

## *Supplementary Material*

# **Genome-wide association study of wood anatomical and morphological traits in *Populus trichocarpa***

**Hari B. Chhetri, Anna Furches, David Macaya-Sanz, Alejandro Riveros Walker, David Kainer, Piet Jones, Anne E. Harman-Ware, Timothy J. Tschaplinski, Daniel Jacobson, Gerald A. Tuskan<sup>2</sup>, Stephen P. DiFazio\***

**\* Correspondence:**

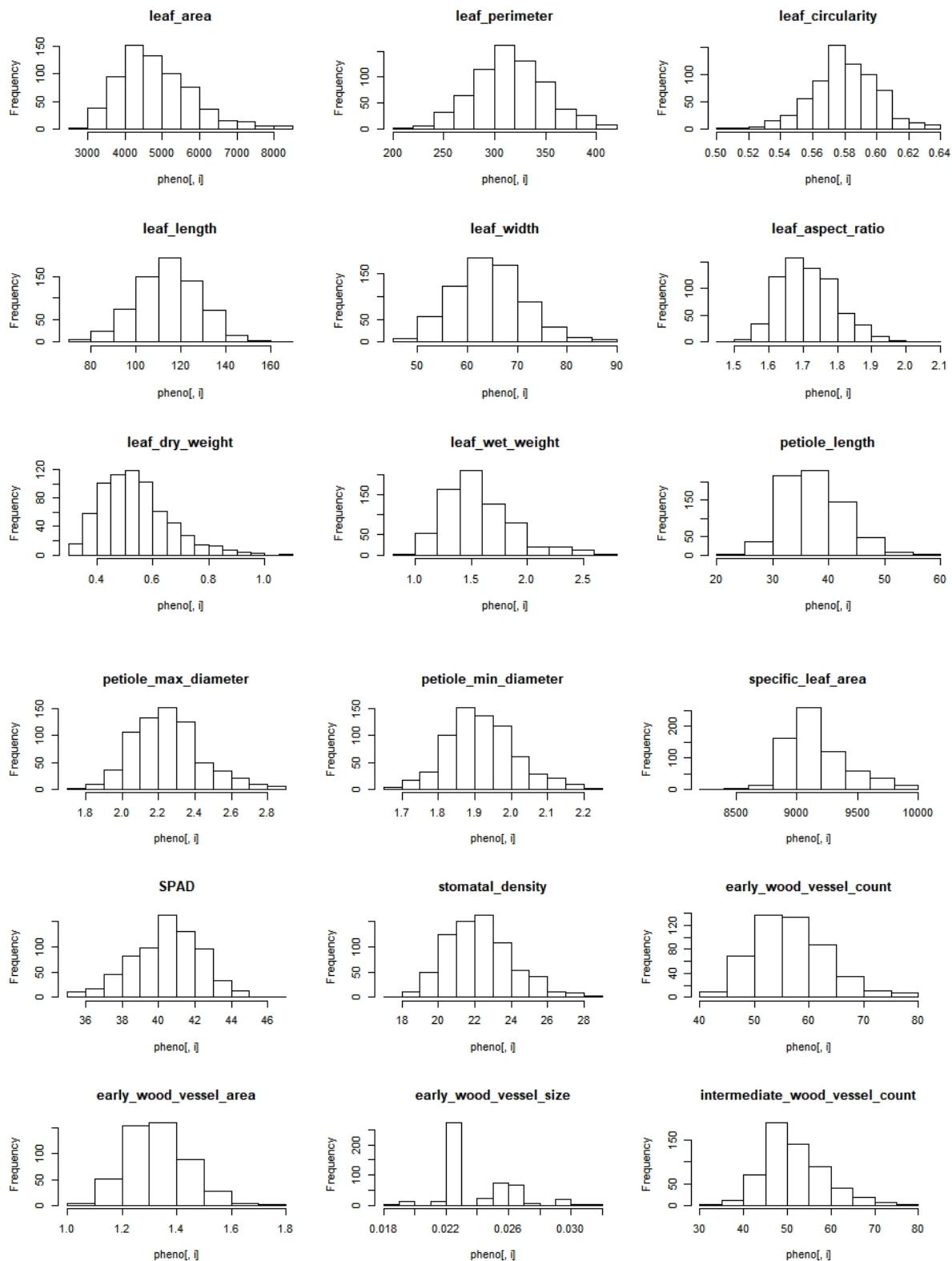
Stephen DiFazio

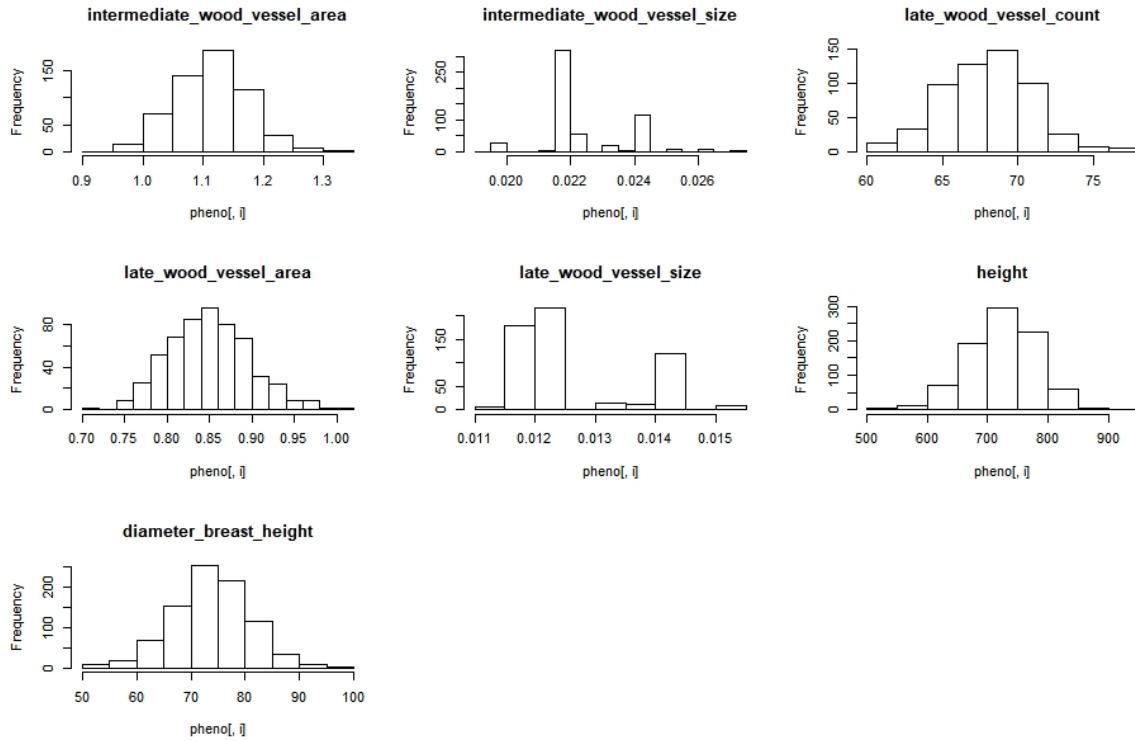
[spdifazio@mail.wvu.edu](mailto:spdifazio@mail.wvu.edu)

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## **1      Supplementary Figures**



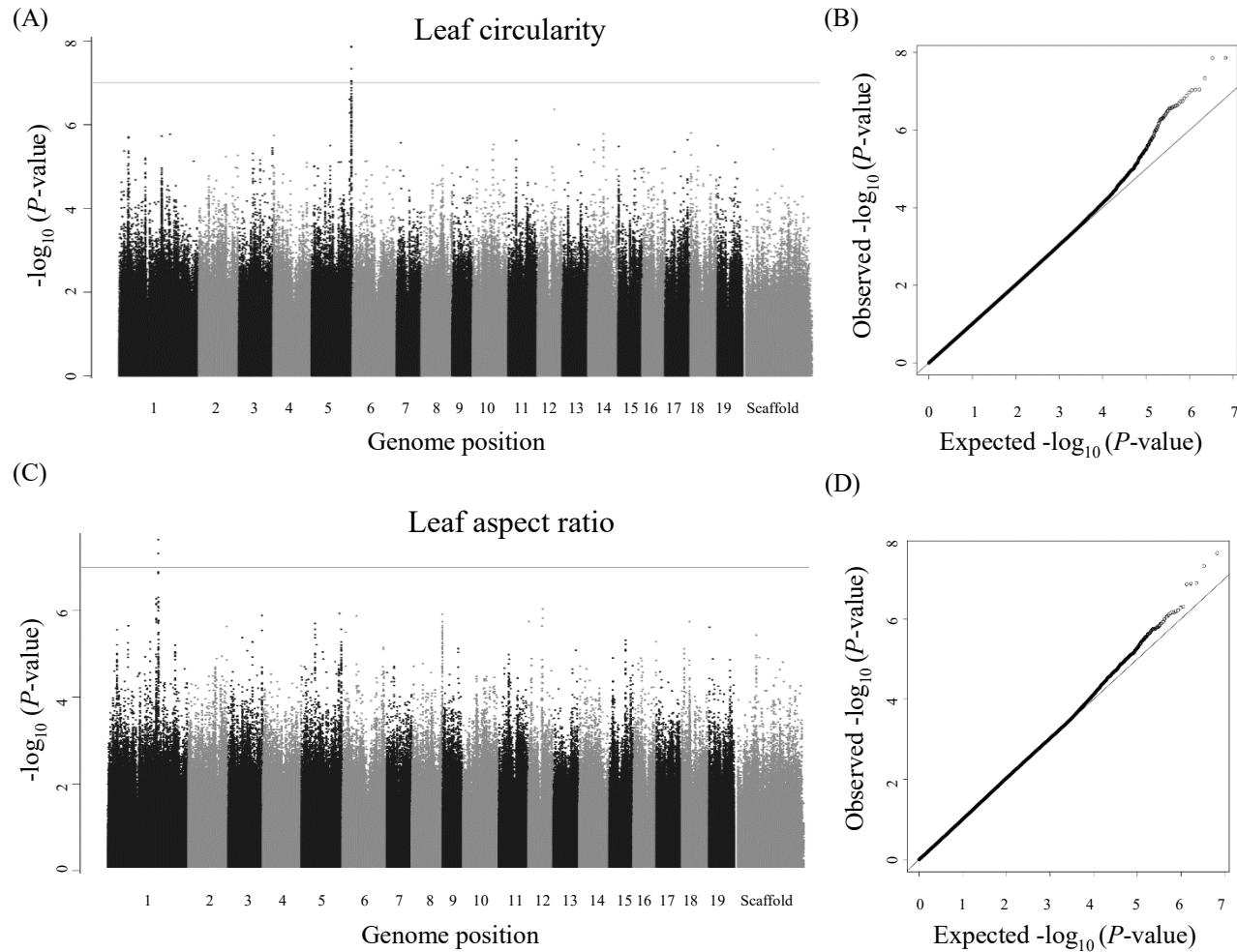


**Figure S1.** Histograms showing distribution of BLUP adjusted phenotypic values (BLUPs added to the mean phenotypic value) for each of the 25 morphological and anatomical traits used in GWAS study.

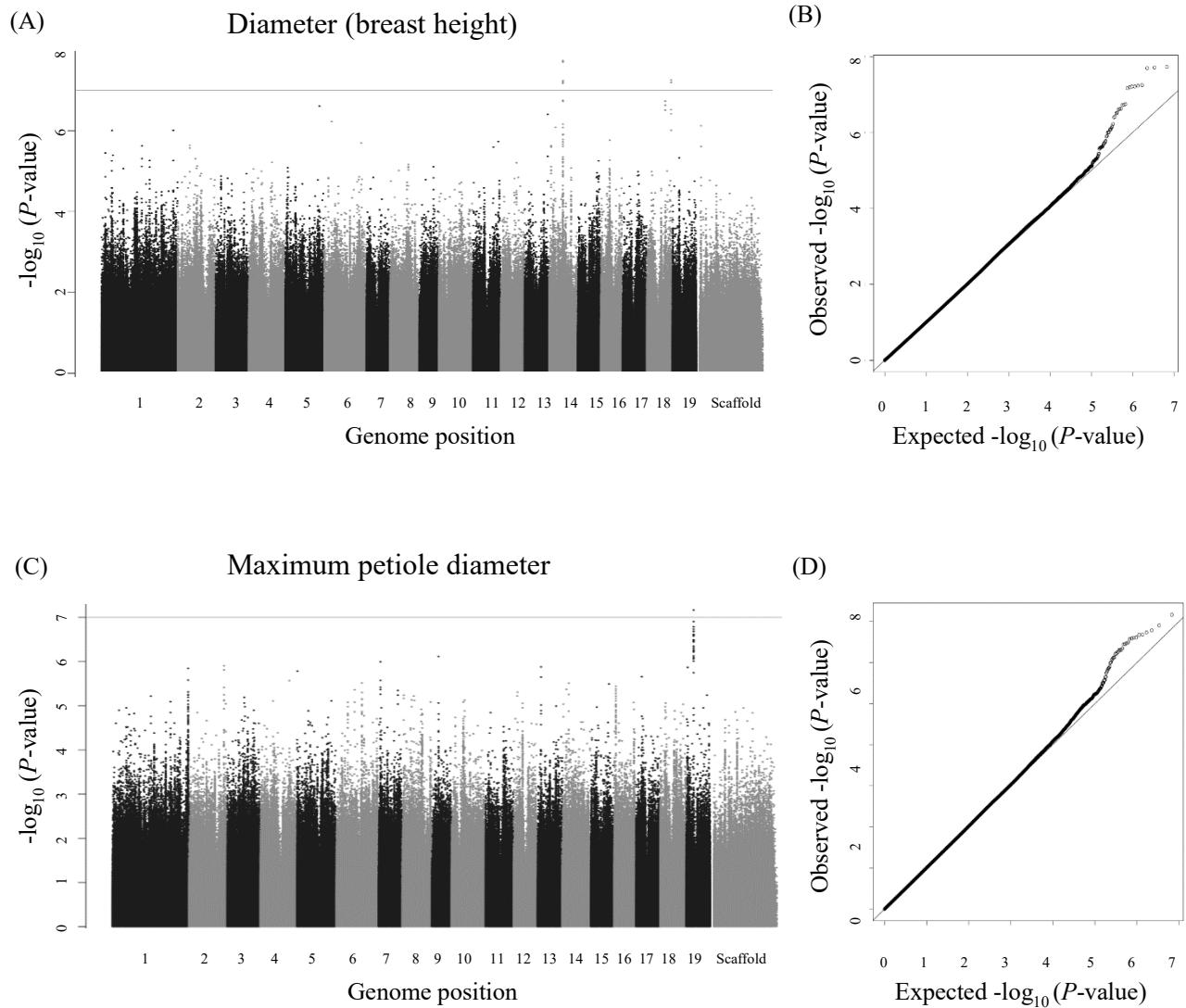
Trait	Diameter (breast height)	Height	MORPHOLOGY	WOOD ANATOMY
Diameter (breast height)				
Height				
Leaf area	0.025 <b>-0.039</b> Leaf area			
Leaf perimeter	0.042 <b>-0.012</b> 0.052 Leaf perimeter			
Leaf circularity	-0.046 <b>-0.073</b> 0.064 Leaf circularity			
Leaf length	0.031 <b>-0.016</b> 0.089 0.058 -0.243 Leaf length			
Leaf width	0.004 <b>-0.057</b> 0.058 0.003 0.179 0.004 Leaf width			
Leaf aspect ratio	0.069 <b>-0.078</b> -0.244 -0.084 -0.054 0.117			
Leaf dry weight	0.006 <b>-0.057</b> 0.094 0.080 -0.039 0.068 0.840 -0.131 Leaf dry weight			
Leaf wet weight	0.051 <b>-0.022</b> 0.034 0.086 0.014 0.848 0.877 <b>-0.187</b> 0.931 Leaf wet weight			
Leaflet length	-0.064 <b>-0.081</b> 0.601 0.620 -0.054 0.631 0.619 -0.144 0.707 0.566 Leaflet length			
Petiole diameter (max)	0.029 <b>-0.047</b> 0.819 0.793 0.029 0.753 0.782 -0.181 0.837 0.880 0.551 Petiole diameter (max)			
Petiole diameter (min)	0.034 <b>-0.032</b> 0.525 0.535 0.009 0.077 0.021 0.132 0.742 0.848 0.348 Petiole diameter (min)			
Specific leaf area	0.008 <b>-0.007</b> 0.008 0.004 <b>-0.059</b> 0.021 <b>-0.079</b> <b>-0.322</b> <b>-0.104</b> <b>-0.347</b> -0.257 -0.039 Specific leaf area			
SPAD	0.124 <b>0.105</b> 0.241 0.307 <b>-0.214</b> 0.375 0.215 0.179 0.398 0.279 0.455 0.253 0.148 <b>-0.499</b> SPAD			
Stomatal density	-0.112 <b>-0.063</b> <b>-0.198</b> <b>-0.174</b> <b>-0.043</b> <b>-0.157</b> 0.028 <b>-0.187</b> <b>-0.234</b> <b>-0.058</b> <b>-0.220</b> <b>-0.224</b> -0.045 -0.047 Stomatal density			
Early wood vessel count	-0.065 <b>-0.003</b> <b>-0.065</b> <b>-0.058</b> <b>-0.002</b> <b>-0.046</b> -0.060 <b>-0.058</b> <b>-0.081</b> <b>-0.058</b> <b>-0.067</b> <b>-0.062</b> <b>-0.047</b> <b>-0.004</b> <b>-0.057</b> <b>-0.052</b> <b>-0.078</b> 0.037 Early wood vessel count			
Early wood vessel area	0.243 <b>0.266</b> <b>0.047</b> <b>0.047</b> <b>0.030</b> <b>0.037</b> 0.059 <b>-0.027</b> <b>0.015</b> 0.041 <b>0.054</b> 0.039 <b>0.013</b> 0.030 0.028 <b>-0.030</b> <b>-0.020</b> -0.096 0.557 Early wood vessel area			WOOD ANATOMY
Early wood vessel size	0.296 <b>0.246</b> <b>0.059</b> <b>0.054</b> <b>0.035</b> <b>0.018</b> 0.075 <b>-0.113</b> <b>0.026</b> 0.026 <b>0.066</b> 0.037 <b>0.003</b> 0.102 0.026 <b>-0.049</b> <b>-0.080</b> -0.105 <b>-0.432</b> 0.345 Early wood vessel size			
Intermediate wood vessel count	<b>-0.305</b> <b>-0.217</b> <b>-0.028</b> <b>-0.025</b> <b>-0.015</b> <b>-0.007</b> -0.032 <b>0.049</b> <b>-0.014</b> -0.011 <b>-0.057</b> <b>-0.017</b> <b>-0.010</b> <b>-0.089</b> -0.142 <b>0.049</b> 0.140 0.056 <b>0.449</b> 0.023 <b>-0.366</b> Intermediate wood vessel count			
Intermediate wood vessel area	-0.052 <b>0.018</b> <b>0.020</b> <b>0.036</b> <b>-0.054</b> 0.042 0.022 <b>0.030</b> <b>0.020</b> 0.040 <b>0.010</b> 0.031 <b>0.006</b> -0.057 -0.067 -0.013 0.107 0.029 <b>0.268</b> 0.343 0.043 <b>0.610</b> Intermediate wood vessel area			
Intermediate wood vessel size	0.301 <b>0.231</b> <b>0.051</b> <b>0.070</b> <b>-0.070</b> 0.068 0.039 <b>0.033</b> <b>0.036</b> 0.062 <b>0.064</b> 0.053 <b>0.021</b> 0.075 0.109 <b>-0.077</b> <b>-0.106</b> -0.073 <b>-0.189</b> 0.249 <b>0.408</b> <b>-0.496</b> 0.175 Intermediate wood vessel size			
Late wood vessel count	<b>-0.231</b> <b>-0.188</b> <b>-0.013</b> <b>-0.017</b> 0.011 <b>-0.013</b> -0.017 <b>0.017</b> <b>-0.016</b> -0.009 <b>-0.019</b> 0.022 <b>0.019</b> <b>-0.010</b> -0.148 <b>0.060</b> 0.157 0.072 <b>0.349</b> 0.117 <b>-0.217</b> 0.586 0.399 <b>-0.296</b> Late wood vessel count			
Late wood vessel area	-0.038 <b>0.045</b> <b>-0.077</b> <b>-0.067</b> <b>-0.020</b> <b>-0.072</b> -0.083 <b>0.043</b> <b>-0.090</b> -0.038 <b>-0.118</b> <b>-0.036</b> 0.001 <b>0.047</b> -0.142 <b>-0.002</b> 0.167 0.025 <b>0.274</b> <b>0.250</b> <b>-0.034</b> 0.403 <b>0.531</b> 0.035 <b>0.633</b> Late wood vessel area			
Late wood vessel size	0.167 <b>0.191</b> <b>-0.062</b> <b>-0.056</b> <b>-0.025</b> <b>-0.057</b> -0.077 <b>0.037</b> <b>-0.073</b> <b>-0.040</b> <b>-0.105</b> <b>-0.071</b> <b>-0.051</b> 0.064 <b>-0.025</b> <b>-0.068</b> <b>-0.010</b> -0.006 <b>-0.052</b> 0.073 <b>0.145</b> <b>-0.176</b> 0.016 <b>0.265</b> <b>-0.426</b> 0.193 Late wood vessel size			

**Figure S2.** Pairwise correlation ( $r$ ) of *P. trichocarpa* wood anatomical and morphological traits in Clatskanie, Oregon. Shaded values greater than 0.15 or less than -0.15 are significant based on the Bonferroni correction criteria at 5% significance level.

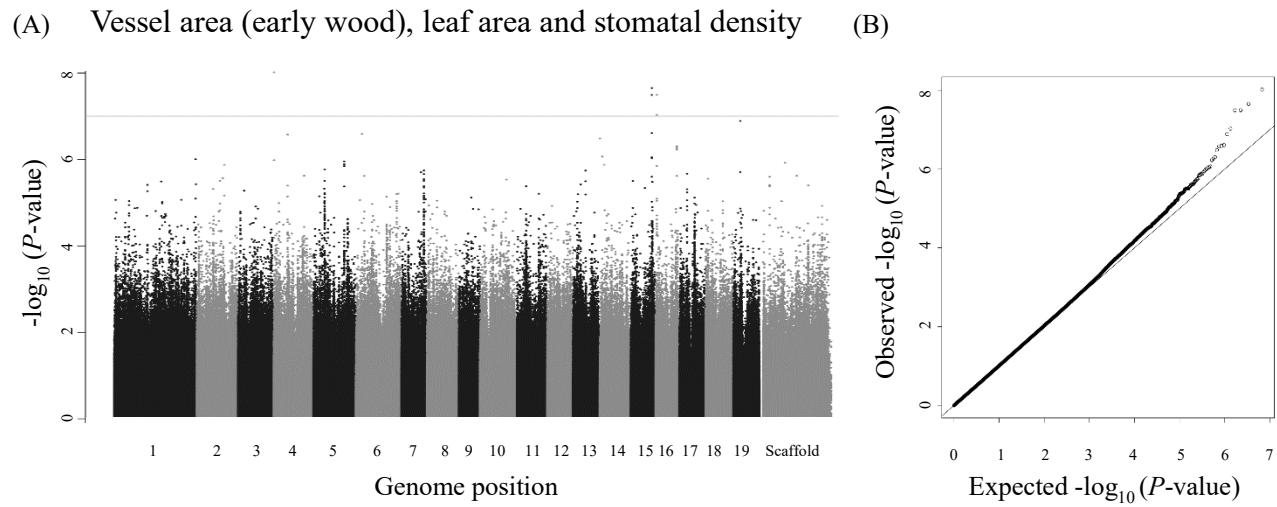




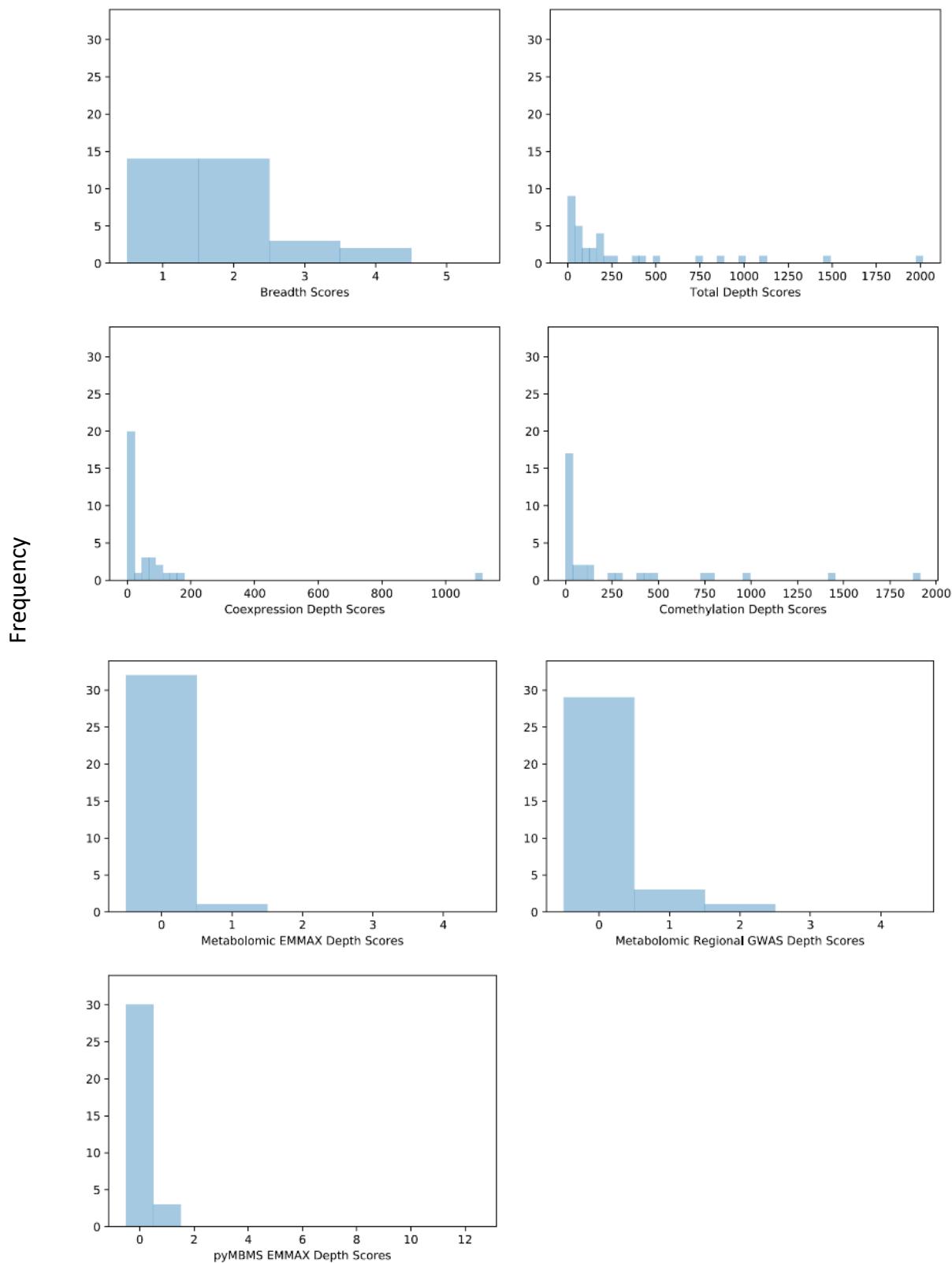
**Figure S5.** Single trait GWAS – Manhattan (left) and QQ plots (right). Numbers 1 to 19 represent chromosomes; scaffolds are the reads that did not align to any of the 19 chromosomes. (A) and (B) Leaf circularity; (C) and (D) Leaf aspect ratio. SNPs above gray line have  $p < 1 \times 10^{-7}$ , which is roughly equivalent to  $\text{FDR} \leq 0.1$  and SNPs above black line (the top line) have  $p < 7.417 \times 10^{-9}$  (the Bonferroni correction threshold), which is roughly equivalent to  $\text{FDR} \leq 0.05$  in this study.



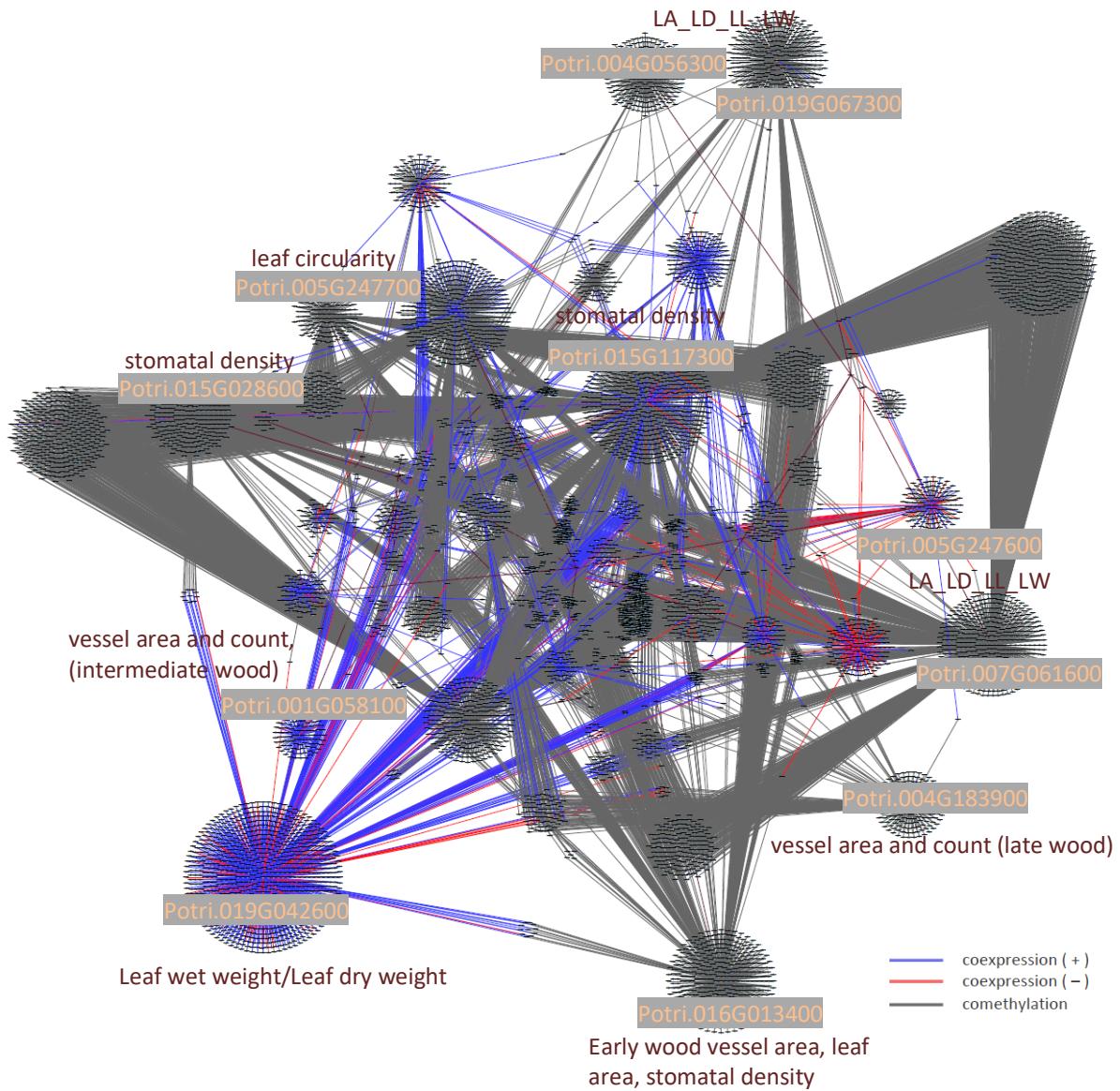
**Figure S6.** Single trait GWAS – Manhattan (left) and QQ plots (right). Numbers 1 to 19 represent chromosomes; scaffolds are the reads that did not align to any of the 19 chromosomes. (A) and (B) Diameter (breast height); (C) and (D) Maximum petiole diameter. SNPs above gray line have  $p < 1 \times 10^{-7}$ , which is roughly equivalent to FDR  $\leq 0.1$  and SNPs above black line (the top line) have  $p < 7.417 \times 10^{-9}$  (the Bonferroni correction threshold), which is roughly equivalent to FDR  $\leq 0.05$  in this study.



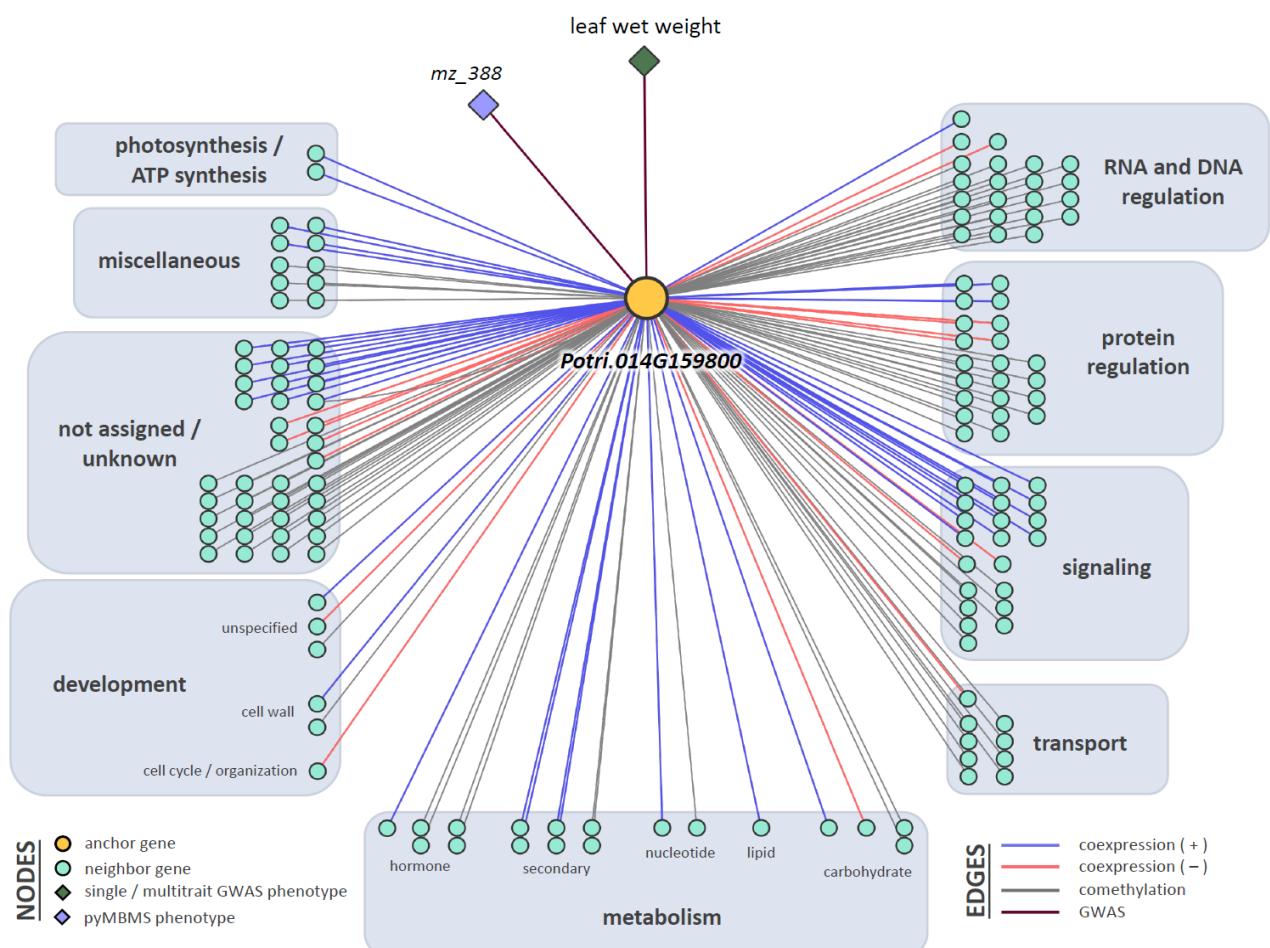
**Figure S7.** Multitrait GWAS – Manhattan (left) and QQ plots (right). Numbers 1 to 19 represent chromosomes; scaffolds are the reads that did not align to any of the 19 chromosomes. (A) and (B) Vessel area (early wood), leaf area and stomatal density. SNPs above gray line have  $p < 1 \times 10^{-7}$ , which is roughly equivalent to  $\text{FDR} \leq 0.1$  and SNPs above black line (the top line) have  $p < 7.417 \times 10^{-9}$  (the Bonferroni correction threshold), which is roughly equivalent to  $\text{FDR} \leq 0.05$  in this study.



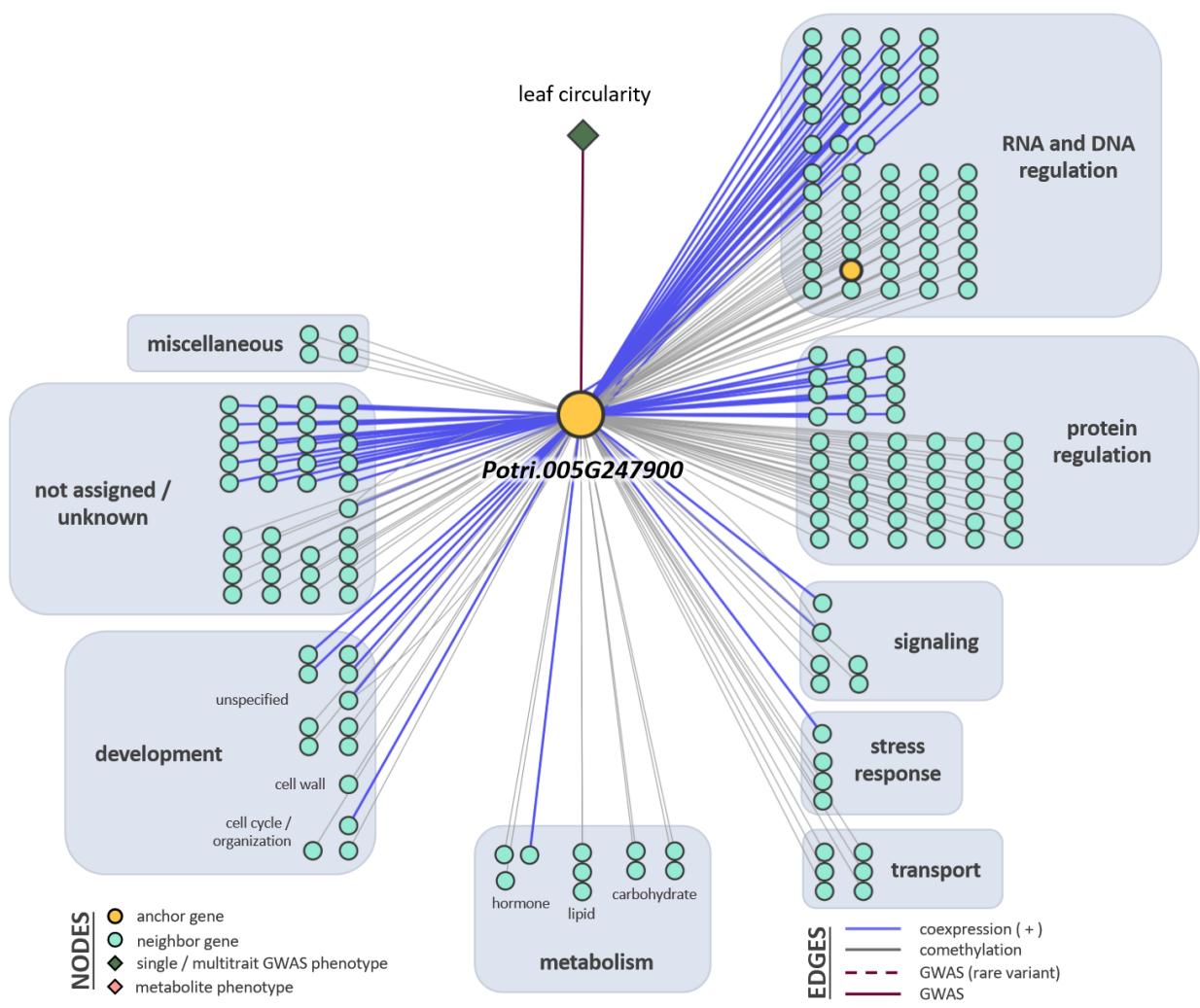
**Figure S8.** Distributions of LOE scores for all single trait and multitrait anchor (GWAS) genes.



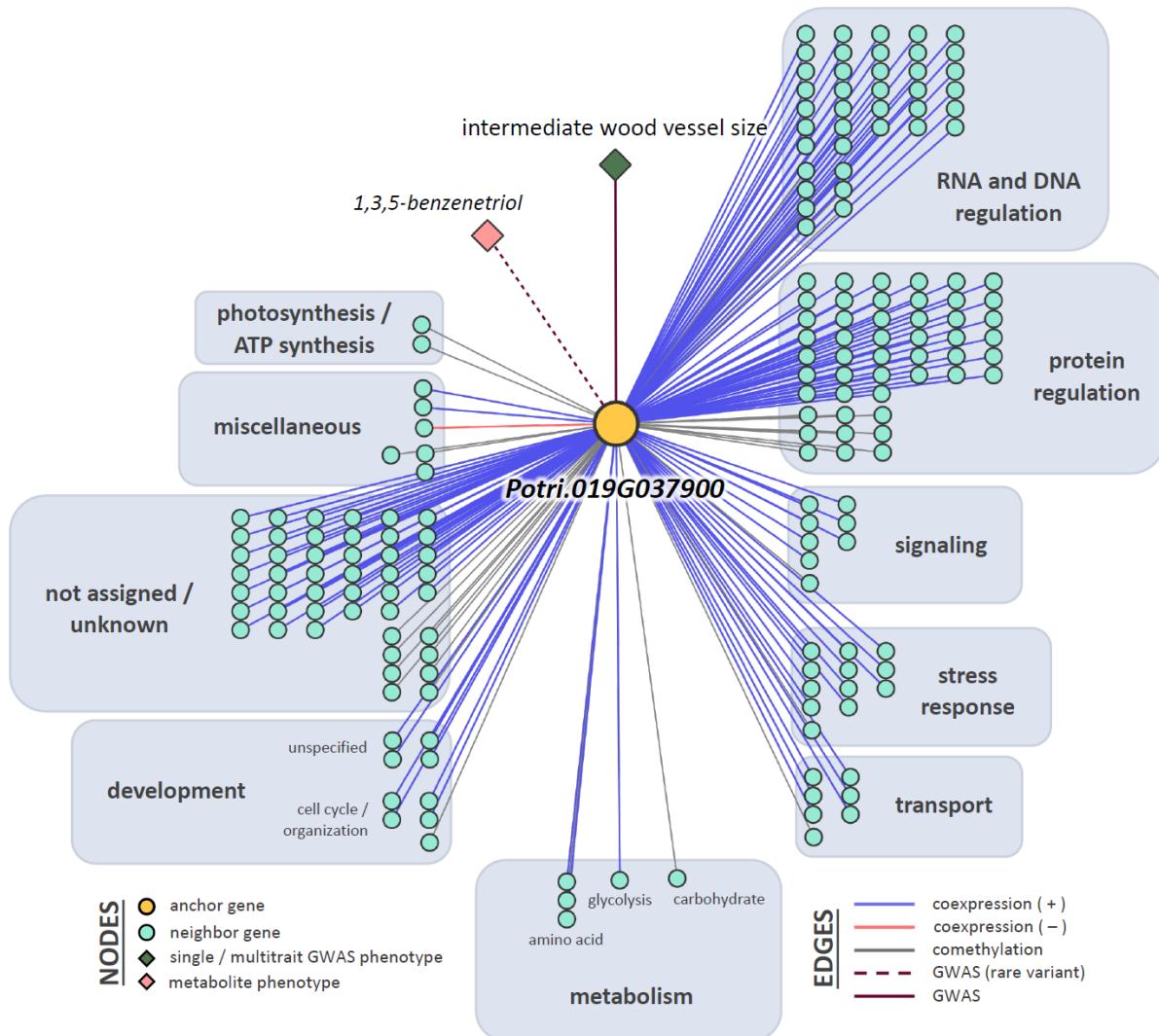
**Figure S9.** One-hop multi-omic network for all single and multitrati GWAS (anchor) genes with major anchor genes highlighted. LA\_LD\_LL\_LW = leaf area, leaf dry weight, leaf length and leaf width. For GWAS edges, solid maroon indicates “traditional” GWAS and dashed maroon indicates RV metabolite GWAS.



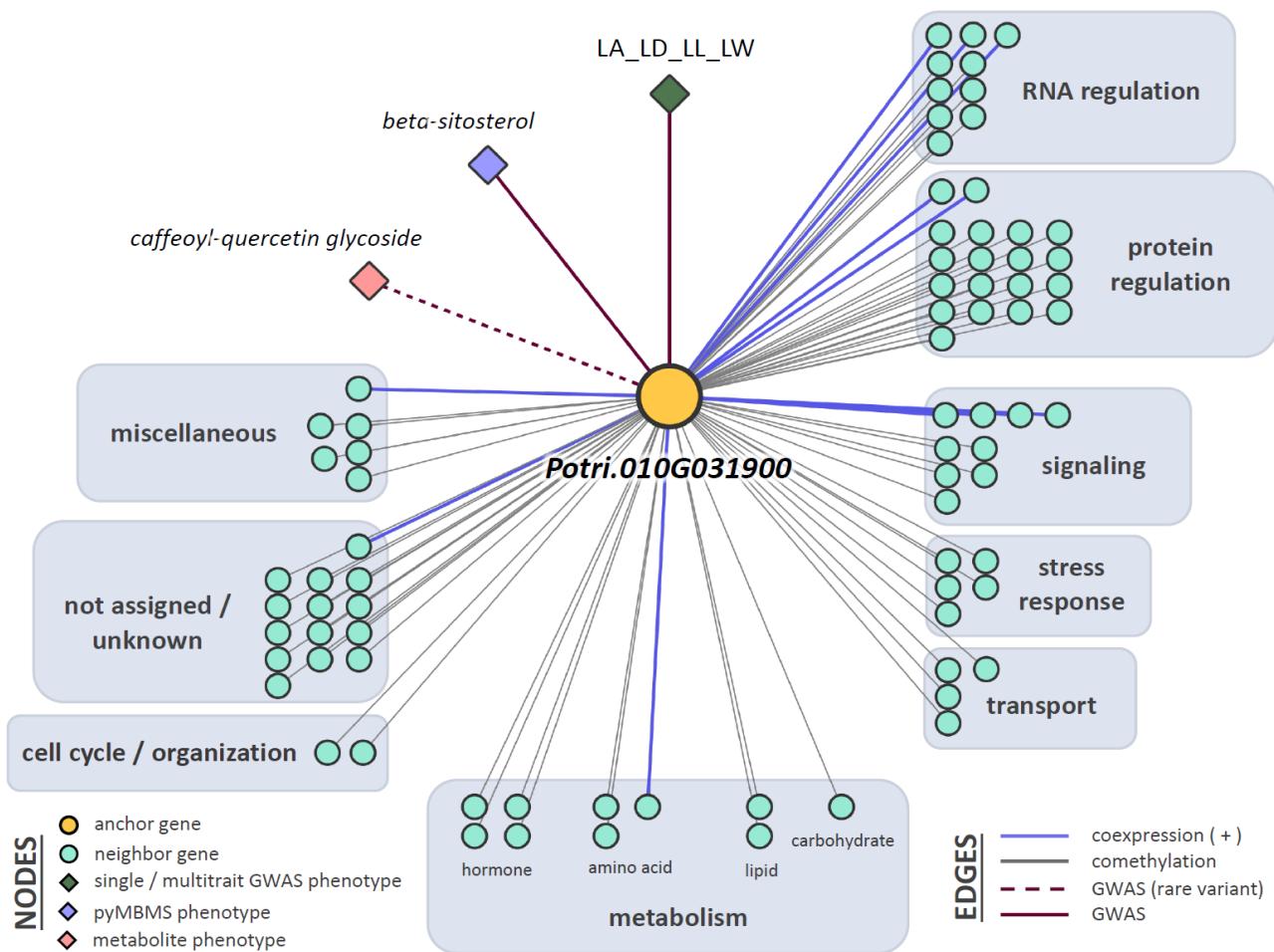
**Figure S10.** One-hop multi-omic network for the gene *Potri.014G159800*, which was associated with leaf wet weight.



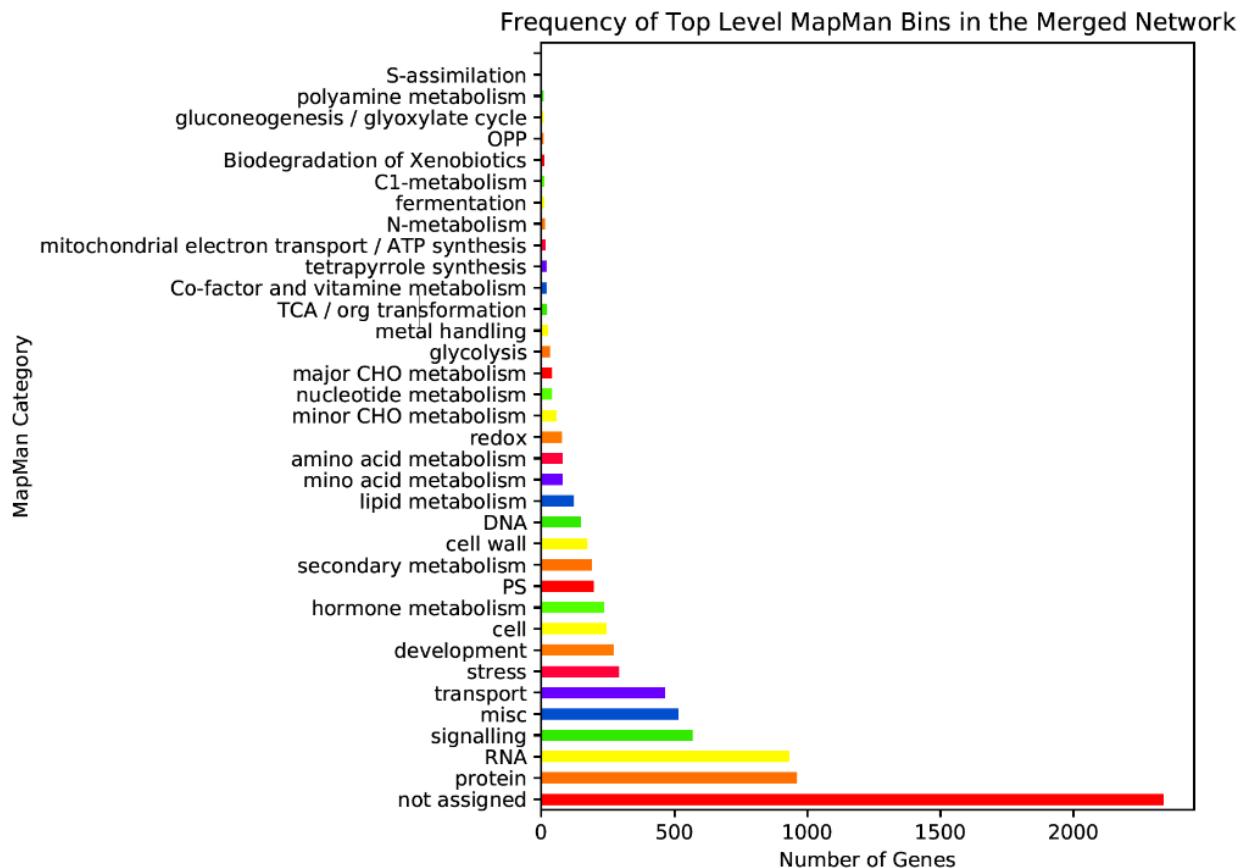
**Figure S11.** One-hop multi-omic network for the gene *Potri.005G247900*, which was associated with leaf circularity.



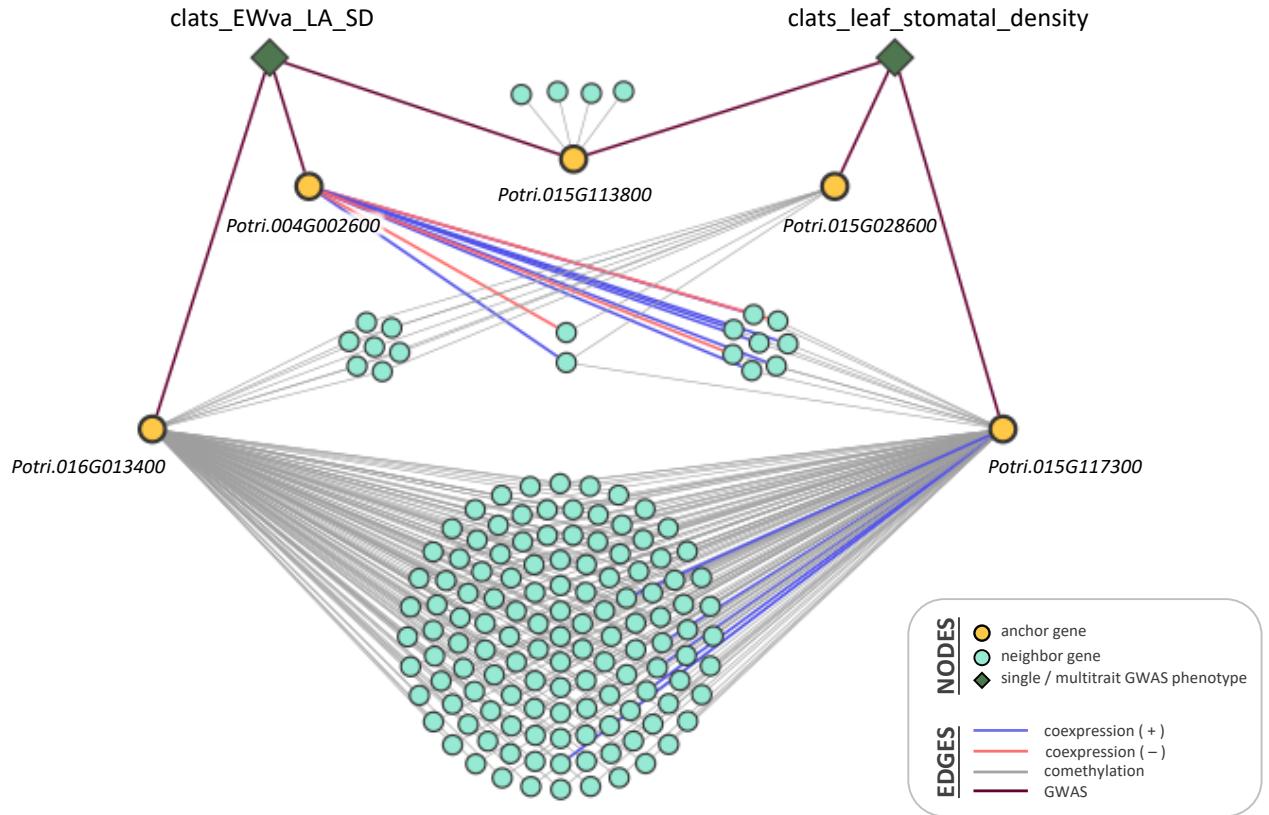
**Figure S12.** One-hop multi-omic network for the gene *Potri.019G037900*, which was associated with intermediate wood vessel size.



**Figure S13.** One-hop multi-omic network for the gene Potri.010G031900, which was associated with the multitrait phenotype leaf area, leaf dry weight, leaf length and leaf width.



**Figure S14.** Mapman annotations for all single trait and multitrait GWAS (anchor) genes.



**Figure S15.** Intersections of subnetworks for single-trait GWAS for Stomatal density and multi-trait GWAS for Early Wood vessel area, Leaf Area, and Stomatal Density.