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OUTGROWING Co-Dep Niceness (Part 3)

Accession	BIN Name	Description	P-C	P-HS	P-E-C	P-E-HS	Fold change P-HS vs. P-C
Solyc10g051390	RNA.RNA binding	RNA-binding glycine-rich protein-1b	2.8	0.1	2.3	0.6	-28.1
Solyc02g068740	PS.photorespiration.glycine cleavage.H protein	Glycine cleavage system H protein 1	1.6	0.1	0.7	0.4	-16.4
Solyc02g068450	transport.p- and v-ATPases.H+-transporting two-sector ATPase	Vacuolar ATPase F subunit	1.5	0.1	0.5	1.0	-14.8
Solyc10g078960	protein.synthesis.ribosomal protein.eukaryotic.60S subunit.L21	60S ribosomal protein L21-like protein	1.3	0.1	1.3	1.0	-12.8
Solyc09g090590	development.unspecified	Calcium binding protein caleosin	1.2	0.1	0.7	0.3	-12.0
Solyc09g091170	not assigned.unknown	NADH dehydrogenase 1 alpha subcomplex subunit 13	1.2	0.1	1.5	0.1	-11.7
Solyc07g005560	protein.synthesis.initiation	Eukaryotic translation initiation factor 5A	1.1	0.1	8.0	0.1	-10.6
Solyc06g069860	Protein.synthesis.ribosomal protein.eukaryotic.60S subunit.L34	60S ribosomal protein L34	1.1	0.1	0.1	0.5	-10.5
Solyc10g007700	signaling.G-proteins	Ras-related protein Rab-2-A	1.0	0.1	0.6	0.9	-9.9
Solyc02g070310	protein.synthesis.ribosomal protein.eukaryotic.60S subunit.L32	Ribosomal protein L32	0.9	0.1	0.5	0.5	-9.5
Solyc09g007180	nucleotide metabolism.phosphotransfer and pyrophosphatases.adenylate kinase	Adenylate kinase	0.7	0.1	0.1	0.1	-7.5
Solyc04g074550	Mitochondrial electron transport/ATP synthesis.cytochrome c oxidase	Cytochrome c oxidase subunit Vib	0.7	0.1	0.1	0.9	-6.8
Solyc05g005160	TCA/org. transformation.other organic acid transformaitons.atp-citrate lyase	ATP-citrate lyase A-2	0.6	0.1	0.8	0.6	-5.9
Solyc10g018060	not assigned.unknown	Elicitor-responsive protein 3	0.6	0.1	0.1	0.3	-5.7
Solyc09g061230	redox.ascorbate and glutathione	Cytochrome b5	2.5	0.4	0.9	2.1	-5.7

The complete dataset can be found in **Supplementary Table S6**. P-C, pollen derived from plants maintained at control conditions; P-HS, pollen derived from plants exposed to HS conditions; P-E-C, pollen derived from plants pre-treated with 1 ppm ethephon followed by maintaining the plants at control conditions; P-E-HS, pollen derived from plants pre-treated with 1 ppm ethephon followed by exposing the plants to HS conditions. The NSAF values were multiplied by 1000. For enabling fold-change calculation, the value '0.1' was arbitrarily chosen to represent zero.

OUTGROWING Co-Dep Niceness (Part 3)



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