

Supplementary Material

Table S1. Detailed ANOVA results for all evaluated parameters in roots of *Solanum lycopersicum* L. cv. Micro-Tom grown for 28 days in OECD soil contaminated by GLY (10 mg kg⁻¹) and/or foliar treated with SNP (200 μ M). Parameters where significant differences ($p \leq 0.05$) were recorded are highlighted at bold.

Parameter	ANOVA
Root length	F (3, 10) = 19.11; $p < 0.01$
Fresh biomass	F (3, 9) = 64.36; $p < 0.01$
NR	F (3, 8) = 3.013; $p > 0.05$
Total protein	F (3, 10) = 19.21; $p < 0.01$
LP	F (3, 9) = 9.339; $p < 0.01$
O ₂ ⁻	F (3, 16) = 12.03; $p < 0.01$
H ₂ O ₂	F (3, 8) = 12.29; $p < 0.01$
Proline	F (3, 12) = 14.17; $p < 0.01$
GSH	F (3, 8) = 46.08; $p < 0.01$
Total AsA	F (3, 8) = 7.842; $p < 0.01$
AsA/DHA	F (3, 13) = 5.991; $p < 0.01$
TAC	F (3, 8) = 4.792; $p < 0.05$
TPC	F (3, 8) = 0.3788; $p > 0.05$
SOD	F (3, 9) = 10.36; $p < 0.01$
CAT	F (3, 9) = 13.45; $p < 0.01$
APX	F (3, 8) = 23.07; $p < 0.01$
GST	F (3, 9) = 50.51; $p < 0.01$

Table S2. Detailed ANOVA results for all evaluated parameters in shoots of *Solanum lycopersicum* L. cv. Micro-Tom grown for 28 days in OECD soil contaminated by GLY (10 mg kg⁻¹) and/or foliar treated with SNP (200 μ M). Parameters where significant differences ($p \leq 0.05$) were recorded are highlighted at bold.

Parameter	ANOVA
Fresh biomass	F (3, 10) = 13.74; $p < 0.01$
NR	F (3, 8) = 7.339; $p < 0.05$
Total protein	F (3, 27) = 3.902; $p < 0.05$
LP	F (3, 8) = 14.02; $p < 0.01$
O₂⁻	F (3, 17) = 20.23; $p < 0.01$
H ₂ O ₂	F (3, 10) = 1.127; $p > 0.05$
Proline	F (3, 9) = 28.02; $p < 0.01$
GSH	F (3, 15) = 14.83; $p < 0.01$
Total AsA	F (3, 9) = 7.387; $p < 0.01$
AsA/DHA	F (3, 9) = 5.532; $p < 0.05$
TAC	F (3, 8) = 1.445; $p > 0.05$
TPC	F (3, 5) = 49.63; $p < 0.01$
SOD	F (3, 9) = 2.874; $p > 0.05$
CAT	F (3, 7) = 27.41; $p < 0.01$
APX	F (3, 9) = 7.474; $p < 0.01$
GST	F (3, 9) = 6.360; $p < 0.05$

Table S3. Detailed ANOVA results for productivity-related parameters of *Solanum lycopersicum* L. cv. Micro-Tom grown for 28 days in OECD soil contaminated by GLY (10 mg kg⁻¹) and/or foliar treated with SNP (200 μ M). Parameters where significant differences ($p \leq 0.05$) were recorded are highlighted at bold.

Parameter	ANOVA
Number of flowers	F (3, 10) = 4.444; $p < 0.05$
Number of fruits	F (3, 14) = 4.370; $p < 0.05$
Fruit fresh biomass	F (3, 16) = 0.571; $p > 0.05$