

Supplementary Document 2

Statistical analyses

Doc S2 Table 1. Statistical analysis (ANOVA/Tukey-Kramer) of growth parameters of wild-type and mutant plants. Growth rates of leaf rosettes were determined for plants grown in soil under SL-conditions ($\text{PDF} = 150 \mu\text{mol}\cdot\text{m}^{-2}\cdot\text{s}^{-1}$) in the long-day (Figure 2C, main article). Root growth was assessed for plants grown on 1/2MS agar (Figure 2D, main article). For the generative growth parameters plants were grown as described in Table 1 (main article). The biotypes are denoted, **a** = Ws-2, **b** = *xpt-1*, **c** = Col-0, **d** = *tpt-2*, **e** = *tpt-2/xpt-1*, **f** = amiRNA:*XPT tpt-2* #3, and **g** = amiRNA:*XPT tpt-2* #4. DAS = days after sowing. The significance levels of $P < 0.05$ or $P < 0.01$ are indicated by light or dark blue colors.

7 comparisons

Doc S2 Table 2. Statistical analysis (ANOVA/Tukey-Kramer) of Chl a fluorescence parameters of wild-type and mutant plants impaired in XPT and TPT as well as starch biosynthesis. The growth conditions and experimental setup is described in Supplementary Table 2. The biotypes are denoted, **a** = Ws-2, **b** = Col-0, **c** = *tpt-2*, **d** = *tpt-2/xpt-1*, **e** = amiRNA:XPT *tpt-2* #3, **f** = amiRNA:XPT *tpt-2* #4, **g** = *adg1-1/tpt-2*, **h** = *xpt-1*, **i** = *tpt-1*, and **j** = *tpt-1/xpt-1*. The significance levels of P < 0.05 or P < 0.01 are indicated by light or dark blue colors.

7 Comparisons

PS Parameters	<i>b vs a</i>	<i>c vs a</i>	<i>d vs a</i>	<i>e vs a</i>	<i>f vs a</i>	<i>g vs a</i>	<i>c vs b</i>	<i>d vs b</i>	<i>e vs b</i>	<i>f vs b</i>	<i>g vs b</i>	<i>d vs c</i>	<i>e vs c</i>	<i>f vs c</i>	<i>g vs c</i>	<i>e vs d</i>	<i>f vs d</i>	<i>g vs d</i>	<i>f vs e</i>	<i>g vs e</i>	<i>f vs g</i>
Long-day HL																					
F_v/F_m																					
Φ_{PSII}																					
Relative ETR ₍₇₀₀₎																					

4 Comparisons

<i>h vs a</i>	<i>i vs a</i>	<i>j vs a</i>	<i>i vs h</i>	<i>j vs h</i>	<i>i vs j</i>

Doc S2 Table 3. Statistical analysis (ANOVA/Tukey-Kramer) of rosette leaf parameters, i.e. photosynthetic pigments and protein contents as well as specific fresh weights. Plants were grown under HL- (PFD = 300 $\mu\text{mol}\cdot\text{m}^{-2}\cdot\text{s}^{-1}$; Tabel 3, main article) or LL-conditions (PFD = 30 $\mu\text{mol}\cdot\text{m}^{-2}\cdot\text{s}^{-1}$; Supplementary Table 3). The biotypes are denoted, **a** = Ws-2, **b** = Col-0, **c** = *tpt-2*, **d** = *tpt-2/xpt-1*, **e** = amiRNA:*XPT tpt-2 #3*, **f** = amiRNA:*XPT tpt-2 #4*, **g** = *adg1-1/tpt-2*, **h** = *xpt-1*, **i** = *tpt-1*, and **j** = *tpt-1/xpt-1*. The significance levels of P < 0.05 or P < 0.01 are indicated by light or dark blue colors.

7 Comparisons

4 Comparisons

Doc S2 Table 4. Statistical analysis (ANOVA/Tukey-Kramer) of carbohydrate contents in rosette leaves of wild-type and mutant plants. The plants were grown in soil under HL-conditions (PFD = 300 $\mu\text{mol}\cdot\text{m}^{-2}\cdot\text{s}^{-1}$) in the long-day (Figure 4, main article). The biotypes are denoted, **a** = Ws-2, **b** = xpt-1, **c** = tpt-1, **d** = tpt-1/xpt-1, **e** = Col-0, **f** = tpt-2, **g** = tpt-2/xpt-1, and **h** = adg1-1/tpt-2. The significance levels of P < 0.05 or P < 0.01 are indicated by light or dark blue colors.

8 Comparisons

Doc S2 Table4 (continued)

Time course	<i>b vs a</i>	<i>c vs a</i>	<i>d vs a</i>	<i>e vs a</i>	<i>f vs a</i>	<i>g vs a</i>	<i>h vs a</i>	<i>c vs b</i>	<i>d vs b</i>	<i>e vs b</i>	<i>f vs b</i>	<i>g vs b</i>	<i>h vs b</i>	<i>d vs c</i>	<i>e vs c</i>	<i>f vs c</i>	<i>g vs c</i>	<i>h vs c</i>	<i>e vs d</i>	<i>f vs d</i>	<i>g vs d</i>	<i>h vs d</i>	<i>f vs e</i>	<i>g vs e</i>	<i>h vs e</i>	<i>g vs f</i>	<i>h vs f</i>	<i>h vs g</i>
C Glc																												
1 h																												
4 h																												
8 h																												
12 h																												
15 h																												
17 h																												
20 h																												
23 h																												
D Fru																												
1 h																												
4 h																												
8 h																												
12 h																												
15 h																												
17 h																												
20 h																												
23 h																												

a = Ws-2, **b** = xpt-1, **c** = tpt-1, **d** = tpt-1/xpt-1, **e** = Col-0, **f** = tpt-2, **g** = tpt-2/xpt-1, and **h** = adg1-1/tpt-2

Doc S2 Table4 (continued)

Time course	<i>b</i> vs <i>a</i>	<i>c</i> vs <i>a</i>	<i>d</i> vs <i>a</i>	<i>e</i> vs <i>a</i>	<i>f</i> vs <i>a</i>	<i>g</i> vs <i>a</i>	<i>h</i> vs <i>a</i>	<i>c</i> vs <i>b</i>	<i>d</i> vs <i>b</i>	<i>e</i> vs <i>b</i>	<i>f</i> vs <i>b</i>	<i>g</i> vs <i>b</i>	<i>h</i> vs <i>b</i>	<i>d</i> vs <i>c</i>	<i>e</i> vs <i>c</i>	<i>f</i> vs <i>c</i>	<i>g</i> vs <i>c</i>	<i>h</i> vs <i>c</i>	<i>e</i> vs <i>d</i>	<i>f</i> vs <i>d</i>	<i>g</i> vs <i>d</i>	<i>h</i> vs <i>d</i>	<i>f</i> vs <i>e</i>	<i>g</i> vs <i>e</i>	<i>h</i> vs <i>e</i>	<i>g</i> vs <i>f</i>	<i>h</i> vs <i>f</i>	<i>h</i> vs <i>g</i>
E Suc																												
1 h																												
4 h																												
8 h																												
12 h																												
15 h																												
17 h																												
20 h																												
23 h																												

a = Ws-2, *b* = xpt-1, *c* = tpt-1, *d* = tpt-1/xpt-1, *e* = Col-0, *f* = tpt-2, *g* = tpt-2/xpt-1, and *h* = adg1-1/tpt-2

Doc S2 Table 5. Statistical analysis (ANOVA/Tukey-Kramer) of metabolite contents in leaves of wild-type and mutant plants. The plants were grown in soil under HL-conditions (PFD = 300 $\mu\text{mol}\cdot\text{m}^{-2}\cdot\text{s}^{-1}$) in the long-day (see Figure 5, and Supplementary Table 5). Samples were taken at the middle of the dark period (**A**), or the beginning (**B**), and middle of the light period (**C**). The biotypes are denoted, **a** = Ws-2, **b** = *xpt-1*, **c** = *tpt-1/xpt-1*, **d** = Col-0, **e** = *tpt-2*, and **f** = *tpt-2/xpt-1*. The significance levels of P < 0.05 or P < 0.01 are indicated by light or dark blue colors.

6 Comparisons

A Middle of the dark period (5 h in the dark)

Metabolites	<i>b vs a</i>	<i>c vs a</i>	<i>d vs a</i>	<i>e vs a</i>	<i>f vs a</i>	<i>c vs b</i>	<i>d vs b</i>	<i>e vs b</i>	<i>f vs b</i>	<i>d vs c</i>	<i>e vs c</i>	<i>f vs c</i>	<i>e vs d</i>	<i>f vs d</i>	<i>f vs e</i>
3-PGA				■			■		■		■		■		
DHAP															
S-7-P		■				■									
Ri5P															
Xu5P+Ru5P	■				■	■			■	■	■				■
Fru6P															
Glc6P															
Glc1P															
UDPG															
Glycerol-3-P															
ATP															
AMP															
Asn						■									
Phe					■				■			■			
Tyr									■						
Trp													■		

B Beginning of the light period (1 h in the light)

Doc S2 Table 5 (continued)

C Middle of the light period (8 h in the light)

	<i>b vs a</i>	<i>c vs a</i>	<i>d vs a</i>	<i>e vs a</i>	<i>f vs a</i>	<i>c vs b</i>	<i>d vs b</i>	<i>e vs b</i>	<i>f vs b</i>	<i>d vs c</i>	<i>e vs c</i>	<i>f vs c</i>	<i>e vs d</i>	<i>f vs d</i>	<i>e vs f</i>
3-PGA															
DHAP															
S-7-P															
Ri5P															
Xu5P+Ru5P		■			■	■				■	■				■
Fru6P															
Glc6P															
Glc1P															
UDPG															
Glycerol-3-P											■				
ATP															
AMP															
Asn															
Phe	■	■									■				
Tyr				■				■			■		■		■
Trp															

a = Ws-2, **b** = xpt-1, **c** = tpt-1/xpt-1, **d** = Col-0, **e** = tpt-2, and **f** = tpt-2/xpt-1.

Doc S2 Table 6 Statistical analysis (ANOVA/Tukey-Kramer) of metabolite contents in leaves of wild-type and mutant plants. The plants were grown in soil under HL-conditions (PFD = 300 $\mu\text{mol}\cdot\text{m}^{-2}\cdot\text{s}^{-1}$) in the long day (see Figure 5, and Supplementary Table 5). Samples were taken at the middle of the dark period (**A**), or the beginning (**B**) and middle of the light period (**C**). The biotypes are denoted, **a** = Ws-2, **b** = *xpt-1*, **c** = *tpt-1/xpt-1*, **d** = Col-0, **e** = *tpt-2*, and **f** = *tpt-2/xpt-1*. The significance levels of P < 0.05 or P < 0.01 are indicated by light or dark blue colors.

A Middle of the dark period (5 h in the dark)

6 Comparisons

Doc S2 Table 6 (continued)

B Beginning of the light period (1 h in the light)

Metabolite	b vs a	c vs a	d vs a	e vs a	f vs a	c vs b	d vs b	e vs b	f vs b	d vs c	e vs c	f vs c	e vs d	f vs d	e vs f
Glu															
Asp															
Ser															
Gly															
Thr															
Pro															
Phe															
Tyr															
Val															
Leu															
Ile															
Lys															
Met															
β -Ala															
Glycerol															
Mannitol															
Myoinositol															
Sorbitol															
Putrescine															
(Iso) Citrate															
2-Oxoglutarate															
Succinate															
Fumarate															
Malate															
Glycolate															
Lactate															
Glycerate															
Gluconate															
Maleate															
Malonate															
Shikimate															
Quinate															
Glc															
Fru															
Suc															
Maltose															
Mannose															
Xylose															

a = Ws-2, **b** = xpt-1, **c** = tpt-1/xpt-1, **d** = Col-0, **e** = tpt-2, and **f** = tpt-2/xpt-1.

Doc S2 Table 6 (continued)

C Middle of the light period (8 h in the light)

Metabolites	b vs a	c vs a	d vs a	e vs a	f vs a	c vs b	d vs b	e vs b	f vs b	d vs c	e vs c	f vs c	e vs d	f vs d	e vs f
Glu															
Asp															
Ser															
Gly															
Thr															
Pro															
Phe															
Tyr															
Val															
Leu															
Ile															
Lys															
Met															
β -Ala															
Glycerol															
Mannitol															
Myoinositol															
Sorbitol															
Putrescine															
(Iso) Citrate															
2-Oxoglutarate															
Succinate															
Fumarate															
Malate															
Glycolate															
Lactate															
Glycerate															
Gluconate															
Maleate															
Malonate															
Shikimate															
Quinate															
Glc															
Fru															
Suc															
Maltose															
Mannose															
Xylose															

a = Ws-2, b = xpt-1, c = tpt-1/xpt-1, d = Col-0, e = tpt-2, and f = tpt-2/xpt-1.

Doc S2 Table 7. Statistical analysis (ANOVA/Tukey-Kramer) of soluble amino acid contents in leaves of wild-type and mutant plants.

The plants were grown in soil under HL-conditions ($\text{PFD} = 300 \mu\text{mol}\cdot\text{m}^{-2}\cdot\text{s}^{-1}$) in the long-day (Figure 6C, main article). Samples were taken either 4 h in the dark (**A, C**) or 8 h in the light (**B, D**). The data represents the mean \pm SE of $n = 4$ to 5 independent replicates. The biotypes are denoted, **a** = Ws-2, **b** = Col-0, **c** = *tpt-2*, **d** = *tpt-2/xpt-1*, **e** = amiRNA:*XPT tpt-2 #3*, **f** = amiRNA:*XPT tpt-2 #4*, **g** = *adg1-1/tpt-2*, **h** = *xpt-1*, **i** = *tpt-1*, and **j** = *tpt-1/xpt-1*. The significance levels of $P < 0.05$ or $P < 0.01$ are indicated by light or dark blue colors.

A 4 h dark (absolute)

7 Comparisons

	<i>b vs a</i>	<i>c vs a</i>	<i>d vs a</i>	<i>e vs a</i>	<i>f vs a</i>	<i>g vs a</i>	<i>c vs b</i>	<i>d vs b</i>	<i>e vs b</i>	<i>f vs b</i>	<i>g vs b</i>	<i>d vs c</i>	<i>e vs c</i>	<i>f vs c</i>	<i>g vs c</i>	<i>e vs d</i>	<i>f vs d</i>	<i>g vs d</i>	<i>f vs e</i>	<i>g vs e</i>	<i>f vs g</i>
Amino acids																					
Glu																					
Gln																					
Asp																					
Asn																					
Ala																					
Ser																					
Gly																					
Thr																					
His																					
Arg																					
Phe																					
Tyr																					
Trp																					
Val																					
Leu																					
Ile																					
Lys																					
Σ																					

	<i>h vs a</i>	<i>i vs a</i>	<i>j vs a</i>	<i>i vs h</i>	<i>j vs h</i>	<i>i vs j</i>
h vs a						
i vs a						
j vs a						
i vs h						
j vs h						
i vs j						

Doc S2 Table 7. (continued)

B 8 h light (absolute)

	<i>b vs a</i>	<i>c vs a</i>	<i>d vs a</i>	<i>e vs a</i>	<i>f vs a</i>	<i>g vs a</i>	<i>c vs b</i>	<i>d vs b</i>	<i>e vs b</i>	<i>f vs b</i>	<i>g vs b</i>	<i>d vs c</i>	<i>e vs c</i>	<i>f vs c</i>	<i>g vs c</i>	<i>e vs d</i>	<i>f vs d</i>	<i>g vs d</i>	<i>f vs e</i>	<i>g vs e</i>	<i>f vs g</i>
Amino acids																					
Glu																					
Gln																					
Asp																					
Asn																					
Ala																					
Ser																					
Gly																					
Thr																					
His																					
Arg																					
Phe																					
Tyr																					
Trp																					
Val																					
Leu																					
Ile																					
Lys																					
Σ																					

a = Ws-2, **b** = Col-0, **c** = *tpt-2*, **d** = *tpt-2/xpt-1*, **e** = amiRNA:*XPT tpt-2 #3*, **f** = amiRNA:*XPT tpt-2 #4*, **g** = *adg1-1/tpt-2*, **h** = *xpt-1*, **i** = *tpt-1*, and **j** = *tpt-1/xpt-1*.

Doc S2 Table 7. (continued)

C 4 h dark (relative)

	<i>b vs a</i>	<i>c vs a</i>	<i>d vs a</i>	<i>e vs a</i>	<i>f vs a</i>	<i>g vs a</i>	<i>c vs b</i>	<i>d vs b</i>	<i>e vs b</i>	<i>f vs b</i>	<i>g vs b</i>	<i>d vs c</i>	<i>e vs c</i>	<i>f vs c</i>	<i>g vs c</i>	<i>e vs d</i>	<i>f vs d</i>	<i>g vs d</i>	<i>f vs e</i>	<i>g vs e</i>	<i>f vs g</i>
Amino acids																					
Glu																					
Gln																					
Asp																					
Asn																					
Ala																					
Ser																					
Gly																					
Thr																					
His																					
Arg																					
Phe																					
Tyr																					
Trp																					
Val																					
Leu																					
Ile																					
Lys																					

a = Ws-2, **b** = Col-0, **c** = *tpt-2*, **d** = *tpt-2/xpt-1*, **e** = amiRNA:XPT *tpt-2* #3, **f** = amiRNA:XPT *tpt-2* #4, **g** = *adg1-1/tpt-2*, **h** = *xpt-1*, **i** = *tpt-1*, and **j** = *tpt-1/xpt-1*.

Doc S2 Table 7. (continued)

D 8 h light (relative)

	b vs a	c vs a	d vs a	e vs a	f vs a	g vs a	c vs b	d vs b	e vs b	f vs b	g vs b	d vs c	e vs c	f vs c	g vs c	e vs d	f vs d	g vs d	f vs e	g vs e	f vs g	
Amino acids																						
Glu																						
Gln																						
Asp																						
Asn																						
Ala																						
Ser																						
Gly																						
Thr																						
His																						
Arg																						
Phe																						
Tyr																						
Trp																						
Val																						
Leu																						
Ile																						
Lys																						

a = Ws-2, b = Col-0, c = tpt-2, d = tpt-2/xpt-1, e = amiRNA:XPT tpt-2 #3, f = amiRNA:XPT tpt-2 #4, g = adg1-1/tpt-2, h = xpt-1, i = tpt-1, and j = tpt-1/xpt-1.

Doc S2 Table 8. Statistical analysis (ANOVA/Tukey-Kramer) of elemental analyses (N, C) in rosette leaves as well as fw/dw-ratios of wild-type and mutant plants. The plants were grown in soil under HL-conditions (PFD = 300 $\mu\text{mol}\cdot\text{m}^{-2}\cdot\text{s}^{-1}$) in the long-day (Table 3, main article). The biotypes are denoted, **a** = Ws-2, **b** = xpt-1, **c** = tpt-1, **d** = tpt-1/xpt-1, **e** = Col-0, **f** = tpt-2, **g** = tpt-2/xpt-1, and **h** = adg1-1/tpt-2. The significance levels of P < 0.05 or P < 0.01 are indicated by light or dark blue colors.

8 Comparisons

Doc S2 Table 9. Statistical analysis (ANOVA/Tukey-Kramer) of feeding Suc, Gln or both on rosette growth of wild-type and mutant plants. Plants were grown under HL-conditions (i.e. a PFD of $300 \mu\text{mol}\cdot\text{m}^{-2}\cdot\text{s}^{-1}$) in the long-day on 1/2 Ms agar plates in the absence or presence of 50 mM Suc, 2 mM Gln, or both. under HL. (Figure 7, main article; Supplementary Figures 2 and 3). The biotypes are denoted, **a** = Ws-2, **b** = *tpt-1/xpt-1*, **c** = Col-0, **d** = *tpt-2/xpt-1*, **e** = *gpt2-1/tpt-2/xpt-1*, and **f** = *adg1-1/tpt-2*. DAS = days after sowing. The significance levels of $P < 0.05$ or $P < 0.01$ are indicated by light or dark blue colors.

6 Comparisons

Time course rosette growth	<i>b vs a</i>	<i>c vs a</i>	<i>d vs a</i>	<i>e vs a</i>	<i>f vs a</i>	<i>c vs b</i>	<i>d vs b</i>	<i>e vs b</i>	<i>f vs b</i>	<i>d vs c</i>	<i>e vs c</i>	<i>f vs c</i>	<i>e vs d</i>	<i>f vs d</i>	<i>f vs e</i>	Time course rosette growth	<i>b vs a</i>	<i>c vs a</i>	<i>d vs a</i>	<i>e vs a</i>	<i>f vs a</i>	<i>c vs b</i>	<i>d vs b</i>	<i>e vs b</i>	<i>f vs b</i>	<i>d vs c</i>	<i>e vs c</i>	<i>f vs c</i>	<i>e vs d</i>	<i>f vs d</i>	<i>f vs e</i>		
½ MS																	+Suc																
11 DAS																11 DAS																	
15 DAS																15 DAS																	
19 DAS																19 DAS																	
22 DAS																22 DAS																	
+Gln																	+Suc + Gln																
11 DAS																11 DAS																	
15 DAS																15 DAS																	
19 DAS																19 DAS																	
22 DAS																22 DAS																	

6 Comparisons

Doc S2 Table 10. Statistical analysis (ANOVA/Tukey-Kramer) of feeding Suc, Gln or both on rosette growth of wild-type and mutant plants. Plants were grown under HL-conditions (i.e. a PFD of $300 \mu\text{mol}\cdot\text{m}^{-2}\cdot\text{s}^{-1}$) in the long-day on 1/2 Ms agar plates in the absence or presence of 50 mM Suc, 2 mM Gln, or both. under HL. (Figure 7, main article; Supplementary Figures 2 and 3). The conditions are denoted, **a** = 1/2 MS, **b** = 50 mM Suc, **c** = 2 mM Gln, and **d** = 50 mM Suc and 2 mM Gln. DAS = days after sowing. The significance levels of $P < 0.05$ or $P < 0.01$ are indicated by light or dark blue colors.

4 Comparisons		4 Comparisons		4 Comparisons																	
Time of growth	<i>b vs a</i>	<i>c vs a</i>	<i>d vs a</i>	<i>c vs b</i>	<i>d vs b</i>	<i>d vs c</i>	Time of growth	<i>b vs a</i>	<i>c vs a</i>	<i>d vs a</i>	<i>c vs b</i>	<i>d vs b</i>	<i>d vs c</i>	Time of growth	<i>b vs a</i>	<i>c vs a</i>	<i>d vs a</i>	<i>c vs b</i>	<i>d vs b</i>	<i>d vs c</i>	
Ws-2						tpt-1/xpt-1						Col-0									
11 DAS							11 DAS							11 DAS							
15 DAS							15 DAS							15 DAS							
19 DAS							19 DAS							19 DAS							
22 DAS							22 DAS							22 DAS							
tpt-2/xpt-1						gpt2-1/tpt-2/xpt-1						adg1-1/tpt-2									
11 DAS							11 DAS							11 DAS							
15 DAS							15 DAS							15 DAS							
19 DAS							19 DAS							19 DAS							
22 DAS							22 DAS							22 DAS							

Doc S2 Table 11. Statistical analysis (ANOVA/Tukey-Kramer) of feeding Suc, Gln or both on modulated Chl a fluorescence parameters in wild-type and mutant plants. Plants were grown under HL-conditions (i.e. a PFD of $300 \mu\text{mol}\cdot\text{m}^{-2}\cdot\text{s}^{-1}$) in the long-day on 1/2 Ms agar plates in the absence or presence of 50 mM Suc, 2 mM Gln, or both. (Figure 7, main article; Supplementary Figure 4). The biotypes are denoted, **a** = Ws-2, **b** = *tpt-1/xpt-1*, **c** = Col-0, **d** = *tpt-2/xpt-1*, **e** = *gpt2-1/tpt-2/xpt-1*, and **f** = *adg1-1/tpt-2*. The significance levels of $P < 0.05$ or $P < 0.01$ are indicated by light or dark blue colors.

6 Comparisons

6 Comparisons

Doc S2 Table 12. Statistical analysis (ANOVA/Tukey-Kramer) of feeding Suc, Gln or both on modulated Chl a fluorescence parameters in wild-type and mutant plants. Plants were grown under HL-conditions (i.e. a PFD of $300 \mu\text{mol}\cdot\text{m}^{-2}\cdot\text{s}^{-1}$) in the long-day on 1/2 Ms agar plates in the absence or presence of 50 mM Suc, 2 mM Gln, or both. (Figure 7, main article; Supplementary Figure 4). The conditions are denoted, **a** = 1/2 MS, **b** = 50 mM Suc, **c** = 2 mM Gln, and **d** = 50 mM Suc and 2 mM Gln. The significance levels of $P < 0.05$ or $P < 0.01$ are indicated by light or dark blue colors.

4 Comparisons						4 Comparisons						4 Comparisons						
PS parameters	<i>b vs a</i>	<i>c vs a</i>	<i>d vs a</i>	<i>c vs b</i>	<i>d vs b</i>	PS parameters	<i>b vs a</i>	<i>c vs a</i>	<i>d vs a</i>	<i>c vs b</i>	<i>d vs b</i>	PS parameters	<i>b vs a</i>	<i>c vs a</i>	<i>d vs a</i>	<i>c vs b</i>	<i>d vs b</i>	<i>d vs c</i>
Ws-2						tpt-1/xpt-1						Col-0						
F_v/F_m	Dark Blue	Dark Blue	Dark Blue	Dark Blue	White	Dark Blue	Dark Blue	White	Dark Blue	White	Dark Blue	Dark Blue	Light Blue	White	Dark Blue	Light Blue	Dark Blue	
Φ_{PSII}	Light Blue	Dark Blue	Dark Blue	Dark Blue	White	Dark Blue	Light Blue	White	Dark Blue	White	Dark Blue	Light Blue	Light Blue	White	Dark Blue	Dark Blue	White	
tpt-2/xpt-1						gpt2-1/tpt-2/xpt-1						adg1-1/tpt-2						
F_v/F_m	Dark Blue	Dark Blue	Dark Blue	Dark Blue	White	Dark Blue	Dark Blue	White	Dark Blue	White	Dark Blue	Dark Blue	Light Blue	White	Dark Blue	Dark Blue	Dark Blue	
Φ_{PSII}	Dark Blue	White	Dark Blue	Dark Blue	White	Dark Blue	White	White	Dark Blue	Light Blue	Dark Blue	Light Blue	Light Blue	White	Dark Blue	Dark Blue	Dark Blue	

Doc S2 Table 13. Statistical analysis (ANOVA/Tukey-Kramer) of spectroscopic analysis of thylakoid membranes isolated from wild-type and mutants plants. Plants were grown in soil under HL-conditions (PFD = 300 $\mu\text{mol}\cdot\text{m}^{-2}\cdot\text{s}^{-1}$) in the long-day. Functional components of PSII, PSI and the Cyt b₆/f complex were determined by spectroscopic methods using thylakoids isolated from the plants and were either referred to a leaf area- (Figure 8B, main article) or Chl basis (Supplementary Figure 5). The biotypes are denoted, **a** = Col-0, **b** = *tpt-2*, **c** = *tpt-2/xpt-1*, **d** = amiRNA:*XPT tpt-2 #4*, **e** = Ws-2, **f** = *xpt-1*, **g** = *tpt-1*, and **h** = *tpt-1/xpt-1*. The significance levels of P < 0.05 or P < 0.01 are indicated by light or dark blue colors.

8 Comparisons