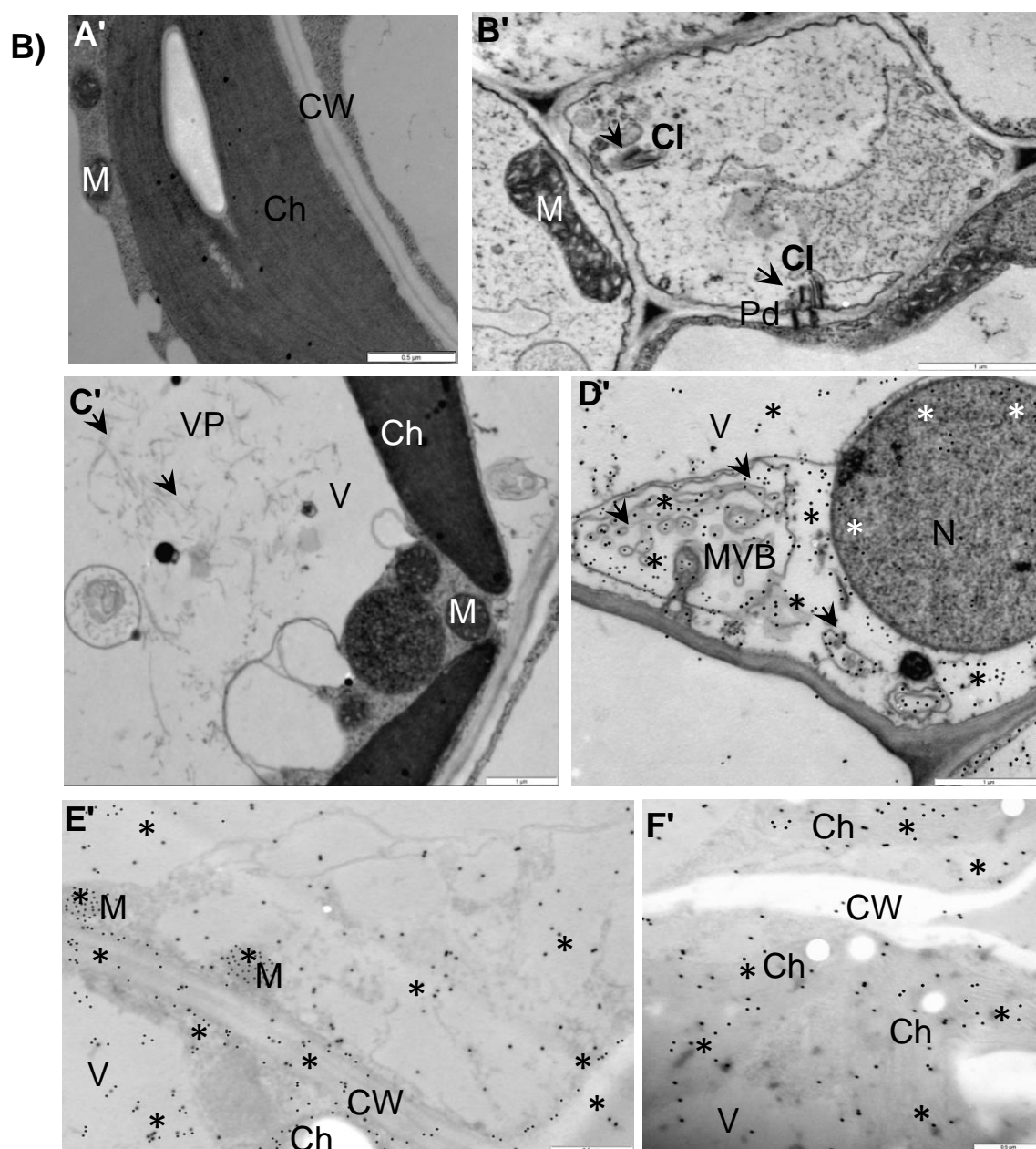
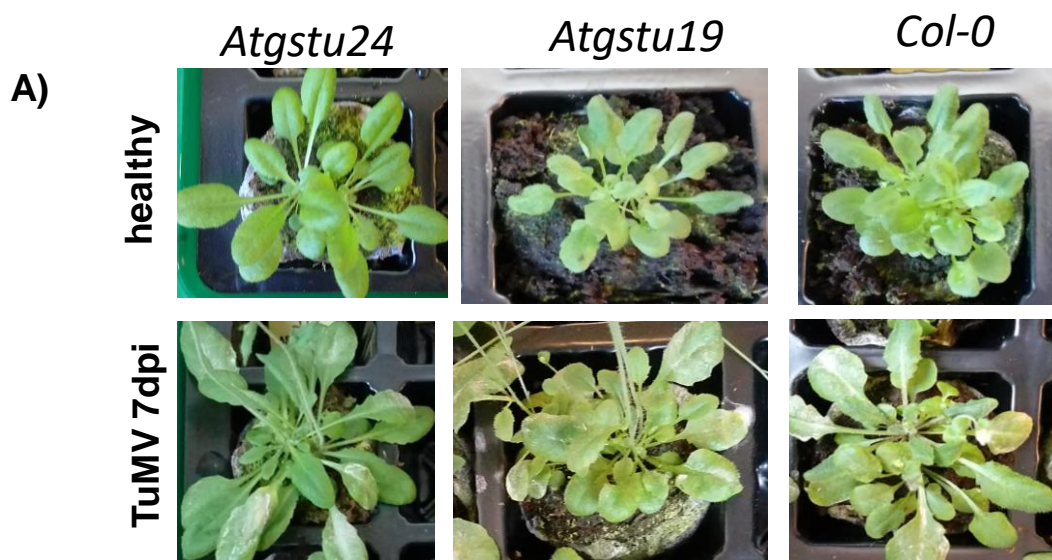


**Table S1. TuMV detection assessment using DAS-ELISA on *Arabidopsis thaliana* wild type and mutant leaves at 3 and 7 dpi. Values presented are mean optical density (OD<sub>405nm</sub>) values. Absence of virus is marked by (-). Presence of TuMV (+) is confirmed in samples with mean OD<sub>405nm</sub> above estimated cut off point: 0.1412.**

Sample	Mean OD <sub>450nm</sub>	Presence (+)/absence of the virus (-)
buffer	0.0001	-
mock-inoculated Col-0 (3dpi)	0.0305	-
mock-inoculated <i>Atgstu19</i> (3dpi)	0.0213	-
mock-inoculated <i>Atgstu24</i> (3dpi)	0.0200	-
TuMV-inoculated Col-0 (3dpi)	0.8820	+
TuMV-inoculated <i>Atgstu19</i> (3dpi)	0.7519	+
TuMV-inoculated <i>Atgstu 24</i> (3dpi)	0.5200	+
mock-inoculated Col-0 (7dpi)	0.0501	-
mock-inoculated <i>Atgstu19</i> (7dpi)	0.0521	-
mock-inoculated <i>Atgstu 24</i> (7dpi)	0.0402	-
TuMV-inoculated Col-0 (7dpi)	1.425	+
TuMV-inoculated <i>Atgstu19</i> (7dpi)	3.321	+
TuMV-inoculated <i>Atgstu24</i> (7dpi)	0.693	+
mock-inoculated Col-0 (14dpi)	0.0600	-
mock-inoculated <i>Atgstu19</i> (14dpi)	0.0632	-
mock-inoculated <i>Atgstu 24</i> (14dpi)	0.0606	-
TuMV-inoculated Col-0 (14dpi)	1.939	+
TuMV-inoculated <i>Atgstu19</i> (14dpi)	4.026	+
TuMV-inoculated <i>Atgstu24</i> (14dpi)	0.299	+



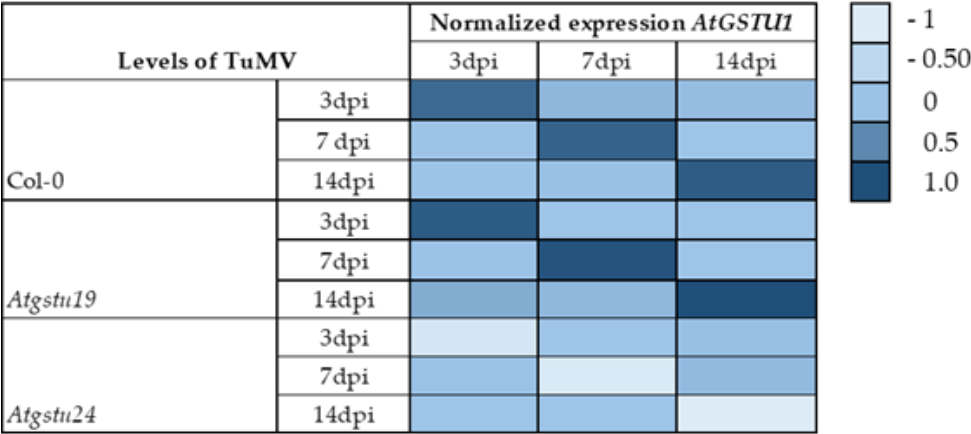
Supplementary Figure S1

### Supplementary Figure S1

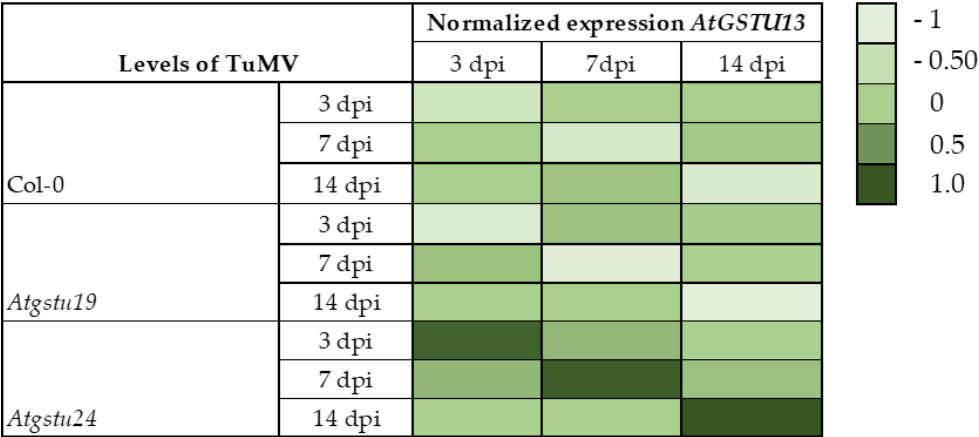
**Panel A)** - *Arabidopsis thaliana Atgstu24, Atgstu19* knockout mutant plants and Col-0 wild type plants: healthy plants, and 7 dpi after TuMV inoculation;

**Panel B)** – Ultrastructural changes in Col-0 wild type inoculated leaves: **(A')** Unchanged mesophyll cell from Col-0 mock-inoculated leaf. CW-cell wall, Ch-chloroplast, M-mitochondria. Scale bar 0.5µm; **(B')** TuMV cytoplasmic inclusions (CI, arrows) in phloem parenchyma cells 7 days after TuMV inoculation. M-mitochondria, Pd-plasmodesmata. Scale bar 1µm. **(C')** Virus particles (VP, arrows) in vacuole (V) of mesophyll cell. Ch-chloroplast, M-mitochondria. Scale bar 1µm. **(D')** Gold particles indicating TuMV (\*) in multivesicular bodies (MVB, arrows) and in nucleus (N) in mesophyll cell 7 dpi after TuMV inoculation. Scale bar 1µm. **(E')** Gold granules indicating GSH (\*) deposition along cell wall (CW), in mitochondria (M) and chloroplasts (Ch) in phloem Col-0 7dpi TuMV-inoculated leaf. Scale bar 0.5 µm. **(F')** GSH (\*) deposition in changed chloroplasts (Ch) in Col-0 mesophyll cell 14 days after TuMV inoculation. Scale bar 0.5 µm.

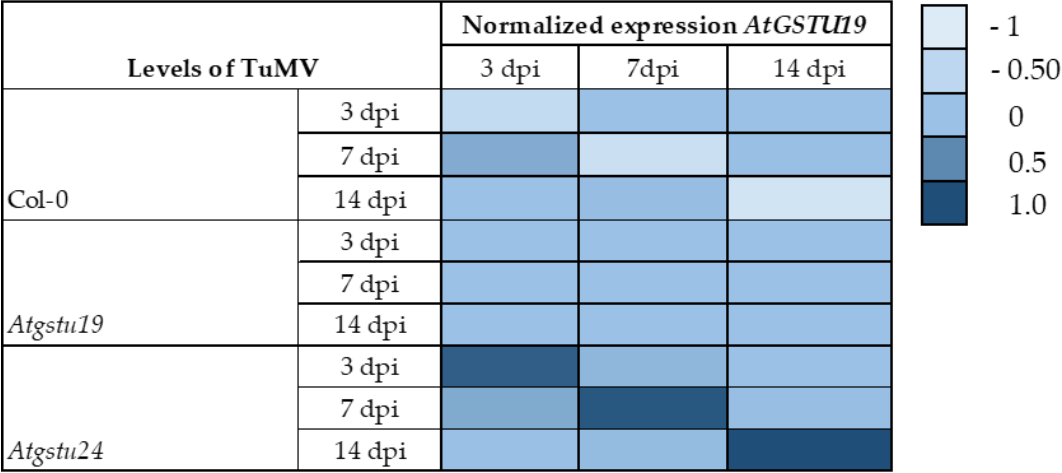
**Table S2.** Heatmap of PCC for *AtGSTU1* normalized expression (based on *AtEf1α* and *AtF-Box*) and *TuMV* levels in virus-inoculated **Col-0**, *Atgstu19* and *Atgstu24* plants from 3 to 14 dpi. PCC matrix values are presented pairwise for specific cell compartments in specific time dpi and marked with colors, from very dark blue (PCC = 1) to bright blue (PCC = -1).



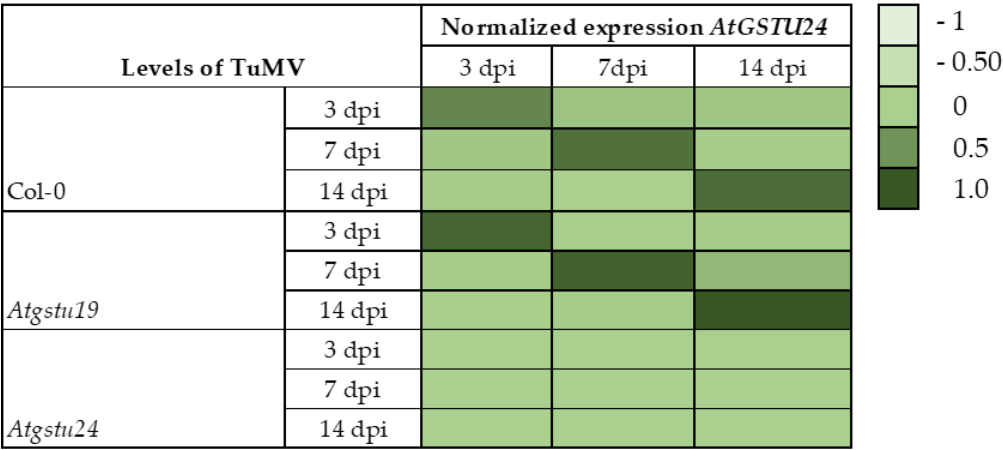
**Table S3.** Heatmap of PCC for *AtGSTU13* normalized expression (based on *AtEf1α* and *AtF-Box*) and *TuMV* levels in virus-inoculated **Col-0**, *Atgstu19* and *Atgstu24* plants from 3 to 14 dpi. PCC matrix values are presented pairwise for specific cell compartments in specific time dpi and marked with colors, from very dark blue (PCC = 1) to bright blue (PCC = -1).



**Table S4.** Heatmap of PCC for *AtGSTU19* normalized expression (based on *AtEf1α* and *AtF-Box*) and *TuMV* levels in virus-inoculated **Col-0**, *Atgstu19* and *Atgstu24* plants from **3 to 14 dpi**. PCC matrix values are presented pairwise for specific cell compartments in specific time dpi and marked with colors, from very dark blue (PCC = 1) to bright blue (PCC = -1).



**Table S5.** Heatmap of PCC for *AtGSTU24* normalized expression (based on *AtEf1α* and *AtF-Box*) and *TuMV* levels in virus-inoculated **Col-0**, *Atgstu19* and *Atgstu24* plants from **3 to 14 dpi**. PCC matrix values are presented pairwise for specific cell compartments in specific time dpi and marked with colors, from very dark blue (PCC = 1) to bright blue (PCC = -1).



**Table S6.** Primer sequences for RT-qPCR analyses

Genes	Forward Primer	Reverse Primer	Concentration in Reaction (μM)
<i>Investigated</i>			
<i>TuMV-CP</i>	5'- CCGGAATTC ATGRTITGGT GYATIGAIAA YGG-3'	5'- CGCGGATCCGCIG YYTTCATYTGIRIIW KIGC-3'	0.5
<i>AtGSTU1</i>	5'- GCAGTGAGG GGATGTATTC -3'	5'- TTTCGTAGGCAAG AAGTATCT-3'	0.5
<i>AtGSTU13</i>	5'- CGCAAAGCA AAAGTTCAA TGT-3'	5'- TGGCACAAAACA CAGACAAAT-3'	0.5
<i>AtGSTU19</i>	5'- ATGATGCTC AGAGGAAGG TG-3'	5'- ATAGCCAAAGTC ATCGCCAC-3'	0,5
<i>AtGSTU24</i>	5'- AAGGTGAGG AGCAAGAAG CA-3'	5'- ACATACCCAAAA GTTTCGTCTC-3'	0.5
<i>Reference</i>			
<i>AtEF1a</i>	5'- CACCACTGG AGGTTTTGAG G-3'	3'- TGGAGTATTGGG GGTGGT-5'	0.5
<i>AtF-Box</i>	5'- GCTTGCACA CGCCATATC AAT-3'	3'- TGGATTTTACCAC CTTCCGCA-5'	0.5

**Table S7.** Conditions of the RT-qPCR for the reference genes (\*).

Program	Parameters
Preliminary denaturation	95 °C for 5 min
Amplification (35 cycles)	95 °C for 10 s
	58 °C for 10 s
	72 °C for 20 s *
Melting curve	65–95 °C; 0.1 °C/s