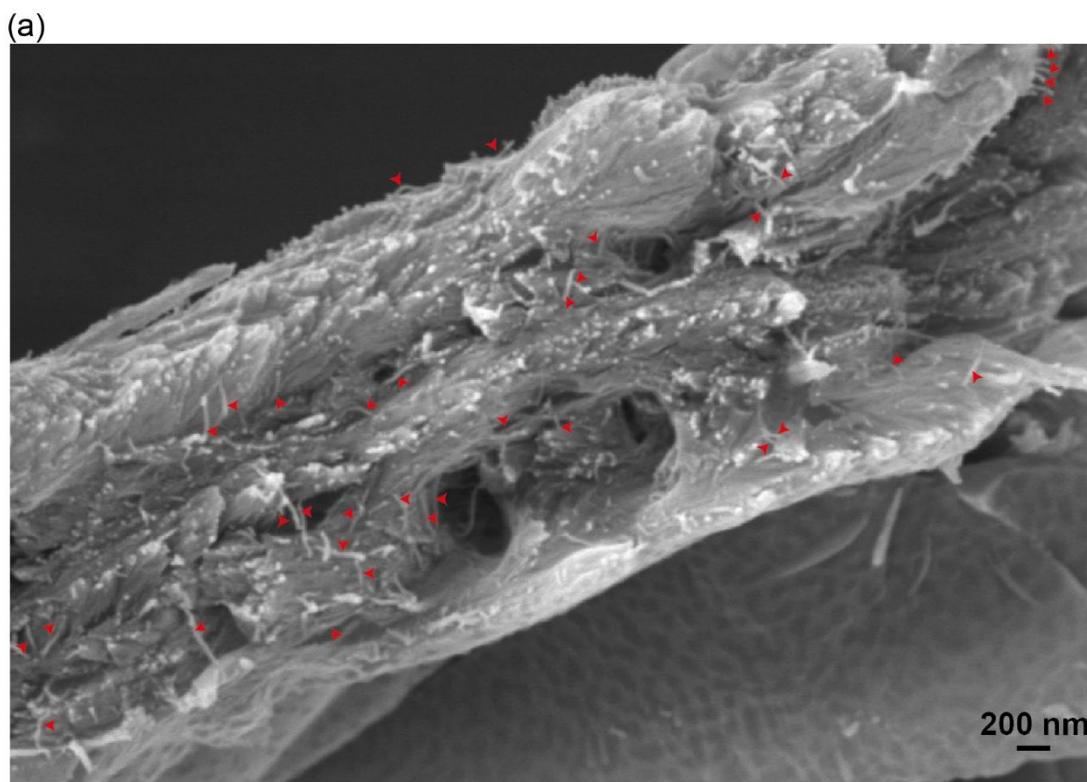
755 **Figure 2. cryo-SEM analysis of spruce stem sections.** (a) to (f) Representative images of stem
756 sections of one-year-old spruce branch at different magnifications. Red arrows indicate tracheids (b),
757 macrofibril bundles (c and d) and individual macrofibrils (e and f). Scale bars are provided for each
758 image.

759 **Figure 3. cryo-SEM analysis of poplar stem sections** (a) to (e) Representative images of stem
760 sections of *in vitro* grown poplar trees at different magnifications. Red arrows show vessels (b) and
761 macrofibrils (c and e). Yellow arrows indicate fibre cells (b). Higher magnification images (c, d and
762 e) are presented for vessels. Scale bars are provided for each image. (f) Diameter of spruce tracheid
763 cell wall fibrils compared to these observed in poplar vessel cell walls. For each bar 150 individual
764 fibrils were measured. Boxplots mark the median and show between 25th and 75th percentile of the
765 data. *** denotes $p \leq 0.00001$ in Student's t-test.

766 **Figure 4. Analysis of Arabidopsis stem sections and fibrous cellulose.** (a) to (c) Imaging of WT
767 vessels at increasing magnification (d) Imaging of fibrous cellulose standard from cotton linters shows
768 cell wall fibrils with an appearance similar to structures seen *in planta*. (e) Imaging of individual
769 vessels in WT plants. (f) Imaging of individual vessels in *irx3* plants. (g) and (h) Macrofibrils are
770 detectable in WT Arabidopsis and are absent in *irx3* secondary cell walls. Red arrows indicate the
771 macrofibril structures throughout the figure. Scale bars are provided for each image.

772 **Figure 5. Analysis of macrofibrils in mutant Arabidopsis plants.** Representative image of (a) *irx9*,
773 (b) *irx10*, (c) *esk1*, (d) *4c11*, (e) *lac4* and (f) *csla2/3/9* Arabidopsis macrofibrils. Scale bar corresponds
774 to 200 nm on each image. Red arrows show macrofibrils (g) Quantification of macrofibril diameter in
775 WT and mutant Arabidopsis plants. N = 150. Boxplots mark a median and show between 25th and 75th
776 percentile of the data. *** denotes $p \leq 0.00001$, ** denotes $p \leq 0.0001$, * denotes $p \leq 0.05$ in Tukey
777 test following ANOVA when compared to WT, ns indicates lack of statistically significant difference.
778 Additional images of each genotype are shown in figures S8 - S13.

779



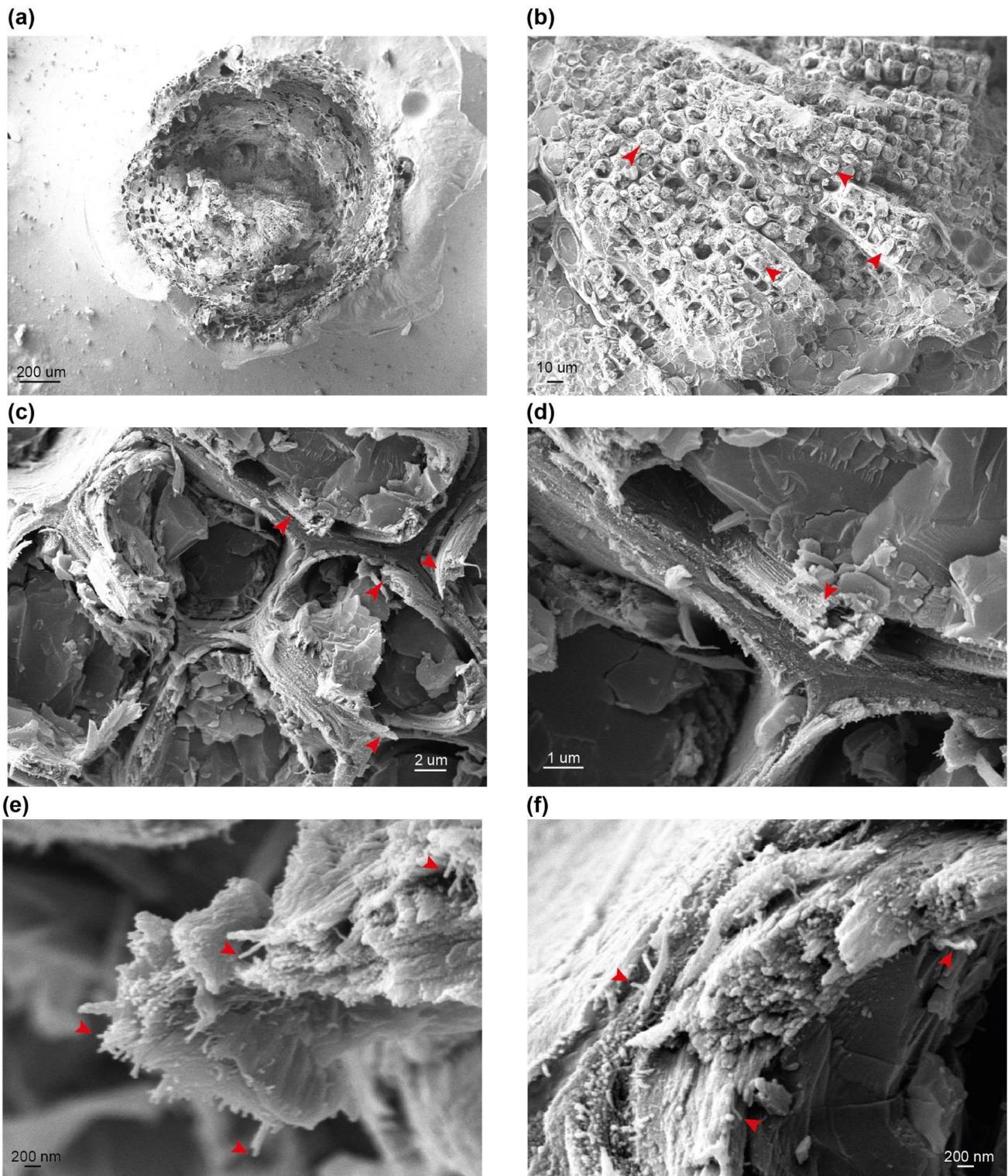
780

781 Figure 1

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Cell wall macrofibrils are composed of cellulose, xylan and lignin



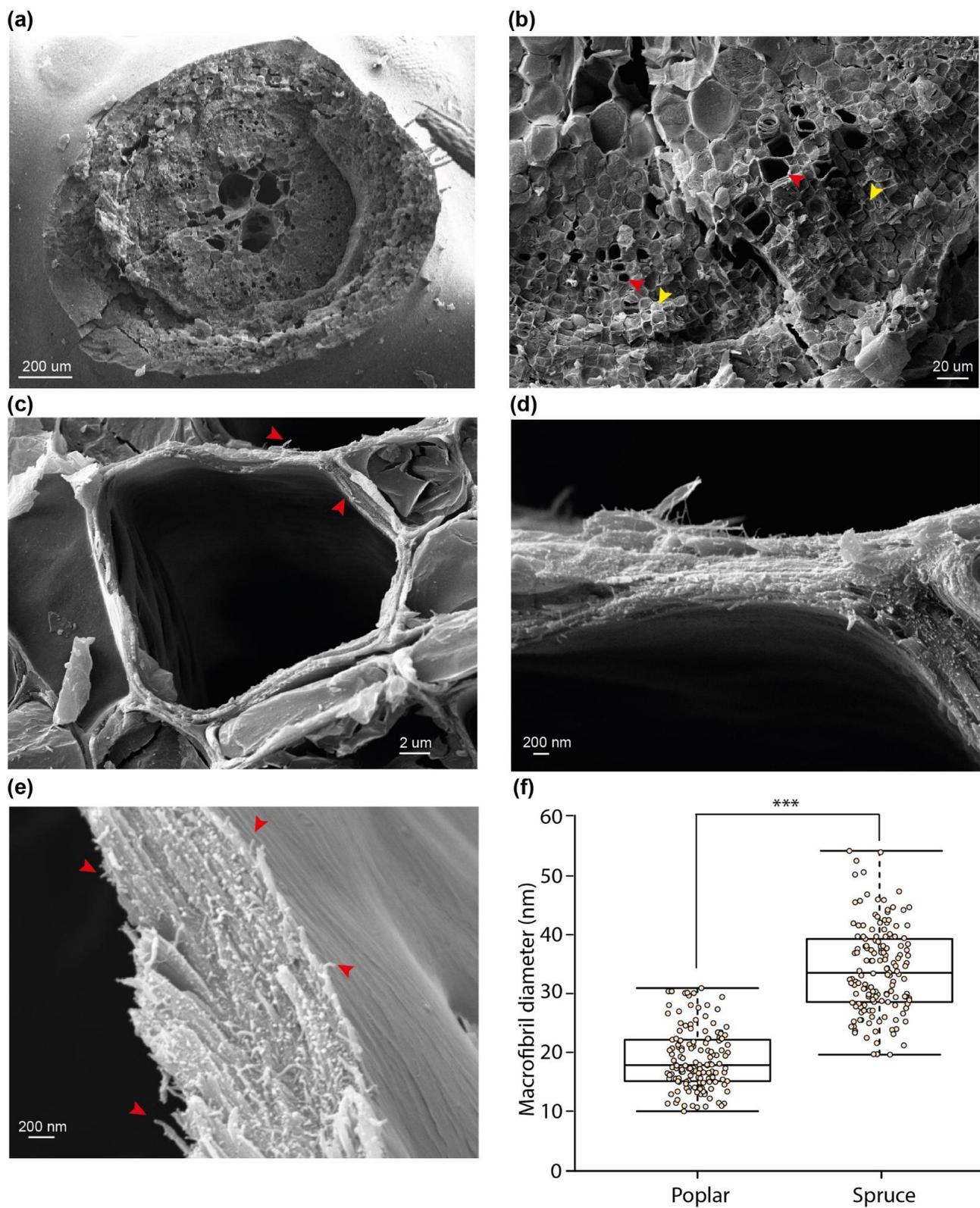
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785 Figure 2

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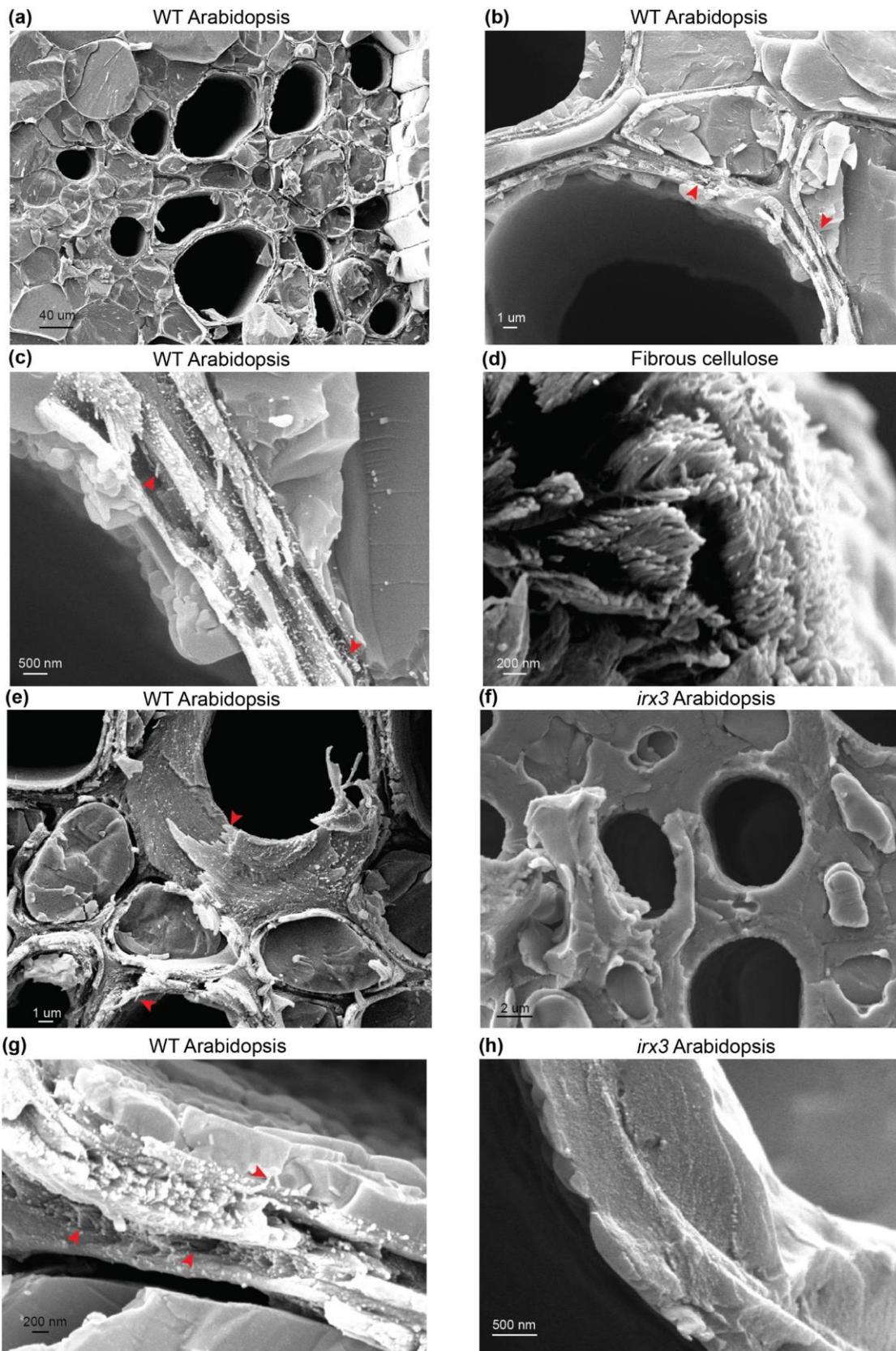
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790 Figure 3

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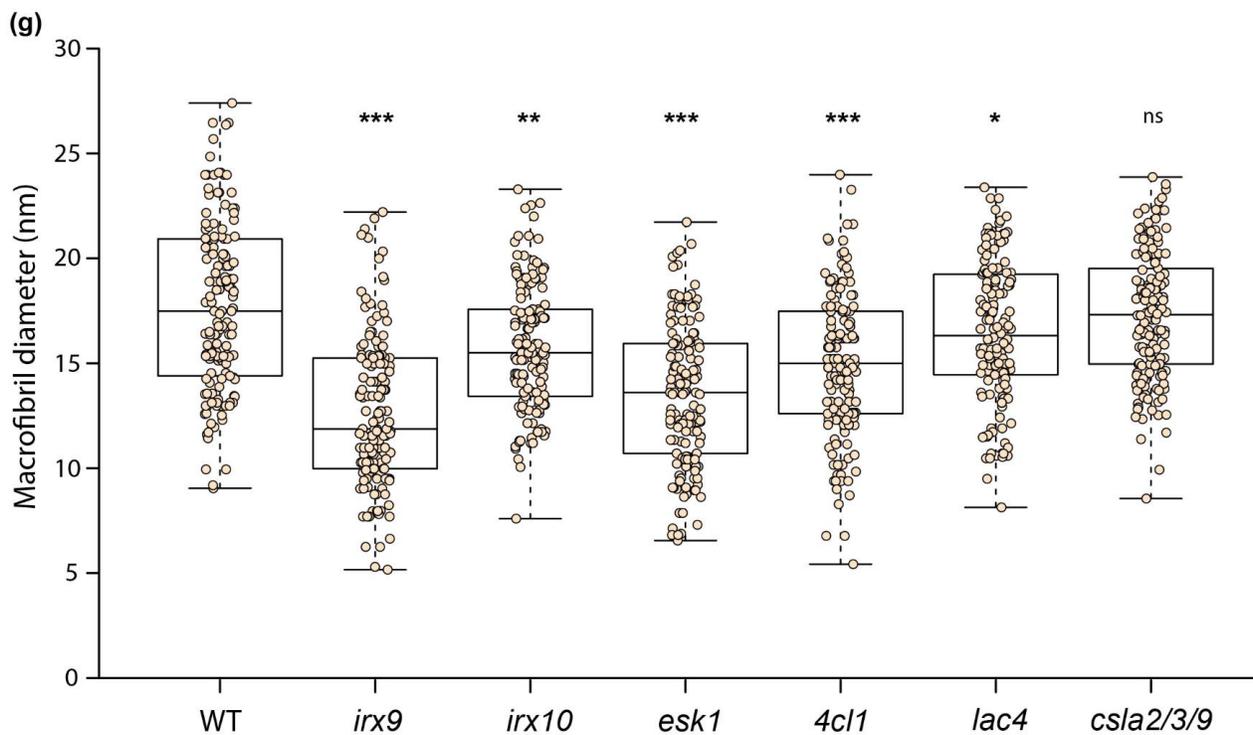
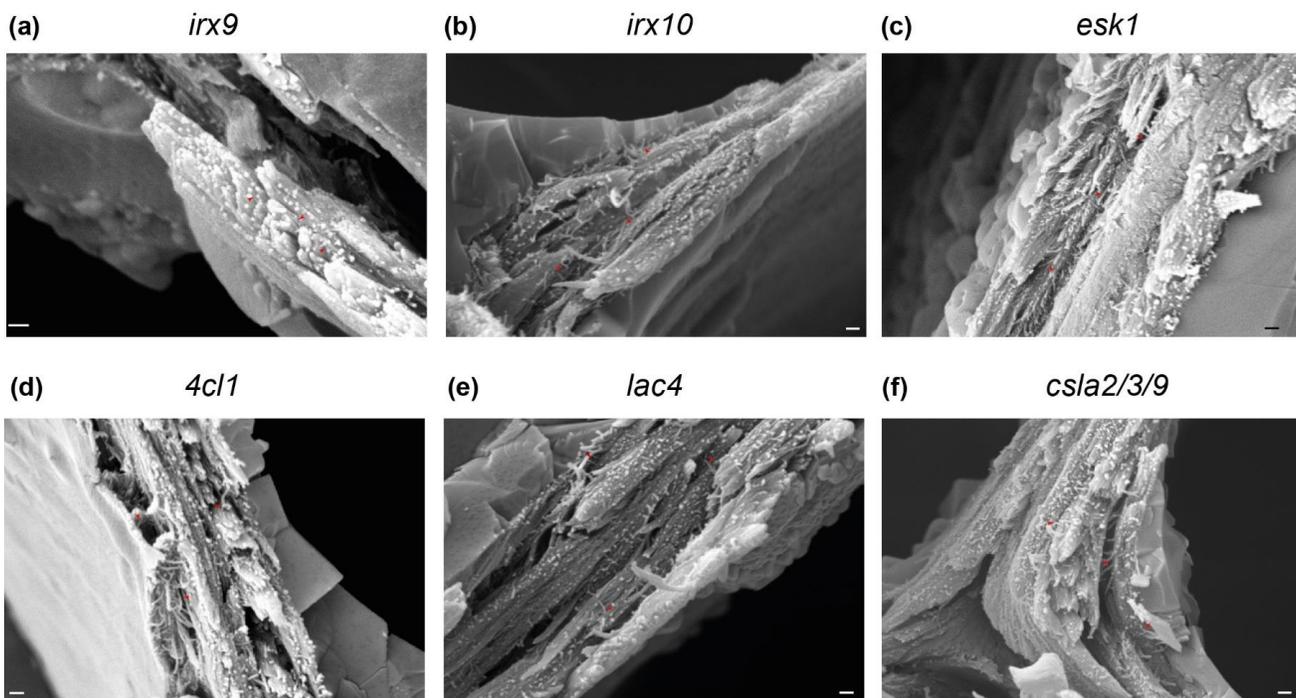
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Cell wall macrofibrils are composed of cellulose, xylan and lignin



793

794 Figure 4



Median macrofibril diameter (nm)

WT	<i>irx9</i>	<i>irx10</i>	<i>esk1</i>	<i>4cl1</i>	<i>lac4</i>	<i>csla2/3/9</i>
17.5	11.9	15.5	13.6	15.0	16.3	17.3

795

796

797 Figure 5

798

Cell wall microfibrils are composed of cellulose, xylan and lignin

799