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Stem xylem traits and wood formation affect sex-specific responses to drought and rewatering in Populus cathayana

Liu, Miao

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1 Tree Physiology

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3	Stem xylem traits and wood formation affect sex-specific responses to drought
4	and re-watering in Populus cathayana
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6	Miao Liu ¹ , Yang Zhao ¹ , Yuting Wang ¹ , Helena Korpelainen ² , Chunyang Li ^{1,*}
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8	
9	¹ College of Life and Environmental Sciences, Hangzhou Normal University, Hangzhou
10	311121, China
11	² Department of Agricultural Sciences, Viikki Plant Science Centre, University of
12	Helsinki, P.O. Box 27, FI-00014, Finland
13	
14	
15	* Corresponding author: Chunyang Li, E-mail address: <u>licy@hznu.edu.cn</u>
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18	Head title: Sexual differences in drought and re-watering responses
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Abstract The increased frequency and intensity of drought pose great threats to the 23 survival of trees, especially in dioecious tree species, which usually display sex-related 24 25 differences in mortality and biased sex ratios. The present study explored the roles of sex-specific differences in xylem anatomy and function and carbon metabolism in 26 drought resistance and drought recovery in dioecious Populus cathayana Rehder. We 27 showed that the sex-specific drought resistance and subsequent recovery were closely 28 linked to the xylem anatomy and carbon metabolism. Females had a greater xylem 29 vessel area per vessel, biomass and hydraulic efficiency (a higher theoretical hydraulic 30 31 conductivity) under well-watered conditions. In contrast, males had a lower xylem lumen area, but greater vessel numbers, and a higher cell wall thickness and leaf 32 photosynthesis, suggesting a conservative water-use strategy (a lower theoretical 33 34 hydraulic conductivity) and drought resistance under drought stress. The recovery of photosynthetic ability and leaf water potential after drought in males was largely 35 dependent on the recovery of xylem function and the regulation of the xylem 36 37 carbohydrate metabolism. Additionally, numbers of up-regulated genes related to xylem cell wall biogenesis and cell turgor maintenance were greater in males relative 38 to females under drought stress and subsequent re-watering, which facilitated drought 39 resistance and xylem function restoration in males. These results suggested that sex-40 specific drought resistance and restoration were related to xylem anatomy and function, 41 carbohydrate metabolism and cell turgor maintenance. 42

Key words: dioecy, drought, re-watering, xylem anatomy, hydraulic efficiency,
hydraulic safety

Increasingly frequent and intensive drought events have been linked with plant mortality. For instance, drought impacts growth and distribution of tree populations not only during the drought period, but there may be lasting effects afterwards even for years (Barigah et al., 2018; Kannenberg and Phillips, 2020). Therefore, it is critical to elucidate the physiological and molecular mechanism causing the drought-induced plant damage and recovery after drought.

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As previously discovered, drought-induced tree mortality is caused by hydraulic failure 54 and carbon starvation (Rodriguez-Zaccaro and Groover, 2019). Leaf photosynthesis is 55 56 widely considered as the major carbon source of plants in response to drought. Maintaining the continued water supply for leaf photosynthesis is critical for plant 57 survival and growth in dry environment (Sapes et al., 2019). In woody plants, leaf 58 59 photosynthesis is closely associated with the plant's hydraulic system, which can supply water to leaves (Martorell et al., 2014). In woody plants, the drought-induced hydraulic 60 conductivity of xylem coincides with stomatal closure and the reduced leaf 61 photosynthetic capacity (Martorell et al., 2014). The recovery of leaf photosynthesis 62 largely depends on the restoration of xylem structure and functions (Martorell et al., 63 2014). Moreover, the non-structural carbohydrate (NSC) pools in the wood limit the 64 increasing xylem tension and maintain the hydraulic integrity of xylem (De 65 Baerdemaeker et al., 2017; De Roo et al., 2020). The ability and extent to refill 66

embolized vessels after re-watering plants are also associated with NSC pools (Brunetti
et al., 2020; Kannenberg and Phillips, 2020). However, the regulation of the wood
xylem structure and NSC dynamics in response to drought and subsequent re-watering
is unclear, especially in dioecious plant species.

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72 Drought increases xylem tension in woody plants. Stomatal closure reduces water loss and xylem tension, which cannot prevent xylem embolism during extended drought and 73 consequent plant mortality (Kannenberg and Phillips, 2020). Variation in the 74 75 morphology and arrangement of xylem elements shares a basic structural principle for the reduction of xylem embolism and adaptation to drought (Fonti et al., 2010; Torres-76 Ruiz et al., 2017). It has been suggested that the anatomical traits of xylem could 77 78 successfully predict hydraulic function (Hacke et al., 2015). According to Hagen-Poiseuillle's law, the rate of water flow is proportional to the fourth power of the 79 tracheid lumen diameter (Hacke and Sperry, 2001). The hydraulic efficiency in plants 80 81 with higher lumen diameter is more sensitive to xylem embolism under drought (Olano et al., 2017). In contrast, the increased wall thickness relative to the lumen area tends 82 to indicate a more conservative strategy, with less conductivity and correspondingly 83 lower vulnerability to embolism, especially under drought stress (Rungwattana and 84 85 Hietz, 2018).

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The anatomical traits of xylem are suggested to be developmentally plastic, which reflects great morphological diversity across plant lineages and the adaptation of

hydraulic traits to different environmental conditions (Martorell et al., 2014; Hultine et al., 2016). Moreover, plants can utilize NSC for osmotic regulation to maintain cell turgor and/or refill embolized xylem vessels under drought and/or re-watering after drought (Secchi and Zwieniecki, 2012). Despite the established association between the physiological plasticity of xylem and water flow, very little is known about the molecular mechanism underlying xylem traits under drought and re-watering, especially in dioecious plants.

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Wood growth is considered as an ideal system for studying xylem anatomy and function 97 and for providing implications for the carbon cycling under drought and re-watering. 98 Wood stores immense amounts of carbon with a slow carbon turnover, and the growth 99 100 is closely correlated with the net primary productivity (Anderegg et al., 2015). The regulation of xylem traits and xylem NSC under drought and recoveryafter drought 101 varies among plant species and genotypes, as well as depending on the severity and 102 duration of drought (Olano et al., 2017; Falchi et al., 2020). Tree species exhibit major 103 interspecific and intraspecific variation in hydraulic conductivity along an 104 environmental gradient, especially under drought (Burton et al., 2017; Islam et al., 105 2018). 106

Variation in hydraulic conductivity may show opposite patterns in males and females
of dioecious species if variation is tightly associated with sex (Olano et al., 2017).
Dioecious plants exhibit sexual dimorphism in response to abiotic stress (Liu et al.,
2020; Liu et al., 2021a). Female plants usually have a higher reproductive effort than

males, since females produce flowers, seeds and fruits, and they require contrasting
water-use strategies compared to males (Hultine et al., 2016; Liu et al., 2021b). Indeed,
the female trunk exhibits such anatomical traits of xylem that relate to a larger hydraulic
efficiency, while male anatomy shows a more conservative strategy, especially under
drought conditions, for instance, in the dioecious conifer *Juniperus thurifera* (Olano et
al., 2017). However, much is still unknown.

In this study, we used *Populus cathayana* females and males as a model species to elucidate the sex-specific tolerance and responses to drought and subsequent rewatering based on the structure and function of wood xylem. We aimed to answer three main questions: (1) do wood xylem traits differ between sexes under drought and re-watering; (2) how is the trade-off between traits related to hydraulic efficiency in males and females; (3) how is the xylem structure of males and females regulated by drought and re-watering after drought stress at the molecular level.

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125

- 126 Materials and Methods
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128	Plant material and experimental design	ļ

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P. cathayana cuttings were gathered from different female and male trees in riparian
and valley flat habitats of the Qinghai Province, China (30°67′N, 104°06′E), as
explained in detail by Liu et al. (2020). The cuttings were grown in a glasshouse with

133	natural light. After growing for 4 weeks, 32 healthy and uniform seedlings (about 30
134	cm) were transplanted into 10-l plastic pots with 10 kg of soil. The soil contained 1.75
135	g kg ⁻¹ total N, 122.56 mg kg ⁻¹ available P, 476.46 mg kg ⁻¹ available K, 106.33 mg kg ⁻¹
136	$NO_3^{-}-N$, 52.69 mg kg ⁻¹ $NH_4^{+}-N$, and 33.32 g kg ⁻¹ soil organic matter. After 6 weeks, 16
137	seedlings of each sex were completely randomized and divided into well-watered
138	condition, CK; drought, D; drought + re-watering 7 d, DW7; drought + re-watering 14
139	d, DW14. Firstly, all seedlings of each sex were randomly divided into two groups (CK,
140	D) and cultivated for 6 weeks. Subsequently, 8 plants (CK, D) of each sex were
141	harvested. The remaining plants were re-watered to 100% field capacity for 7 d and 14
142	d. In this study, we mainly explored sexual differences in responses to drought and re-
143	watering. For CK, the pots were weighed each day to maintain 100% field capacity. For
144	D, the pots were weighed each day to maintain 30% field capacity.

146 *Gas exchange and leaf water potential measurements*

Gas exchange parameters were measured on fully developed mature leaves between 09:00 and 11:00, and 14:00 and 16:30. Measurements were performed using a portable photosynthesis measuring system (Ll-6400; Li-Cor Inc., Lincoln, NE, USA) equipped with an integrated leaf chamber fluorometer (LI-6400-40). The conditions for measuring leaf gas exchange parameters were as follows: relative humidity of 75%, airflow rate of 500 μ mol s⁻¹, CO₂ concentration in the cuvette (*C*_a) of 400 μ mol mol⁻¹ and the saturating photon quantum flux density (PPFD) of 1200 μ mol m⁻² s⁻¹ (with 90%

155	red light and 10% blue light). After measuring leaf gas exchange, sampled leaves were
156	cut for the analysis of leaf water potential early in the morning in a pressure chamber
157	(PMS Instruments, Albany, OR, USA).

159 Microscopic observations and anatomical measurements

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After the determination of photosynthetic parameters, stems were cut into segments (1– 161 2 mm²) and immediately immersed in FAA solutions. Afterwards, the samples were 162 rinsed three times with ultrapure water, and dehydrated with a series of ethanol 163 gradients. Samples were embedded in glycol methacrylate. Semi-thin cross-sections 164 were stained with safranin-fast green and viewed under a light microscope (Nikon 165 166 Corporation, Kyoto, Japan) with 20× magnification. The thickness, size and density of leaf palisade and spongy tissues, and the sum of spongy and palisade tissue cell area 167 per transverse section were measured using light micrographs. 168

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170 *Analysis of wood composition by fourier transform infrared spectroscopy (FTIR)*

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The developing xylem was obtained according to Teichmann et al. (2008). Briefly, stem segments were sampled at about two-thirds above the stem base. Only developing xylem was harvested (mature wood increases toward the tree base). Developing xylem was sampled by scraping the surface of the xylem with a razor after the tree bark was removed. Afterwards, the scraped developing xylem was immediately put into liquid nitrogen. Subsequently, the samples were divided into two parts. One part of the
samples was dried with a vacuum freeze dryer for 100 h. The freeze-dried sample
powder was pressed against the diamond crystal of an attenuated total reflectance
device and the infra-red spectra were determined with a FTIR spectrometer Nicolet iS5.
The scanning range was 400-4000 cm⁻¹ wavenumber. The remaining samples were kept
at -80 °C until the subsequent transcriptome sequencing.

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184 *RNA extraction and sequencing*

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Frozen tissues of developing xylem samples (about 100 mg) were ground with a ball 186 mill in liquid nitrogen. Subsequently, total RNA was extracted with a RNA extraction 187 188 kit (TaKaRa MiniBEST Plant RNA Extraction Kit, TaKaRa, Otsu, Japan), and genomic DNA was digested using DNase I according to the manufacturer's instructions (Takara, 189 Otsu, Japan). RNA yield and purity were assessed with the NanoDrop 190 191 spectrophotometer 2000 (NanoDrop Technologies Inc.). The concentration of RNA was assessed using the Agilent 2100 Bioanalyzer (Agilent Technologies Inc.). The synthesis 192 of cDNA was performed with random primers and reverse transcriptase. Double-strand 193 cDNA was synthesized using a SuperScript double-stranded cDNA synthesis kit 194 (Invitrogen, CA) according to the manufacturer's protocols, and the sequencing 195 libraries were synthesized with the TruSeqTM RNA sample preparation kit (Illumina, 196 USA). Subsequently, the cDNA products were purified and amplified with Phusion 197 DNA polymerase (NEB). The libraries were sequenced with the Illumina HiSeq 4000 198

Systems, each with three biological replicates. Sequencing reads were purified by 199 removing adaptors, regions with more than 5% unknown N bases and low-quality reads 200 201 using the SOAPnuke software (BGI, Shenzhen, China). Clean reads were aligned with of Р. the reference trichocarpa 202 genome (http://plants.ensembl.org/Populus trichocarpa/Info/Index). The mapping reads were 203 constructed with StringTie software (v1.04). 204

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206 *Data processing and analysis*

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We identified and calculated differentially expressed genes with transcripts per kilobase 208 of the exon model per million mapped reads (TPM). The abundances of genes were 209 210 quantified with RSEM (http://deweylab.biostat.wisc.edu/rsem/). The coding sequence of P. trichocarpa was blasted against the closest Arabidopsis homolog (AGI 211 identification) with PoplarGene (http://bioinformatics.caf.ac.cn/PoplarGene/gene) and 212 213 annotated using the Arabidopsis Information Resource genome (https://www.arabidopsis.org/). The analysis of differential expression was performed 214 with eight groups (FCK, FD, FDW7, FDW14, MCK, MD, MDW7, MDW14). The 215 thresholds (Padj-value ≤ 0.05 and $|\log 2$ fold-change| ≥ 1) were set to determine 216 significant differences in gene expression. The sequencing data were submitted to 217 NCBI (BioProject accession number: SUB9706978). 218

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220 *Quantitative PCR analyses*

222	About 100 mg frozen tissues of developing xylem samples were used to extract total
223	RNA with a RNA extraction kit (TaKaRa MiniBEST Plant RNA Extraction Kit,
224	TaKaRa, Otsu, Japan). RNA yield and purity were assessed with the NanoDrop
225	spectrophotometer 2000 (NanoDrop Technologies Inc.). The first cDNA strand was
226	synthesized using PrimeScript reverse transcription (RT) reagent kits (Takara)
227	following the manufacturer's protocols. Quantitative reverse transcription polymerase
228	chain reactions (qRT-PCR) were conducted in a 25 μl reaction volume with pairs of
229	gene-specific primers (Table S1) using a SYBR®Premix EX Taq kit (TliRNaseH Plus)
230	(TaKaRa) following the manufacturer's instructions. The relative expression levels
231	were calculated according to $2^{-\Delta\Delta Ct}$ (Liu et al., 2017). TUB4.1 was chosen as the
232	housekeeping gene (He et al., 2013).

Statistical analyses

Differences among means within treatments were revealed using the SPSS software (version 22.0). Before analyses of variance (ANOVAs), the data were checked for normality. Differences between means were analyzed by Duncan's tests at a significance level of P < 0.05

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253	Results
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255	Sexual differences in plant growth and tolerance to drought and re-watering
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257	The dry weight of leaves, stems and roots decreased by 78%, 76% and 60%,
258	respectively, in females, and by 60%, 72% and 50%, respectively, in males under

drought stress relative to controls (Table 1). The dry weight of leaves, stems and roots

in females increased by 20%, 13% and 42%, respectively, after 7 d re-watering (DW7),

and increased by 85%, 21% and 57%, respectively, after 14 d of re-watering after

- drought stress (DW14), when compared to the drought treatment (Table 1). In males,
- the dry weigh of leaves, stems and roots increased by 46%, 21% and 82%, respectively,

at 7 d re-watering, and by 119%, 72% and 204%, respectively, at re-watering 14 d after
drought stress when compared to drought (Table 1).

266

Both the photosynthesis rate and leaf gas conductance decreased significantly in 267 females compared to males under drought conditions (Fig. 1a, b). The leaf 268 photosynthesis and gas conductance recovered in females and males at 7 d and 14 d re-269 watering after drought stress. In males, the leaf photosynthesis completely recovered to 270 the control level (without drought stress) at 7 d re-watering after drought stress (Fig. 271 272 1a). The leaf gas conductance in males recovered to 87% of control at 7 d re-watering and 100% of control at 14 d re-watering after drought stress. In contrast, the leaf 273 photosynthesis of females did not completely recover to the control level at 7 d or even 274 275 at 14 d re-watering after drought stress. The gas conductance of females was lower at 7 d of re-watering than in controls but recovered to the control level after 14 d of re-276 watering (Fig. 1b). Leaf water potential significantly decreased by drought stress in 277 278 both sexes, and the decrease was greater in females than in males (Fig. 1c). Leaf water potential reached 229% of the control level in drought-stressed females and 127% in 279 drought-stressed males after 7 d of re-watering. Leaf water potential was fully 280 recovered to the control level after 14 d of re-watering in drought-stressed females and 281 282 males.

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The proportions of xylem length relative to the radius of wood in the cross-section werehigher in males than in females under all treatments (Fig. 1d). Drought stress did not

affect the proportion of xylem length relative to the radius of wood in the cross-section of females, while drought stress lowered the xylem proportion in males relative to controls. The proportions of xylem length relative to the radius of wood in the crosssection increased in drought-stressed males after 7 d of re-watering, but not in droughtstressed females (Fig. 1d). The proportions of xylem length relative to the radius of wood were higher in both sexes at day 14 of re-watering after drought stress compared to drought stress.

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294 Sexual differences in anatomical traits of wood xylem

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The vessel number of both sexes was higher under drought stress, and at 7 d and 14 d 296 297 of re-watering after drought compared to well-watered controls (Fig. 2a). The decreased vessel numbers in females observed by 7 d and 14 d re-watering after drought were 298 similar. In males, the decrease in the number of vessels induced by re-watering after 299 300 drought was greater at 14 d than at 7 d. The lumen area per vessel was higher in females than in males under all treatments (Fig. 2b). The lumen area per vessel decreased by 301 drought stress, but increased by 7 d and 14 d of re-watering after drought in both sexes 302 when compared to controls. Re-watering after drought increased the lumen area per 303 vessel more at 14 d than that at 7 d in females. In males, the lumen area per vessel 304 increased similarly by 7 d re-watering and 14 d re-watering after drought when 305 compared to drought (Fig. 2b). When compared to drought, 7 d re-watering did not 306 affect the cell wall thickness in either sex, whereas the cell wall thickness decreased at 307

308	14 d re-watering after drought in both sexes (Fig. 2c). The predicted hydraulic
309	conductivity in males decreased by drought stress, but completely recovered to control
310	levels at 7 d and 14 d re-watering after drought (Fig. 2d). In females, the predicted
311	hydraulic conductivity was not affected by drought stress, but it decreased similarly at
312	7 d or 14 d re-watering after drought (Fig. 2d).

- 313
- 314 Sexual differences in wood NSC dynamics
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Overall, the levels of total sugar, sucrose and NSC were higher in male wood compared 316 to females under drought and subsequent re-watering. Drought increased the levels of 317 total sugar, sucrose and NSC in leaves of both sexes, which gradually decreased by re-318 319 watering at 7 d and 14 d after drought in both sexes (Fig. 3a, b, d). Females had higher wood starch levels than males under drought and re-watering at 7 d and 14 d after 320 drought (Fig. 3c). Drought increased wood starch concentrations in both sexes, 321 322 whereaswood starch concentrations decreased similarly at 7 d and 14 d re-watering after drought in females and males (Table 1). 323

325 *Fourier transform infrared spectroscopy spectra of xylem*

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The fourier transform infrared spectroscopy spectra were used to obtain the chemical fingerprint of the molecular wood composition in the newly formed xylem of *P. cathayana* females and males under drought and re-watering. The drought and re-

watering differently affected the wood xylem composition of both sexes (Fig. 4a, b; 330 Table S2). PCA analysis was performed to study the original absorbance data. The PC1 331 332 and PC2 accounted for 48% and 47% of the variation, respectively. Peaks at 1505, 1034, 1596 and 1462 cm⁻¹ were key factors contributing to PC1, whereas 1508, 1157, 1110, 333 1425 and 1375 cm⁻¹ were key factors contributing to PC2. The spectra showed a 334 pronounced difference in major wood compounds, such as lignin, cellulose, lignin and 335 carbohydrate, of the developing xylem between drought and control conditions in both 336 sexes. Additionally, the spectra showed significant differences in the developing xylem 337 338 in both sexes and pronounced changes in the xylem structure induced by drought and re-watering 7 d after drought stress, including hemicellulose (peak 1), pection, cellulose, 339 lignin and carbohydrates, as well as the deformation in cellulose and hemicellulose 340 341 (peak 5). At 14 d re-watering after drought stress, the PCA did not distinguish differences in the xylem structure between females and males. 342

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344 *Transcriptome analysis of developing xylem*

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Transcriptomic changes in the developing xylem were analyzed to study the response mechanism in xylem development to drought and re-watering in both sexes. The sequencing data coincided well with the data obtained by qRT-PCR (Fig. S1). The numbers of differentially expressed genes (DEGs) equaled 5262 (2632 down-regulated and 2630 up-regulated) between FW (female, well-watered) and MW (male, wellwatered), 9210 (4145 down-regulated and 5065 up-regulated) between FD (female, drought) and MD (male, drought), 6339 (3210 down-regulated and 3129 up-regulated)
between FDR7 and MDR7, and 5999 (2704 down-regulated and 3295 up-regulated)
between FDR14 and MDR1 (Fig. 5a, c). Specifically, there were 3283 common DEGs
between "FW versus MW" and "FD versus MD", 3509 common DEGs between "FD
versus MD" and "FDR7 versus MDR7", and 3283 common DEGs between "FD versus
MD" and "FDR14 versus MDR14" (Fig. 5a, c).

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In the total analysis, the responses of FW versus MW were not significant for up-359 regulated and down-regulated genes (Fig. 5b). Significantly up-regulated and down-360 regulated genes existed between "FD versus MD". Furthermore, up-regulated genes 361 were found between "FD versus MD", mainly those related to metabolism, and cell 362 363 wall-related and cell cycle processes (Fig. 5c, d). Up-regulated genes between "FD versus MD" were mainly involved in the cell wall biosynthesis, cell wall polysaccharide 364 biosynthesis and metabolism, xylan and lignin biosynthesis and metabolism, and 365 carbohydrate metabolism (Fig. 5c, d). In contrast, down-regulated genes enriched 366 between "FD versus MD" were mainly involved in responses to stress, transport, 367 metabolism and signaling processes (Fig. S2). These down-regulated genes between 368 "FD versus MD" were mainly involved in abiotic stress responses, such as heat, osmotic 369 and salt stresses, and protein modification processes (Fig. S2). Up-regulated genes 370 between "FDW7 versus MDW7" were enriched in cell cycle, cellular organization, 371 signaling and transport, such as mitotic cell cycle processes and regulation, regulation 372 of microtubule cytoskeleton organization, organic acid transport and protein 373

374	phosphorylation. Down-regulated genes between "FDW7 versus MDW7" were
375	enriched in development, catabolism, metabolism, proteostasis and stress responses,
376	such as phloem development, aminoglycan catabolic process, the regulation of
377	peptidase activity and responses to fatty acid (Fig. 5d). Up-regulated genes between
378	"FDW14 versus MDW14" were mainly involved in organic acid transport, protein
379	modification and regulation of the cellular biosynthetic process. Down-regulated genes
380	between "FDW14 versus MDW14" were mainly enriched in protein phosphorylation,
381	defense, sugar metabolism, regulation of mitotic cell cycle and microtubule-based
382	movement (Fig. S2).
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387	Discussion
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389	Sex-specific anatomical traits of xylem in response to drought
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391	Populus females and males exhibit sex-specific differences in responses to abiotic stress,
392	probably associated with sex-specific demands of nutrients and water (Liu et al., 2021a;
393	Liu et al., 2021b). Some studies have suggested that females have higher reproductive
394	costs than males in some dioecious plants due to the demand for carbon and water
395	during fruiting and seed set (Olano et al., 2017; Liu et al., 2021b). Therefore, females

tend to improve photosynthetic carbon gain and water uptake via higher hydraulic 396 conductivity and stomatal conductance under high-resource conditions (Hultine et al., 397 2016; Olano et al., 2017). Such a better ability for resource acquisition (including water) 398 in females is easily subjected to carbon starvation and hydraulic failure under drought 399 stress (Hultine et al., 2016). Indeed, female poplars have a lower leaf water potential, 400 photosynthetic rate and leaf water use efficiency, and thus they suffer more damage 401 from drought stress than males (Han et al., 2013; Yu et al., 2018). In contrast, males of 402 some dioecious plants show superior performance, which is associated with a more 403 404 conservative water use strategy to maximize drought stress tolerance (Olano et al., 2017). Interestingly, the sex-specific water acquisition and utilization might disappear 405 under well-watered conditions in *P. cathayana* (Chen et al., 2014), further suggesting 406 407 that sexual dimorphism in dioecious poplars is environment-dependent.

408

Drought-induced tree mortality is mainly attributable to carbon starvation or hydraulic 409 410 failure or both (Mitchell et al., 2013; Choat et al., 2018). Moreover, the photosynthetic carbon gain of leaves is associated with the hydraulic system in woody plants (Deans 411 et al., 2020; Qi et al., 2021). The hydraulic system of plants has been reported to 412 determine the ability of plants to cope with drought among plants with similar or 413 different genetic backgrounds (Blackman et al., 2017; Kannenberg et al., 2019). It has 414 been suggested that the tradeoff between hydraulic efficiency and safety affects plants' 415 tolerance to drought stress (Olano et al., 2017; Yao et al., 2021). Plant species with a 416 higher hydraulic conductance generally exhibit greater vulnerabilities to embolism 417

(Scoffoni et al., 2017). In contrast, plants with a hydraulic safety strategy emphasize the ability to maintain xylem conductivity with decreasing water potential levels (Olano et al., 2017; Scoffoni et al., 2017). Indeed, females with a greater theoretical hydraulic conductance are more sensitive to drought stress than males (Fig. 2). We found that females have a lower leaf water potential and photosynthesis but a greater theoretical hydraulic conductance further demonstrating the tradeoff between hydraulic efficiency and safety (Figs 1-2).

425

426 Adjustments in the hydraulic system can be assessed based on the xylem anatomy and function (Fonti et al., 2010). In this study, we found that water conditions modified 427 xylem anatomic traits of stem, affected the tradeoff between hydraulic efficiency and 428 429 safety, and caused sex-specific responses to water conditions (Fig. 2). In a previous study on P. cathayana, the secondary sexual dimorphisms in physiological characters 430 possibly disappeared under well-watered conditions (Chen et al., 2014). Here, we found 431 that such disappearance of secondary sexual dimorphism under well-watered 432 conditions appeared to be associated with the anatomic traits of stem xylem (Fig. 2). 433

434

Under well-watered conditions, females had a greater lumen area per vessel and predicted hydraulic conductivity. The proportion of xylem was greater in males than in females, although they had a lower lumen area per vessel and predicted hydraulic conductivity in males (Figs 1-2). The greater lumen area per vessel in males than females appeared to increase the total amount of water transported into shoots, thus

compensating for their lower hydraulic conductivity. Under well-watered conditions, 440 there was no significant difference in leaf potential between the sexes (Fig. 1), further 441 442 demonstrating the lack of significant secondary sexual dimorphism in *P. cathavana* in no-stress conditions. However, under drought stress, females still tended to adopt a 443 greater hydraulic efficiency (a higher theoretical hydraulic conductivity) to maximize 444 water transport. The increased lumen area per vessel in females elevates the risks of 445 cavitation and air embolism, thus easily resulting in hydraulic failure under drought, as 446 exhibited by the lower photosynthesis and leaf water potential compared to males. In 447 448 contrast, males preferred to adopt a more conservative water use strategy (a lower theoretical hydraulic conductivity) with a lower lumen area per vessel to maximize the 449 stress tolerance under drought stress. 450

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452 Sex-specific responses to drought and subsequent re-watering

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454 It has been suggested that the time for completing the recovery of physiological functioning is essential for tree survival. As proposed by Rehschuh et al. (2020), the 455 recovery ability and time in woody tree species after re-watering largely depends on 456 whether drought impairs the critical physiological processes, such as carbon uptake and 457 water transport. Here, males quickly recovered from drought after re-watering. The 458 extent of recovery in the leaf photosynthesis, leaf water potential and gas conductance 459 were greater in males than in females after 7 d with a favorable water status (Fig. 1). 460 Importantly, the treatment of 14 d re-watering after drought fully recovered leaf 461

photosynthesis and gas conductance in males, but did not fully recover those in females 462 (Fig. 1). As suggested by Brodribb and Cochard (2009), no hysteresis is expected when 463 the recovery of gas exchange is restricted by leaf water potential. In females, the leaf 464 water potential was fully restored within 14 d, but leaf photosynthesis did not fully 465 recover. The occurrence of hysteresis in females within 14 d recovery after drought 466 suggested that other non-hydraulic limitations probably continued to limit the leaf 467 photosynthesis after complete re-watering, and this needs to be explored in future 468 studies. 469

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As we discussed above, the recovery time largely depended on the extent of problems 471 in key physiological processes, such as carbon uptake and water transport. The carbon 472 473 uptake may be assessed by the leaf photosynthesis and gas conductance. Males with a faster rate of recovery from the drought stress may be associated with less impairment 474 in key physiological processes, such as the leaf photosynthesis and gas conductance, 475 476 when compared to females (Fig. 1). Recent studies have revealed that the xylem anatomical traits and function-associated hydraulic failure play an important role in 477 lowering survival during drought and subsequent recovery (Creek et al., 2018). 478 Additionally, the restoration of xylem function may be associated with modifications 479 in anatomical traits (Secchi et al., 2017). The lumen area per vessel and predicted 480 hydraulic conductivity of wood recovered to the control level in females at 7 d and 14 481 d re-watering after drought (Fig. 2). Interestingly, the vessel number was significantly 482 higher in both sexes under 7 d or 14 d re-watering after drought. This phenomenould 483

be associated with the legacy effects of drought. Some studies have suggested that
drought has long-lasting legacy effects on the ecosystem and plant growth (Arnone et
al., 2008; Kaisermann et al., 2017). However, the mechanism underlying such effects
on the xylem vessel number remains to be elucidated in future studies.

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It was noteworthy that the leaf water potential did not fully recover to the control levels 489 in either sex at 7 d re-watering after drought despite the full recovery of the predicted 490 hydraulic conductivity by this time (Figs, 1-2). This could be associated with the 491 492 carbohydrate metabolism during the embolism repair. According to the current active embolism removal models, the starch hydrolysis-induced osmotica accumulation in the 493 apoplast facilitated the refilling of the vessels and, thus, becoming functional (Secchi 494 495 and Zwieniecki, 2012; Pagliarani et al., 2019). In this study, we found that the NSC, including sugar and sucrose, was higher in males than in females (Fig. 3; Table 1). The 496 increased sugar contents in the male wood probably strengthened the ability of trees to 497 498 repair xylem functionality after drought stress. Importantly, the total sugar, sugar and NSC decreased at 7 d or 14 d re-watering after drought, but the contents were higher in 499 males than in females. Actually, the concentration of starch in the wood was lower in 500 males than females under all treatments (Fig. 3; Table 1). The lower starch 501 502 concentration in the male wood compared to females was probably attributable to starch hydrolysis into sugars to regulate the apoplastic water potential and the recovery of 503 504 empty vessel function. Indeed, males had higher sugar concentrations, and higher gene expression levels related to starch degradation in the wood compared to females (Fig. 505

3; Table 1). These results suggested that the xylem for drought stress recovery wasprobably associated with sugar metabolism in the wood.

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509 Transcriptome regulation of developing xylem underlying the molecular mechanism to
510 salt stress and re-watering

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Xylem anatomy and functions, as well as carbohydrate metabolism play critical roles 512 in drought tolerance and hydraulic recovery (Olano et al., 2017; Brunetti et al., 2020). 513 514 The sex-specific transcriptional regulation in response to drought further suggested that females and males adopt different strategies to cope with drought (Fig. 5). The vessel 515 development is initiated from the vessel differentiation and then continued through the 516 517 production of thick and lignified cell walls (Rodriguez-Zaccaro and Groover, 2019). The increased expression of auxin transporter, such as AUX1, PIN1 and PIN5, in males 518 under drought stress can facilitate the differentiation of vessel elements and enhance 519 520 vessel production (Fig. 6). Expression levels of genes related to the xylem cell fate, such as VND1, TED6, LBD15, members of the NAC transcription factor family, and 521 MYB transcription factors, were higher in males than in females under drought, in spite 522 of their reduced expression by drought stress relative to controls and re-watering at 523 either 7 d or 14 d (Fig. 6; Fig. S3). Such down-regulated genes induced by drought 524 stress in both sexes are probably associated with the lower proportion of xylem in the 525 526 cross-sections of stem (Fig. 1).

In addition, the vessel thickness affected the hydraulic conductance of stem (Olano et 528 al., 2017; Scoffoni et al., 2017). Plants with a smaller vessel diameter usually exhibit a 529 more conservative hydraulic strategy and stronger tolerance to drought stress, and the 530 smaller vessel diameter is usually associated with secondary cell wall deposition in 531 plants (Olano et al., 2017). Expression levels of genes related to secondary cell 532 formation, such as MYB transcription factors, NST1 (secondary wall thickening factors), 533 cellulose synthase family IRX1, IRX6 and IRX9 (Endo et al., 2015; Taylor-Teeples et 534 al., 2015), were higher in males than females under drought stress, in spite of the 535 536 reduced expression relative to controls (Fig. 6). Such gene expression modes under drought were consistent with the greater cell wall thickness and lower lumen area in 537 males compared to females (Fig. 2). It was noted that there appears to be inconsistency 538 539 in the expression patterns of genes related to cell wall synthesis and cell thickness in both sexes under drought stress. Such inconsistency may be explained by the large 540 demand for cell wall materials needed for cell expansion (Rodriguez-Zaccaro and 541 Groover, 2019). The expansion from 5 to 50 µm for the tracheary element needs 10 542 times more cell wall material to maintain a constant cell wall thickness (Rodriguez-543 Zaccaro and Groover, 2019). As shown in Fig. 2, the lumen area per vessel significantly 544 decreased by drought in both sexes when compared to the controls. The reduced 545 expression of genes related to cell wall synthesis and metabolism in both sexes under 546 drought stress relative to the controls was probably associated with the reduced demand 547 548 for cell wall material to cell expansion.

549

To recover growth after drought, plant cells must regain their ability to loosen their 550 walls and, over a longer period, incorporate new polymers to avoid wall failure (Rui 551 552 and Dinneny, 2020). In this study, re-watering after drought differentially recovered the expression of genes related to cell wall biosynthesis in both sexes (Fig. 6). The cell wall 553 and the microtubule cytoskeleton are also tightly linked through a mechanical feedback 554 loop to achieve growth and morphogenesis in plants (Eng and Sampathkumar, 2018). 555 The up-regulation of genes related to xylan, actin cytoskeleton organization, lignin 556 synthesis and metabolism, secondary cell wall biogenesis and metabolism, and 557 558 cytoskeleton organization in both sexes at 7 d and 14 d re-watering promoted the cell wall function recovery and probably turgor restoration (Fig. 5). The up-regulation of 559 genes related to xyloglucan metabolisms, such as expansins (EXPs), xyloglucan endo-560 561 β -transglucosylase/hydrolases (XETs/XTHs) and endo-1,4- β -glucanases (EGases), promote the cell wall extensibility (Sasidharan et al., 2011). The higher expression level 562 of genes related to the cell wall and the microtubule cytoskeleton induced by drought 563 564 in both sexes suggested that males had a better ability to recover cell wall expansiondriving cell enlargement and optimal plant growth. Interestingly, the sexual difference 565 in DEGs enriched in cell wall metabolism and cytoskeleton organization at 7 d re-566 watering disappeared at 14 d re-watering after drought, further demonstrating that 567 sexual dimorphism largely depends on abiotic stress factors (Chen et al., 2014; Liu et 568 al., 2020). 569

Drought reduced the plant growth by causing various physiological changes, including 571 loss of turgor (Le Gall et al., 2015). Ions and solutes absorbed by plants could lower 572 573 cells' osmotic potential to maintain turgor and withstand drought (Patakas et al., 2002). Increased transcript levels of some ABC transporters induced by drought in males might 574 contribute to the accumulation of ions and solutes, and to the maintenance of osmotic 575 potential. In addition, the up-regulation of some ion transporters induced by drought in 576 males relative to the controls, such as sulfate transporter, potassium transporter AKT2, 577 ADT4 and calcium transporter PHO1, facilitate the maintenance of turgor and drought 578 579 resistance (Patakas et al., 2002). Plasma membrane ATPases are primary active transporters of cations that maintain steep concentration gradients of ions (Morth et al., 580 2011). The up-regulation of genes related to ATP synthesis and activities facilitated the 581 582 proton-dependent transport, which was responsible for metabolites and other ion transport, and drought stress tolerance in males (Fig. S3). 583

584

585 In addition, the increased NSC storage plays a critical role in osmotic regulation to maintain turgor and/or refill the embolized xylem under drought and re-watering after 586 drought (Pagliarani et al., 2019). Under drought stress, the xylem sugar accumulation 587 is derived from either starch degradation in parenchyma cells or translocation from 588 phloem to parenchyma rays (Pagliarani et al., 2019). The pathway of starch degradation 589 is regulated by phosphoglucan water dikinase, phosphoglucan phosphatases starch 590 excess 4 (SEX4), disproportionating enzyme (DPE1/2) (Chia et al., 2004), starch 591 debranching enzymes (LSA3, LDA) and β -amylases (BAM1/3) (Fulton et al., 2008). 592

Additionally, maltose excess protein (MEX1) transports glucose and maltose to the 593 cytosol (Cho et al., 2011; Fan et al., 2018). In our study, the up-regulation of GWD1, 594 *PWD1*, SEX4, *DPE1*, *DPE2* and *MEX1* genes induced by drought stress and subsequent 595 re-watering in males relative to females was consistent with their lower starch content 596 and higher sugar levels compared to females. The up-regulation of starch degradation 597 and sucrose transport in males under drought and re-watering lowered cell osmotic 598 potential to maintain turgor and hydraulic refill, thus enhancing drought tolerance. 599 Interestingly, we found that only the BAM1 gene was up-regulated under drought in 600 601 females but not under subsequent re-watering relative to the controls, further suggesting that females are more sensitive to drought and have greater difficulties to recover from 602 drought relative to males. 603

604

In addition, the up-regulation of membrane sugar transporters, SUT4, in males under 605 drought and subsequent re-watering promoted the apoplastic sugar accumulation and 606 607 aquaporin-mediated water entry into the empty vessels immediately when re-watering after drought. Indeed, we found that most genes encoding the aquaporins, such as 608 PIP1;4, PIP2 and PIP3, were up-regulated in males but not in females at 7 d re-609 watering after drought stress. Interestingly, we found that the expression of genes 610 related to aquaporins, such as PIP1;3, PIP1B, TIP1;3, SIP1;2 and NIP1;2, was higher 611 in females than in males under well-watered and drought stress conditions (higher than 612 in controls). The expression modes of these genes related to aquaporins further 613 demonstrated that females have a greater hydraulic efficiency, both under drought and 614

615	well-watered conditions, which easily makes females more sensitive to drought. These
616	results suggested that males have a better ability to cope with drought and recover better
617	and faster from drought stress compared to females, which is largely dependent on the
618	regulation of the xylem structure and function, as well as on the osmotic maintenance
619	at the molecular level.
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625	Conclusions
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627	The present study showed that males have a greater tolerance to drought and a better
628	ability of recover from drought compared to females, which was associated with sex-
629	specific anatomical traits of xylem and carbohydrate dynamics. Males showed xylem
630	anatomical traits related to a conservative water-use strategy (a lower theoretical
631	hydraulic conductivity) and higher leaf photosynthesis. In contrast, females had a
632	greater hydraulic efficiency over safety (thinner tracheid walls and a larger lumen area
633	per vessel), more sensitivity to drought and a slower recovery of leaf photosynthesis.
634	The transcription analysis conducted for the developing xylem further demonstrated
635	that the tolerance of drought and quickly recovery in males was associated with the
636	processes of cell wall biogenesis, starch metabolism and ion transport.

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641	analysis and writing, Yang Zhao and Yuting Wang contributed to data collection, Helena
642	Korpelainen contributed to the interpretation of data and manuscript preparation, and
643	Chunyang Li (the corresponding author) had the overall responsibility for experimental
644	design and project management.
645	Conflict of interest The authors declare that they have no conflict of interest.
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881 Figure legends

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Figure 1 The photosynthetic rate (a), gas conductance (b), leaf water potential (c) and proportion of xylem (d) in *P. cathayana* females and males under well-watered (CK) or drought stress (D) conditions, and 7 d (DW7) or 14 d re-watering (DW14) after drought stress. Bars indicate means \pm SD (n = 4). Different lower-case letters above bars indicate significant differences at P < 0.05 based on ANOVA followed by Duncan's tests.

Figure 2 The vessel number (a), numeriarea per vessel (b), cell wall thickness (c) and predicted conductivity (d) in the xylems of *P. cathayana* females and males under wellwatered (CK) or drought stress (D) conditions, and 7 d (DW7) or 14 d re-watering after drought stress (DW14). Bars indicate means \pm SD (n = 4). Different lower-case letters above bars indicate significant differences at *P* < 0.05 based on ANOVA followed by Duncan's tests.

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Figure 3 The levels of total sugar (a), sucrose (b), starch (c) and non-structural carbohydrates (NSC) (d) in the xylems of *P. cathayana* females and males under wellwatered (CK) or drought stress (D) conditions, and 7 d (DW7) or 14 d re-watering after drought stress (DW14). Bars indicate means \pm SD (n = 4). Different lower-case letters above bars indicate significant differences at P < 0.05 based on ANOVA followed by 901 Duncan's tests.

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903	Figure 4 FTIR spectra (a) and the corresponding principle component analysis (PCA)
904	(b) plot of the xylems of <i>P. cathayana</i> females and males under well-watered (CK) or
905	drought stress (D) conditions, and 7 d (DW7) or 14 d re-watering after drought stress
906	(DW14). The average spectrum of the developing xylem was plotted ($n = 4$). PCA was
907	conducted with the data of the selected peaks separately for leaves and roots (Table S2).
908	The spectral peaks and their corresponding molecular components were as follows: 1,
909	1738 cm ⁻¹ for vibration of C=O from esterified pectin; 2, 1650 cm ⁻¹ for C-N vibration
910	from protein; 3, 1596 cm for aromatic skeletal vibrations in lignin; 4, 1505 cm ⁻¹ for
911	C=O stretch and aromatic skeletal vibrations in lignin; 5, 1462 cm ⁻¹ C-H deformation,
912	lignin and carbohydrates; 6, 1425 cm ⁻¹ for vibration of COO- from pectin; 7, 1375 cm ⁻
913	¹ for C-H deformation in hemicellulose and cellulose; 8, 1330 cm ⁻¹ for guaiacyl ring
914	condensed plus syringyl ring; 9, 1235 cm ⁻¹ for deformation of cellulose; 10, 1157 cm ⁻¹
915	for vibration of C-C and C-O stretch from carbohydrates (such as soluble sugar,
916	cellulose and hemicellulose; 11, 1111 cm ⁻¹ for C-C or C-O vibration from pectin; 12,
917	1034 cm ⁻¹ for vibration of C-H or C-C and C-O from cellulose and hemicellulose.

Figure 5 Comparative profiles of transcriptional responses to drought stress, or 7 d or
14 d re-watering after drought stress in the developing xylem of *P. cathayana* females
and males. (a) Venn diagrams of common and sex-specific gene ontology (GO)
categories under well-watered (W) or drought stress (D), 7 d (DW7) or 14 d (DW14)

re-watering after drought stress. (b) Circle diagram of GO terms enriched in up-923 regulated genes (above) and down-regulated genes (below) between females and males 924 925 under well-watered or drought stress conditions, and 7 d or 14 d re-watering after drought stress. (c) Total number of differentially expressed genes. Blue bars mean up-926 regulated genes between treatments, and orange bars mean down-regulated genes 927 between treatments. (d) Comparative transcriptional profiling showing the up-regulated 928 genes between females and males. The X-axis indicates the GO term, and the Y-axis 929 indicates the rich factor (the ratio of gene expression level enriched in males relative to 930 931 fe males). The higher rich factor means the greater enrichment. The size of a circle indicates the number of genes in the GO term, and the color of a circle indicates the P-932 value range. By default, the enrichment result of top 20 is displayed on the premise of 933 934 the *P*-value < 0.05. FD, female + drought stress; MD, male + drought stress; MW, male + well-watered; MDW7, male + re-watering 7 d after drought stress; MDW14, male + 935 re-watering 7 d after drought stress; FW, female + well-watered; FDW7, female + re-936 937 watering 7 d after drought stress; FDW14, female + re-watering 14 d after drought stress. 938

Figure 6 Hierarchical clustering of key genes involved in starch auxin transport, tracheary element differentiation, secondary cell wall biogenesis (a), as well as in sucrose transport and starch metabolism (b) in the developing xylem of *P. cathayana* females and males. The columns show the samples, and the rows show the genes. The colors of cells represent the log10 (TPM) values in each sample. Red cells indicate that genes are highly expressed in the samples, and blue cells indicate that genes are lowly

945	expressed in the samples. FD, female + drought stress; MD, male + drought stress; MW,
946	male + well-watered; MDW7, male + re-watering 7 d after drought stress; MDW14,
947	male + re-watering 7 d after drought stress; FW, female + well-watered; FDW7, female
948	+ re-watering 7 d after drought stress; FDW14, female + re-watering 14 d after drought
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1023 Figure 5



1028 Figure 6



Table 1 Leaf water potential (ψ), leaf dry weight (DW), stem DW, root DW, and wood total sugar, sucrose, starch and non-structural carbohydrates (NSC) in *P. cathayana* females (F) and males (M), as affected by well-watered (CK) or drought stress (D) conditions, and 7 d (DW7) and 14 rewatering (DW14) after drought stress. Different letters in the column indicate significant differences between treatments (*P* < 0.05, Duncan's test).

Treatment Leaf ψ (MPa) Leaf DW (g) Stem DW (g) Root DW (g) Wood Sex Total sugar (mg m⁻¹) Sucrose (mg g^{-1}) Starch (mg g^{-1}) NSC (mg g^{-1}) CK -0.315±0.01d 20.54±0.58a 25.65±1.50a 32.07±1.46cd 6.66±2.32e 42.55±2.33b F 5.67±0.59a 11.23±1.38b D F -1.02±0.01a 4.61±0.60g 6.21±0.56d 2.27±0.36c 36.12±1.09bc 23.13±2.08b 17.01±0.59a 53.13±1.59a DW7 F -0.721±0.05b 5.77±0.80f 6.99±0.87d 3.22±0.41b 30.24±1.97de 12.45±1.80d 12.28±0.91b 42.52±2.37b -0.34±0.02d 8.54±0.38d 8.79±0.80c 3.57±0.13b 15.93±2.83f 7.92±1.46e 11.38±0.89b 25.06±2.37d DW14 F CK -0.358±0.03d 12.27±0.47b 12.77±2.15d 1.78±0.28e 29.77±3.67c Μ 21.41±1.72b 3.75±0.41b 28.25±2.50de 28.50±2.49a 10.96±1.57b 57.33±3.46a 4.85±0.47g 5.89±0.67d 1.86±0.67c 46.12±3.45a D Μ -0.792±0.07b DW7 -0.454±0.04c 7.09±0.51e 7.10±0.59d 3.39±0.29b 39.27±5.33b 19.14±1.12c 7.25±0.69c 45.25±5.18b Μ 10.61±0.78c 25.75±3.74e 13.44±2.05d 4.70±0.59d 33.20±2.27c DW14 М -0.296±0.06d 10.12±0.83c 5.66±0.50a

Values are expressed as means \pm SE (n = 4).