Supplementary data

Ultraviolet radiation exposure time and intensity modulate tomato resistance to herbivory through activation of jasmonic acid signaling

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This file contains information on supplementary Fig. S1, S2 and S3, Table S1, S2 and S3.



Figure S1. Spectral irradiance of the UV lamps after wrapping with white cellulose filters. UV lamps emit residual (< 10% of total PAR) blue (λ 400-450 nm), green (λ 520-560), and yellow (λ 560-590 nm) light. PAR levels in the UV growth chamber did not differ from the control (no UV) growth chamber.



Figure S2. Metabolic responses in control and UV-treated Moneymaker tomato plants was determined at 14 days after the start of the light treatments. Leaf metabolites were analyzed by NMR on leaflets collected from the third/fourth youngest leaf. The ¹H NMR spectra was subjected to principal component analysis and resulted in a model with four principal components explaining 85% of the variance. The first two components, explaining 35.5 and 20.8% of the variance, are represented in the graph. The ellipse defines the Hotelling's T2 confidence region (95%).



Figure S3. Thrips-associated damage (i.e. 'silver damage' symptoms) was evaluated in Moneymaker, Castlemart, *def-1* and *odorless-2* (*od-2*) plants that were first subjected to control or high UV irradiance treatments for 28 days and subsequently used for non-choice whole-plant thrips bioassays. Silver damage was evaluated at 7 days after thrips infestation. Supplemental control (no UV) or high UV irradiance conditions were applied for 30 min d⁻¹. Values represent the mean + SEM (n = 7-9 individual plants). Different letters above bars denote significant differences among groups tested by GLM followed by Fisher's LSD test ($P \le 0.05$).

Table S1. Measurements of UV irradiance were carried out at the plant canopy 1	level	at day	1,
3, 7, 10 and 14 after the start of the light treatments. The integrated irradiance (μW	cm ²) w	'as
converted to kJ $m^2 d^{-1}$ to estimate the daily UV doses.			

-	Integrated Irradiance		Total daily UV doses						
	(μW cm ⁻²)			$(\mathbf{kJ} \mathbf{m}^2 \mathbf{d}^{-1})$					
	UV-B	UV-A	UV	15	20	45	60	120	190
Measurements	280-315	315-400	280-400	15		4 5		120	100
	nm	nm	nm	111111	111111	111111	11111	111111	111111
Day 1	7.6	9.25	16.85	0.15	0.30	0.454	0.606	1.212	1.818
Day 3	7.6	9.06	16.66	0.149	0.299	0.449	0.599	1.198	1.797
Day 7	8.77	10.2	19.01	0.175	0.342	0.513	0.684	1.36	2.04
Day 10	9.19	10.76	19.95	0.175	0.359	0.538	0.718	1.436	2.154
Day 14	10.89	12.6	23.49	0.211	0.422	0.634	0.845	1.6	2.4
Averaged	8.81	10.37	19.19	0.172	0.344	0.517	0.69	1.360	2.04

Table S2. Transitions or specific pair of m/z values associated to the precursors and fragment ions of the analytes measured by LC/MS.

Analyte	$Q1 \; [m/z] \rightarrow Q3 \; [m/z]^a$	CE [V]	Standard
ABA	(-)263.13 → 153.00	9	D6-ABA
JA	$(-)209.12 \rightarrow 59.00$	12	D6-JA
JA-Ile	$(-)322.20 \rightarrow 130.00$	19	D6-JA-Ile
SA	(-) 137.02 → 93.00	15	D6-SA
IAA	$(+) 176.07 \rightarrow 130.00$	-14	D5-IAA
D6-ABA	(-) 269.17 → 159.00	10	
D6-JA	$(-)215.15 \rightarrow 59.00$	10	
D6-JA-Ile	(-) $328.24 \rightarrow 130.00$	19	
D6-SA	$(-)141.05 \rightarrow 97.00$	15	
D5-IAA ^b	$(+) 181.10 \rightarrow 135.00$	-14	
	$(+) 181.10 \rightarrow 134.00$	-14	
	$(+) 181.10 \rightarrow 133.00$	-14	
OPDA	(-) $291,00 \rightarrow 165.00$	18	D6-JA-Ile

CE: collision energy

a Resolution: Q1: 0.7, Q3: 22

b Analyzed as the sum of all three transitions

		Forward Primer	Reverse Primer			
Target gene Gene Identification		5' → 3'	5' → 3'			
PR-P6	Solyc00g174340	GTACTGCATCTTCTTGTTTCC A	TAGATAAGTGCTTGATGTGCC			
TD-2	Solyc09g008670	TGCCGTTAAAAATGTCACCA	ACTGGCGATGCCAAAATATC			
JIP-21	Solyc03g098790	ACTCGTCCTGTGCTTTGTCC	CCCAAGAGGATTTTCGTTGA			
Actin	Solyc03g078400	TTAGCACCTTCCAGCAGATGT	AACAGACAGGACACTCGCACT			

 Table S3. Nucleotide sequence of primers used for qRT-PCR analysis.