

Supplementary Material

Table S1. Physicochemical characteristics of the Siro Royal universal substrate (SIRO[®], Portugal) used for the plant assay.

PHYSICOCHEMICAL CHARACTERISTICS	
pH (CaCl ₂)	5.5 – 6.5
Electric conductivity	50 – 100 µS cm ⁻¹
Granulometry	0 – 15 mm
Organic matter	> 70%
NPK	19 – 7 – 10

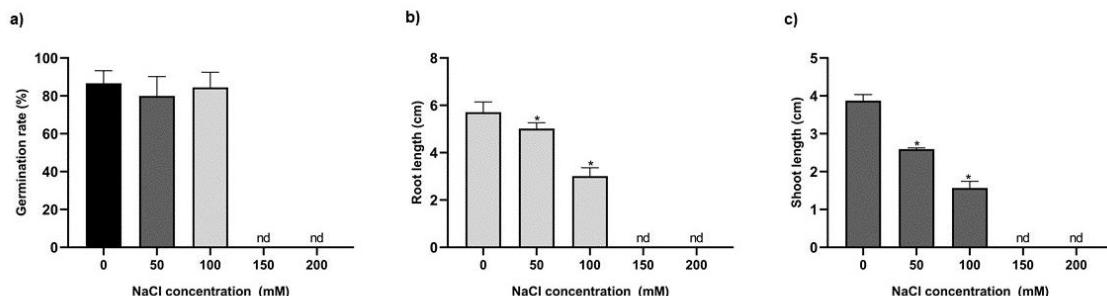


Figure S1. Percentage of germination (a), root length (b) and shoot length (c) of tomato seedlings grown for 7 days in solid MS nutritive medium, supplemented with different concentrations of NaCl (0, 50, 100, 150 and 200 mM). Data presented are mean ± SEM ($n \geq 3$). Asterisks above the error bars indicate significant statistical differences between treatments and the control at $p \leq 0.05$, assessed through Tukey post-hoc test, following a one-way ANOVA.

Table S2. Results of the two-way ANOVA for all evaluated parameters in roots of *Solanum lycopersicum* L. var. *cerasiforme* after 21 days of exposure to 42 °C (4 hours per day) and irrigation with (100 mM) or without NaCl. Parameters where significant differences ($p \leq 0.05$) were recorded are highlighted in bold.

PARAMETER (roots)	Factors		Interaction
	SALT	HEAT	
Length	F (1, 43) = 28.26; $p < 0.0001$	F (1, 43) = 52.66; $p < 0.0001$	F (1, 43) = 0.6012; $p = 0.4424$
Dry weight	F (1, 43) = 36.84; $p < 0.0001$	F (1, 43) = 40.38; $p < 0.0001$	F (1, 43) = 0.5333; $p = 0.4692$
Water content	F (1, 19) = 0.4653; $p = 0.4653$	F (1, 19) = 0.9359; $p = 0.3455$	F (1, 19) = 0.007945; $p = 0.9299$
Na ⁺	F (1, 8) = 656421; $p < 0.0001$	F (1, 8) = 2863; $p < 0.0001$	F (1, 8) = 1834; $p < 0.0001$
K ⁺	F (1, 8) = 19837; $p < 0.0001$	F (1, 8) = 691.1; $p < 0.0001$	F (1, 8) = 2244; $p < 0.0001$
Ca ²⁺	F (1, 8) = 3321; $p < 0.0001$	F (1, 8) = 55.13; $p < 0.0001$	F (1, 8) = 496.1; $p < 0.0001$
Mg ²⁺	F (1, 8) = 165.7; $p < 0.0001$	F (1, 8) = 414.0; $p < 0.0001$	F (1, 8) = 374.4; $p < 0.0001$
O ₂ ⁻	F (1, 8) = 2.350; $p = 0.1638$	F (1, 8) = 0.0004534; $p = 0.9835$	F (1, 8) = 1.930; $p = 0.2022$
H ₂ O ₂	F (1, 8) = 1.694; $p = 0.2293$	F (1, 8) = 61.16; $p < 0.0001$	F (1, 8) = 1.731; $p = 0.2248$
MDA	F (1, 11) = 5.810; $p = 0.0346$	F (1, 11) = 4.635; $p = 0.0544$	F (1, 11) = 0.2805; $p = 0.6069$

Proline	F (1, 9) = 830.0; p < 0.0001	F (1, 9) = 19.13; p = 0.018	F (1, 9) = 18.55; p = 0.0020
Total AsA	F (1, 10) = 0.02314; p = 0.8821	F (1, 10) = 2.693; p = 1318	F (1, 10) = 0.6917; p = 0.4250
AsA	F (1, 10) = 0.06414; p = 0.8052	F (1, 10) = 0.01829; p = 0.8951	F (1, 10) = 0.1727; p = 0.6865
DHA	F (1, 9) = 0.1271; p = 0.7296	F (1, 9) = 1.268; p = 0.2892	F (1, 9) = 0.3393; p = 0.5745
AsA/DHA	F (1, 9) = 0.7880; p = 0.3978	F (1, 9) = 1.967; p = 0.1944	F (1, 9) = 0.1392; p = 0.7177
GSH	F (1, 9) = 2.005; p = 0.1905	F (1, 9) = 18.68; p = 0.0019	F (1, 9) = 5.400; p = 0.0452
Total thiols	F (1, 8) = 3.618; p = 0.0937	F (1, 8) = 79.12; p < 0.0001	F (1, 8) = 1.373; p = 0.2749
Non-protein/ Protein thiols	F (1, 10) = 0.1765; p = 0.6833	F (1, 10) = 6.380; p = 0.0301	F (1, 10) = 1.509; p = 0.2474
SOD	F (1, 11) = 0.006482; p = 0.9373	F (1, 11) = 2.057; p = 0.1793	F (1, 11) = 4.219; p = 0.0645
CAT	F (1, 10) = 15.01; p = 0.0031	F (1, 10) = 9.137; p = 0.0128	F (1, 10) = 7.859; p = 0.0187
APX	F (1, 9) = 8.434; p = 0.0175	F (1, 9) = 43.25; p = 0.0001	F (1, 9) = 0.001389; p = 0.9711
DHAR	F (1, 9) = 7.860; p = 0.0206	F (1, 9) = 58.39; p < 0.0001	F (1, 9) = 24.81; p = 0.0008
GR	F (1, 9) = 2.326; p = 0.1616	F (1, 9) = 13.90; p = 0.0047	F (1, 9) = 34.27; p = 0.0002

Table S3. Results of the two-way ANOVA for all evaluated parameters in shoots of *Solanum lycopersicum* L. var. *cerasiforme* after 21 days of exposure to 42 °C (4 hours per day) and irrigation with (100 mM) or without NaCl. Parameters where significant differences ($p \leq 0.05$) were recorded are highlighted in bold.

PARAMETER (shoots)	Factors		Interaction
	SALT	HEAT	
Length	F (1, 36) = 107.0; p < 0.0001	F (1, 36) = 112.6; p < 0.0001	F (1, 36) = 0.8989; p = 0.3494
Dry weight	F (1, 42) = 59.36; p < 0.0001	F (1, 42) = 76.05; p < 0.0001	F (1, 42) = 1.909; p = 0.1743
Water content	F (1, 19) = 56.06; p < 0.0001	F (1, 19) = 6.082; p = 0.0233	F (1, 19) = 0.1518; p = 0.7011
Na ⁺	F (1, 8) = 1381045; p < 0.0001	F (1, 8) = 82240; p < 0.0001	F (1, 8) = 107573; p < 0.0001
K ⁺	F (1, 8) = 22713; p < 0.0001	F (1, 8) = 3027; p < 0.0001	F (1, 8) = 925.1; p < 0.0001
Ca ²⁺	F (1, 8) = 570.2; p < 0.0001	F (1, 8) = 44.18; p = 0.0002	F (1, 8) = 2424; p < 0.0001
Mg ²⁺	F (1, 8) = 19.00; p < 0.0024	F (1, 8) = 343.6; p < 0.0001	F (1, 8) = 104.0; p < 0.0001
O ₂ ⁻	F (1, 8) = 78.00; p < 0.0001	F (1, 8) = 16.65; p = 0.0035	F (1, 8) = 16.59; p = 0.0036
H ₂ O ₂	F (1, 9) = 42.32; p = 0.0001	F (1, 9) = 38.81; p = 0.0002	F (1, 9) = 26.58; p = 0.0006
MDA	F (1, 9) = 57.70; p < 0.0001	F (1, 9) = 47.30; p < 0.0001	F (1, 9) = 19.71; p = 0.0016
Proline	F (1, 9) = 408.4; p < 0.0001	F (1, 9) = 26.47; p = 0.0006	F (1, 9) = 22.06; p = 0.0011
Total AsA	F (1, 9) = 7.639; p = 0.0220	F (1, 9) = 3.930; p = 0.0788	F (1, 9) = 2.925; p = 0.1214
AsA	F (1, 9) = 0.5398; p = 0.4812	F (1, 9) = 6.613; p = 0.0301	F (1, 9) = 1.180; p = 0.3056
DHA	F (1, 9) = 19.01; p = 0.0018	F (1, 9) = 1.573; p = 0.2414	F (1, 9) = 4.470; p = 0.0636
AsA/DHA	F (1, 10) = 5.243; p = 0.0450	F (1, 10) = 11.76; p = 0.0064	F (1, 10) = 0.004423; p = 0.9483
GSH	F (1, 10) = 32.78; p = 0.0002	F (1, 10) = 0.002158; p = 0.9639	F (1, 10) = 20.78; p = 0.0010
Total thiols	F (1, 11) = 8.443; p = 0.0143	F (1, 11) = 1.035; p = 0.3309	F (1, 11) = 16.34; p = 0.0019
Non-protein/ Protein thiols	F (1, 10) = 52.82; p < 0.0001	F (1, 10) = 58.90; p < 0.0001	F (1, 10) = 34.17; p = 0.0002
SOD	F (1, 10) = 61.00; p < 0.0001	F (1, 10) = 9.535; p = 0.0115	F (1, 10) = 20.44; p = 0.0011
CAT	F (1, 11) = 16.66; p = 0.0018	F (1, 11) = 30.00; p = 0.0002	F (1, 11) = 1.695; p = 0.2196
APX	F (1, 8) = 6.117; p = 0.0385	F (1, 8) = 25.41; p = 0.0010	F (1, 8) = 15.42; p = 0.0044
DHAR	F (1, 10) = 40.70; p < 0.0001	F (1, 10) = 6.185; p = 0.0322	F (1, 10) = 21.95; p = 0.0009
GR	F (1, 11) = 58.88; p < 0.0001	F (1, 11) = 6.330; p = 0.0287	F (1, 11) = 1.646; p = 0.2259

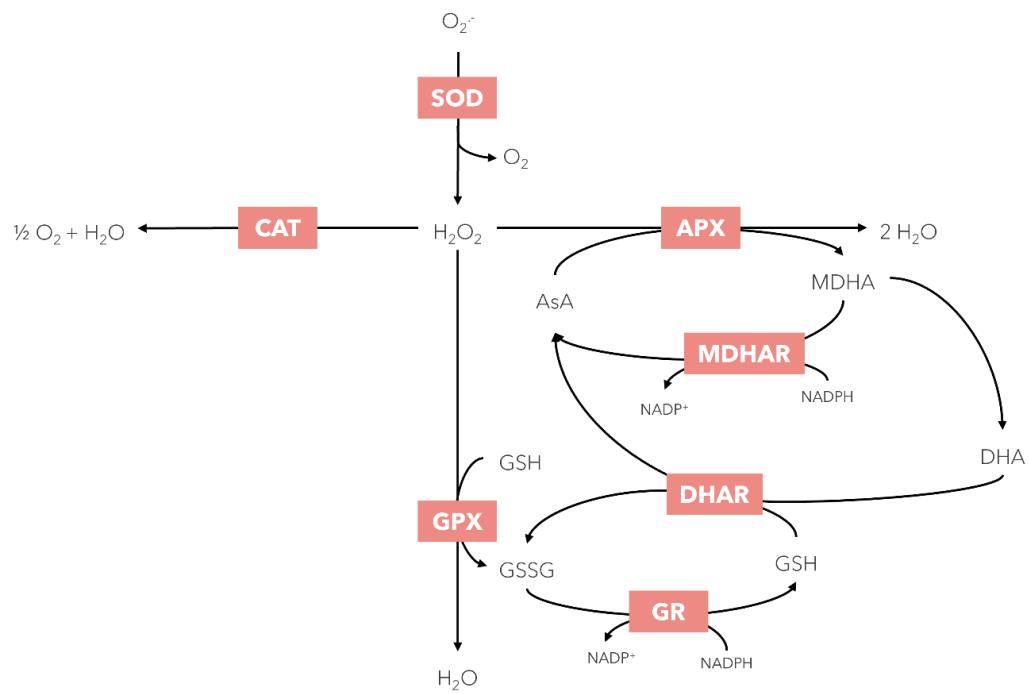


Figure S2. Enzymatic pathways that regulate redox status in plants. Superoxide anion (O_2^-), hydrogen peroxide (H_2O_2), superoxide dismutase (SOD), catalase (CAT), ascorbate peroxidase (APX), monodehydroascorbate reductase (MDHAR), dehydroascorbate reductase (DHAR), glutathione reductase (GR), glutathione peroxidases (GPX), ascorbate (AsA), monodehydroascorbate (MDHA), dehydroascorbate (DHA), oxidized glutathione (GSSG), reduced glutathione (GSH).