

Isoprene enhances leaf cytokinin metabolism and induces early senescence

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Summary

- •Isoprene, a major biogenic volatile hydrocarbon of climate-relevance, indisputably mitigates abiotic stresses in emitting plants. However functional relevance of constitutive isoprene emission in unstressed plants remains contested. Isoprene and cytokinins (CKs) are synthesised from a common substrate and pathway in chloroplasts. It was postulated that isoprene emission may affect CK-metabolism.
- •Using transgenic isoprene-emitting (IE) *Arabidopsis* and isoprene non-emitting (NE) RNA*i* grey poplars (paired with respective NE and IE genotypes), the life of individual IE and NE leaves from emergence to abscission was followed under stress-free conditions. We monitored plant growth rate, above-ground developmental phenotype, modelled leaf photosynthetic energy status, quantified the abundance of leaf CKs, analyzed *Arabidopsis* and poplar leaf transcriptomes by RNA-sequencing in presence and absence of isoprene during leaf senescence.
- •Isoprene emission by unstressed leaves enhanced the abundance of CKs (isopentenyl adenine and its precursor) by >200%, significantly upregulated genes coding for CK-synthesis, CK-signaling and CK-degradation, hastened plant development, increased chloroplast metabolic rate, altered photosynthetic energy status, induced early leaf senescence in both *Arabidopsis* and poplar. IE leaves senesced sooner even in decapitated poplars where source-sink relationships and hormone homeostasis were perturbed.
- •Constitutive isoprene emission significantly accelerates CK-led leaf and organismal development and induces early senescence independent of growth constraints. Isoprene emission provides an early-riser evolutionary advantage and shortens lifecycle duration to assist rapid diversification in unstressed emitters.

Key words: biogenic volatiles, cytokinins, hormones, isoprene, leaf senescence, signalling, transcriptomics, unstressed plants

Introduction

Isoprene (C₅H₈) is the most prominent biogenic volatile organic compound released by the leaves of some of the fastest-growing and most speciose forest trees (Dani et al., 2014a). While isoprene in the atmosphere influences oxidation chemistry, secondary aerosol formation and global radiation budget (McFiggans et al., 2019; Lamkaddam et al., 2021), isoprene also serves emitting plants by mitigating transient heat and oxidative stresses (Behnke et al., 2007; Velikova et al., 2011, Pollastri et al., 2019). Isoprene emission and stress hormones are sensitive to each other and their interactions mediate plant stress responses (Behnke et al., 2010; Parveen et al., 2019; Xu et al., 2020). Over-expression of suppression of isoprene synthase and isoprene fumigation can modulate stress-responsive gene expression (Harvey & Sharkey, 2016; Zuo et al., 2019), although the interactive modes are unclear. Carbon as CO₂ is not a limiting factor for plant growth. Yet, any carbon that is fixed by photosynthesis but not converted into organic compounds (e.g., sugars) under optimal growth conditions is an expensive loss especially for perennials that endure stress and long periods of suboptimal photosynthesis. Isoprene is essentially a 'loss of carbon' fixed by photosynthesis (1 to 2% when unstressed, up to 10% if stressed; Peñuelas & Munné-Bosch, 2005). Thus, it remains unclear why isoprene emission in light occurs continuously in unstressed leaves over their entire lifespan while recent evidence shows continuous emission even in dark-acclimated heterotrophic algae (Dani et al., 2020). All of these suggest fundamental yet unknown house-keeping functions for constitutively emitted isoprene.

Isoprene is made via the plastid-localised methylerythritol phosphate (MEP) pathway by sourcing carbon and energy mainly from photosynthesis. The extended isoprenoid MEP pathway also makes several vital plant hormones (cytokinins, CKs; abscisic acid, ABA; strigolactones, SLs), and accessory photosynthetic and photoprotective pigments (β-carotene and xanthophylls), all directly involved in regulating leaf senescence and lifespan (Dani *et al.*, 2016). In particular, isoprene and CKs are synthesized concurrently from the substrate dimethylallyl diphosphate (DMADP). Adenosine triphosphate (ATP or its dephosphorylated states such as ADP and AMP) combines with DMADP in presence of isopentenyl transferase to give iP (isopentenyladenine)-type CKs, whereas DMADP is directly converted to isoprene by isoprene synthase (Kasahara *et al.*, 2004; Schnitzler *et al.*, 2005). CKs, regarded as plant "youth hormones", regulate leaf

development and expansion, prevent chlorophyll degradation and keep leaves green, maintain chloroplast integrity, and mediate abiotic stress responses (Werner *et al.*, 2001; Zavaleta-Mancera *et al.*, 2007; Nishiyama *et al.*, 2011, Raines *et al.*, 2016).

Given the metabolic proximity and potentially common constraints on the synthesis and functions of isoprene and iP-type CKs, we tested if isoprene affects the course of natural leaf and plant senescence by influencing CK metabolism. We used two verified model plant systems namely (a) the model herbaceous annual *Arabidopsis thaliana* that naturally does not emit isoprene (non-emitting or NE), and *Arabidopsis* genetically transformed to emit isoprene (isoprene-emitting or IE), and (b) the model perennial tree and naturally isoprene-emitting (IE) grey poplar (*Populus x canescens*), and grey poplars where isoprene synthesis and emission was suppressed by RNA-interference (NE). We comprehensively analyzed the impact of constitutive isoprene emission on plant developmental phenotypes and tracked the life of individual isoprene-emitting and non-emitting leaves from emergence to abscission. We measured photosynthesis, modelled photosynthetic energy status of leaves, quantified the changes in the abundance of leaf CKs, and analyzed the poplar and *Arabidopsis* transcriptome by RNA-sequencing in presence and absence of isoprene during leaf senescence.

Materials and Methods

Plant material

Arabidopsis: Arabidopsis thaliana (At) Col-0 ecotype was transformed to obtain isoprene-emitting lines using Agrobacterium and the floral dip method at the plant transformation facility in Michigan State University. The cloning construct included the complete CDS of isoprene synthase (ISPS) from Eucalyptus globulus downstream to the Arabidopsis Rubisco SSU promoter rbcS-1A. Another construct lacking ISPS was used as the empty vector control. Seven independent transgenic lines were obtained (selected on kanamycin) until F3 transgenic seeds were obtained and verified by PCR, and deposited at the Arabidopsis Biological Resource Center (ABRC) in Ohio State University, USA (Zuo et al. 2019). Seeds of At wildtype control (Col-0), empty vector control (line EV-B3), and two isoprene-emitting lines (ISPS-B2 and ISPS-C4) were formally obtained from the ABRC and at that time we only knew that ISPS-B2 and ISPS-C4 lines emitted isoprene. The developmental phenotype and isoprene emission rates were enumerated independently at CNR, Italy. Once in Italy new seed stocks were generated. Seeds were surface sterilised in 70% ethanol, transferred to petri plates containing Murashige and Skoog's agar medium, and vernalised at 4 °C for 48 h. Plates were then transferred to a growth cabinet and allowed to germinate at 18 ± 2 °C, long days (16 h day: 8 h night), light intensity of

100 μ mol m⁻² s⁻¹ from white fluorescent tubes, and 40% relative humidity. Seedlings were individually transplanted to soil-substrate containers placed in plastic trays and watered regularly. Inflorescence and pods were allowed to dry naturally while attached to the plants, and seeds were harvested. An independent set of plants of the four lines was grown under a light intensity of 200 μ mol m⁻² s⁻¹, constituting high-light treatment, keeping all other conditions identical to those used during seed germination.

Poplar: Isoprene-emitting wildtype grey poplars (*Populus x canescens*) along with two transgenic lines in which expression of isoprene synthase (PcISPS) was suppressed by RNA-interference were used. Transgenics were generated and micro-propagated at the Institute of Biochemical Plant Pathology, Helmholtz Centre in Munich (Germany). RNAi-mediated post transcriptional silencing of isoprene synthase in poplars was enabled by the introduction of sense and antisense hairpin sequences (160 bp, highly specific to ISPS) in a binary vector via Agrobacterium mediated transformation (35S: PcISPS-RNAi) (Behnke et al., 2007). Rooted 3-month-old cuttings (15 individuals each of wild type (WT), Empty Vector (EV), RA1 and RA2 isoprene-suppressed lines) were brought to the National Research Council (CNR) research area in Florence (Italy). The saplings were initially grown in 2 L pots containing soil substrate (25%), silica sand (25%), perlite (50% v/v) and slowrelease fertilizer. Young saplings were soon transplanted to 7 L pots (for 2 months) and later transplanted to 18 L pots with the same soil substrate with slow-release fertilizer. Poplars were watered regularly and acclimated to natural seasonal variation in sun light intensity, photoperiod, temperature, and humidity in a CNR experimental facility for genetically modified organisms. After the first season (from April 2018 until December 2018), the main stems were pruned to get stubs (1.5 m) without any leaves or branches. These rooted-stem cuttings from year 1 were transferred to 40 L pots in February of year 2. Budbreak commenced in March, and the second seasonal monitoring of leaf development and senescence went on until December 2019. The microclimate data in the experimental site for the two years of the experiment were gathered from the weather monitoring and modelling centre maintained by Consorzio LaMMA of CNR.

Plant developmental phenotyping

Arabidopsis: The day *Arabidopsis* seeds germinated was noted day 0. Leaf samples were collected at six time points during the plant's lifecycle, classified on the basis of days after germination (DAG) and flowering. Leaves were numbered according to their order of emergence from base to apex, excluding cotyledon leaves. Leaves in position 7, 8, 9, and 10 from the base (the biggest leaves) were marked. A first batch of plants was grown under the same light intensity at which plants germinated (100 μmol photons m⁻² s⁻¹). For this batch, leaves were sampled at 28 DAG (youngest stage sampled), 36 DAG (leaves 7 and 8 fully expanded), 48 DAG

(fully mature plant body, prior to bolting), 56 DAG (early-senescence phase, inflorescence seen in all four lines), 64 DAG (mid-senescence phase), and finally at 76 DAG (late-senescence phase, near-end of lifecycle). Inflorescence was cut and weighed at 56, 64 and 76 DAG. A second batch of *Arabidopsis* plants were grown under 200 μmol photons m⁻² s⁻¹, and leaves in positions 7, 8, 9, and 10 were sampled at 24, 28, 36, and 48 DAG respectively.

Poplar: Poplar leaves were tagged at the time of emergence. Leaves were grouped into three categories. (1) Spring leaves (emerging in May that became lower nodal leaves of the main stem) (2) Summer leaves (emerging in July that became intermediate nodal leaves of the stem), and (3) Autumn leaves (emerging in September, representing mature leaves near the stem apex). Leaf area was measured using a LI-3000 portable area meter (LI-COR Biosciences Inc., USA). Apical extension was measured once every fortnight to once a month, and sub-seasonal trends in apical growth rate were calculated. At the end of the experiment (December), branching pattern was quantified by marking the branches (from base to apex) and by measuring their length and fresh weight post-harvest.

Poplar "no-growth" scenarios: Plants were allowed to grow for one full growing season from April to October, and left undisturbed during natural senescence until December. One-year-old plants were completely pruned (1.5 m from the soil) in the following January, and repotted with new soil in March when they started sprouting. Five plants per each poplar line grew for the entire second growing season without pruning (from April to October, left undisturbed until December). Ten leaves per individual that emerged on the main stem within a span of 10 days in spring (June) were tagged. Another set of six to ten leaves that emerged on the main stem in summer (July) were tagged on the day of emergence for photosynthesis measurements. Ten plants per poplar line grew until middle of July when the apex was cut. Decapitated poplars held onto 15 to 20 fully expanded leaves, whose date of emergence was known and that were 45 ± 10 day old at the time of decapitation. Any new sprouting was removed within a couple of days after emergence. No new branches were allowed to grow for the remaining part of the year.

It is acknowledged that a significant proportion of nutrient recycled from senescing leaves can be supplied to roots and overwintering subterranean storage tissues in many perennial species (Scarascia-Mugnozza *et al.*, 1999; Byne *et al.*, 2020). It is also noted that seasonal senescence in young poplar saplings may appear qualitatively different from typical autumn senescence in adult trees, although both are essentially the same processes, showing high variability among leaves within individuals (Keskitalo *et al.*, 2005; Mattila *et al.*, 2018), and both driven by sink constraints and environmental cues. We did not quantify below-ground changes in this study, since all poplars were grown in nutrient-replete pots. The pots had enough soil volume (avoiding

root limitation) and slow-release fertilizers to support growth for 8 months (avoiding nitrogen deprivation). They were watered adequately throughout the experiment.

Measurements of gas exchange and calculation of energy status and kinetic parameters

Arabidopsis: Net photosynthesis (P_n) , stomatal conductance (g_s) , and intercellular CO_2 concentration (C_i) were measured on fully expanded leaves in positions 8 to 12 (N=5 individual plants) before flowering (40 to 48 DAG) and after flowering (64 to 72 DAG). Measurements were made between 11 am and 3 pm, using a LICOR 6400XT infrared gas analyser (LI-COR Biosciences Inc., USA). Leaves clamped in the circular leaf cuvette (area: 2 cm²) were maintained at 20 °C, light intensity was set to 150 µmol m⁻² s⁻¹ for low-light acclimated plants and 250 µmol m⁻² s⁻¹ for high-light acclimated plants, volatile-free clean air was humidified to achieve ~40% RH (leaf to air vapour pressure deficit was 0.9 to 1.2 kPa) and CO₂ concentration was 400 μmol mol⁻¹.

Poplar. In poplar leaves, all gas exchange measurements were carried out using a LI-COR 6400XT between 10 am and 3 pm, on bright sunny days, and on individual mature leaves at three to four stages of a leaf lifecycle (see above). Leaf temperature was 24 ± 1 °C in April, 28 ± 1 °C in June/July, 24 ± 1 °C in September/October, and 20 ± 1 °C in November/December. In all measurements the relative humidity was maintained between 40 and 65%, ambient CO₂ concentration was 400 μmol mol⁻¹, and the light intensity was set to 1500 μmol m⁻² s⁻¹ (except for leaves senescing in Nov/Dec, when light intensity was set to 1000 μmol m⁻² s⁻¹).

Photosynthesis response to increases and decreases in CO₂ concentrations (40 to 1600 μmol mol⁻¹) was recorded in poplar leaves throughout their lifecycle. $V_{\rm cmax}$ (maximum carboxylation rate of Rubisco) and J(instantaneous electron transport rate) were estimated by fitting net CO₂ assimilation rate or net photosynthesis (P_n) vs. C_i curves using an excel based curve-fitting tool (Sharkey, 2016). Isoprene synthesis is a minor yet important non-photosynthetic and non-photorespiratory energy sink within chloroplasts (Dani et al. 2014b). Hence, the chloroplast energy status of the leaves was quantified using the following equations:

$$R_l = \frac{1}{12} [J - 4(P_n + R_d)] \tag{1}$$

$$J_{c} = \frac{1}{3} [J + 8 (P_{n} + R_{d})]$$

$$J_{o} = \frac{2}{3} [J - 4 (P_{n} + R_{d})]$$

$$v_{o} = \frac{1}{6} [J - \frac{2}{3} (P_{n} + R_{d})]$$
(4)

$$J_o = \frac{2}{3} [J - 4 (P_n + R_d)]$$
 (3)

$$v_o = \frac{1}{6} \left[J - \frac{2}{3} (P_{\rm n} + R_{\rm d}) \right] \tag{4}$$

 $v_{\rm o}$

 $v_c = (P_n + R_d) + \frac{1}{2}v_o$

Term	Description
$P_{\rm n}$	Net rate of CO ₂ uptake per unit of projected leaf area (μmol m ⁻² s ⁻¹)
$C_{\rm i}$	Intercellular CO ₂ concentration (μmol mol ⁻¹)
J	Instantaneous electron transport rate (μmol e- m-2 s-1)
$J_{ m c}$	Proportion of J utilised for carboxylation of RuBP by Rubisco
$J_{ m o}$	Proportion of <i>J</i> utilised for oxygenation of RuBP by Rubisco (photorespiration)
K_{m}	Effective Michaelis-Menten coefficient for carboxylation by Rubisco (at 25 °C)
$R_{\rm d}$	Day mitochondrial respiration rate (μmol m ⁻² s ⁻¹)
R_1	Photorespiration rate (μmol m ⁻² s ⁻¹)
$V_{\rm cmax}$	Maximum rate of RuBP carboxylation by Rubisco (μmol m ⁻² s ⁻¹)
$v_{\rm c}$	Rate of carboxylation by Rubisco

Rate of oxygenation by Rubisco

(5)

Measurements of imaging chlorophyll fluorescence

The maximal quantum yield of chlorophyll fluorescence and photochemical efficiency of photosystem II (F_v/F_m) and the electron transport rate were estimated by saturating-pulse chlorophyll fluorescence imaging using a Walz Imaging PAM (Heinz Walz, Germany). For *Arabidopsis* grown in low-light (100 µmol m⁻² s⁻¹) whole plants (<28 DAG) and leaves cut from fully mature plants (>42 DAG, except WT line) were used during imaging (N=6 whole plants, or individual cut leaves harvested from 4 individuals). In poplars, individual cut leaves (2 leaves of same age from N=3 individuals) were sampled at 30, 60, 90, 120, and 180 days after emergence (DAE) and used for fluorescence measurements. All leaves were dark-acclimated for 30 min before imaging.

Isoprene sampling and quantification

Isoprene sampling was done from fully expanded leaves for *Arabidopsis* one or two leaves per individual (*N*=4 individual plants per line). In poplars, two to three fully expanded mature leaves (leaf age between 30 to 60 days) per individual per line were sampled (*N*=4 to 5 individual plants per line). A portion (300 mL min⁻¹) of the LI-COR cuvette outflow was diverted using a mass flow pump (AC Buck Inc. FL, USA) onto a cartridge filled with absorbents (30 mg each of Carbosieve X and Carbosieve B, Supelco, USA). Isoprene from emitting lines was quantified following the protocol for volatile thermal desorption gas chromatography-mass spectrometry (after Dani *et al.*, 2017). Briefly, an Agilent 5975 gas chromatograph-mass spectrometer (GC-MS; Agilent Technologies, USA) system was fitted with an HP-INNOWax (50 m length, 0.2 mm ID, 0.4 μm film) column. Thermal desorption was executed by a Twister* multipurpose autosampler and TD unit (Gerstel Technologies, Germany) fitted with an e-Trap cryofocussing system (Chromtech, Germany). The GC separation programme was 40 °C for 1 min, reaching 110 °C at 5 °C min⁻¹, held for 10 min, and then increased to 260 °C at 30 °C min⁻¹ and held for 2 min. Isoprene standards were prepared in 2 L Tedlar bags (Sigma-Aldrich, USA) containing nitrogen, and analysed as above. Samples collected from the cuvette headspace containing control lines were treated as zero-isoprene controls.

Extraction and quantification of cytokinins

Cytokinins (CKs) were extracted from fully expanded *Arabidopsis* leaves (position 7 to 10) before flowering under low-light (at 36 and 48 DAG) and high-light treatments (at 28 and 36 DAG). For *Arabidopsis*, leaves from at least four individuals were pooled to get one biological replicate with sufficient leaf mass and four such biological replicates were analysed. This meant ≥10 individuals per line for each treatment group were

sampled. CKs were also extracted from fully mature healthy leaves (45 to 60 DAE) in all poplars. Cut portions of two to three poplar leaves (of similar age) from one individual was pooled to get one true biological replicate and four such biological replicates were analysed. CKs were extracted in acidified aqueous methanol, purified by two solid-phase extraction (SPE) steps and subsequently measured with a High-Performance Liquid Chromatograph coupled to a Tandem Mass Spectrometer (LC-MS/MS, Applied Biosystems, USA, as in Schäfer et al., 2015; Dobrev & Kaminek, 2002). 30 to 400 mg ground plant tissue was extracted twice with 800 μL MeOH: H₂O: HCOOH (15:4:1) at -20 °C. Deuterated internal CK standards in the form of 0.2 ng [²H₆] iPR, 0.2 ng [2H₅] tZR, and 1 ng [2H₅] tZ were supplemented in the first extraction step (standards from OlChemIm s.r.o., Czech Republic). The expanded hormone abbreviations for internal standards are - iPR: isopentenyladenine riboside; iP: isopentenyladenine; and tZR: trans zeatin riboside. Also see the caption under Figure 2 and Table S1 for additional details. Extraction and SPE were performed in 96 Well BioTubes (Arctic White LLC) and 96-Well Deep Well Plates (Thermo Scientific). The first SPE step was performed on a Multi 96 HR-X column (Macherey-Nagel, www.mn-net.com/us/chromatography/) conditioned with extraction buffer. The flow through was collected and the MeOH was evaporated at 42 °C under constant nitrogen flow. Then, 850 µL of 1 M HCOOH was added to the samples and loaded on a Multi 96 HR-XC column (Macherey-Nagel) pre-conditioned with 1 M HCOOH. Sequentially 1 mL each of 1 M HCOOH, MeOH, 0.35 M NH₄OH were added and eluted. Finally, CKs were eluted with 1 mL 0.35 M NH₄OH in 60% MeOH. The second SPE was performed using a Chromabond Multi 96 vacuum chamber. After evaporation, samples were reconstituted in 50 µL 0.1% acetic acid, CKs were chromatographically separated on a Zorbax Eclipse XDB-C18 column (50×4.6 mm, 1.8 µm) at 25 °C fitted to an Agilent 1200 HPLC system (Agilent Technologies, USA). Solvent A (water, 0.05% HCOOH) and solvent B (acetonitrile) mixture was suppled at 1.1 mL min⁻¹ [0 to 0.5 min, 95% A; 0.5 to 5 min, 5 to 31.5% B in A; 5 to 6.5 min, 100% B; and 6.5 to 9 min 95% A]. The LC was coupled to an API 6500 tandem mass spectrometer (Applied Biosystems SCIEX, Germany) equipped with a Turbospray ion source and quadrupole mass analyser. Data was acquired and processed using Analyst 1.6.3 software (AB SCIEX). The MS was in positive ionization mode (MRM modus) to monitor analyte parent to product ion conversion (Table S1). We did not have internal standards for and hence could not quantify the immediate CKnucleotide precursors of active iP and tZ. Hence, those are not reported. CK-riboside levels in *Arabidopsis* leaves sampled just before flowering are given in Table S2.

RNA extraction, library preparation, RNA-seq and differential gene expression analysis

Fully-expanded *Arabidopsis* leaves in position 8, 9, 10 (from the base) were sampled from 8 individuals per each line before flowering i.e., 36 to 40 days after germination (DAG) under low-light treatment, and 28 to 36

DAG under high-light treatment. Summer leaves from poplars were sampled from four individuals (per line) at 60 and 90 days after emergence (DAE). Leaves were frozen in liquid nitrogen and stored at -80 °C. Arabidopsis leaves were pooled (2 to 3 individuals forming one biological replicate) to get sufficient starting material for RNA extraction. Leaves from two Arabidopsis lines (non-emitting EV-B3 control and isopreneemitting ISPS-C4) and two poplar lines (emitting WT control and isoprene-suppressed RA1) were used for RNA-seq. Sampled leaves were ground in liquid nitrogen and total RNA was extracted from approximately 100 mg of ground leaf material (three biological replicates per genotype, per leaf life-stage) using the mirPremier Isolation Kit (Sigma-Aldrich) following the manufacturer's instructions. Ribosomal RNA depleted strand-specific RNA libraries were generated (in triplicate) using the Illumina TruSeq Stranded Total RNA library preparation kit with Ribo-Zero Plant (Illumina Inc., USA). Libraries were quantified using Agilent 2100 Bioanalyzer RNA assay (Agilent Technologies, USA), pooled in equimolar amount and sequenced on a single lane on the Illumina NovaSeq 6000 sequencer to get 150 bp paired-end reads (2×15 million total reads). Images from the instruments were processed following the manufacturer's software pipeline to generate FASTQ sequence files. Adaptor sequences and low quality 3' ends were removed from short reads using respectively CUTADAPT (Martin, 2011) and ERNE-FILTER (Del Fabbro et al., 2013). After trimming, only pairs with both reads longer than 50 bp were retained. Trimmed reads were aligned against the *Populus tremula* v2.2 reference genome (ftp://plantgenie.org/Data/PopGenIE/Populus tremula/v2.2) and the A. thaliana (TAIR10) reference genome (http://www.Arabidopsis.org) using STAR v2.7.2b (Dobin et al., 2013) with default parameters. The htseqcount python utility (Anders et al., 2015) was used to calculate gene-based read count values considering only uniquely mapping reads. The HTSeq count data were used as the input to measure differential gene expression using the Bioconductor package DESeq2 v1.14.1 (Love et al., 2014) implemented in R. The raw counts of each gene were normalised to adjust for different sequencing depths across samples.

Statistical analysis

Normality of observed values (within lines) was tested using the Kolmogorov-Smirnov test. Differences in means among lines for net photosynthesis (CO₂ assimilation rate), electron transport rate, stomatal conductance, above ground biomass, and inflorescence weight (only *Arabidopsis*) were tested by one-way analysis of variance (ANOVA) followed by a post-hoc Tukey's multiple comparison test ($N \ge 10$ biological replicates per line, α =0.05). Differences in poplar photosynthesis and chloroplast energy status were tested using data collected from ≥ 5 leaves per individual at each sampled leaf age ($N \ge 6$ biological replicates per line). Differences in isoprene emission rate and cytokinin abundance was verified by either Kruskal-Wallis H test for comparing medians ($N \ge 4$; α =0.05) or by Games-Howell test for comparing means when variances

were unequal ($N \ge 4$; α =0.05). Leaf sampling details are given under their respective sub-headings under methods. Differentially expressed genes (from DESeq) were identified using pairwise comparisons at an adjusted p-value (FDR) threshold of ≤ 0.01 for determining significance. Transcripts with $\log_2[\text{fold change}] > +1.5$ and <-1.5 at $p_{\text{adj}} < 0.001$ were considered biologically significant, truly sensitive to presence or absence of isoprene, and shortlisted for functional interpretation. The foldchange heatmap was generated using the open-source Morpheus (https://software.broadinstitute.org/morpheus) matrix visualization application. All other statistical tests were carried out using Minitab 18.1 statistical package (Minitab Inc, USA).

Results and Discussion

Isoprene enhances leaf cytokinin metabolism, accelerates plant growth rate, strengthens apical dominance, and induces early flowering, early leaf and plant senescence

Isoprene-emitting (IE) lines of Arabidopsis and poplar showed faster growth (Figs. 1a, 1b, 1e, 1f), early leaf senescence (monitored by early photosynthesis decay, Fig. 1c, 1d, 1g-i), and above-ground plant phenotype that was distinct from non-emitting (NE) Arabidopsis and poplar lines (Fig. 1b, 1f). Significantly higher levels of isopentenyladenine riboside (iPR, Fig. 2b) and its free base derivative isopentenyladenine (iP; Fig. 2c) in the IE leaves of both Arabidopsis and poplar with respect to NE leaves show that isoprene has a direct positive impact on the plastid-localised synthesis of isoprenoid-type CKs via the MEP pathway. Senescence associated declines in net photosynthesis (P_n) , maximum carboxylation capacity of Rubisco (V_{cmax}) , maximum photochemical yield of PSII (Y(II)), and instantaneous photosynthetic electron transport rate (ETR) occurred sooner and were steeper in IE than in NE Arabidopsis and poplar (Fig. 1c, 1d, 1g; Fig. S1a). However, it is surprising that high CK abundance which is normally expected to delay senescence (Gan & Amasino, 1995; Werner et al., 2001), led to early leaf senescence in IE plants. While significant overexpression of genes coding for two-component signal transduction / response regulators involved in CK-signaling in IE leaves suggests greater CK activity, we also observed significant upregulation of cytokinin oxidase/deoxygenase in IE Arabidopsis (AtCKXI and AtCKX6, Fig. 3a) and poplar (PtCKX5 and PtCKX7; Fig. 3b), indicating greater degradation of CKs. Leaf age-specific changes in chloroplast energy status were altered during seasonal senescence in NE poplar leaves (Fig. S1b-d; not measured in Arabidopsis). SIGMA factors (facilitators of chloroplast RNA-polymerase) and other transcription factors involved in chloroplast replication were enriched in IE Arabidopsis (Fig. 3a) and consistent with this, SIGMA factors and transcripts representing corephotosynthetic genes were depleted in NE poplar leaves (Fig. 3b), all indicating faster chloroplast metabolic

rate in presence of isoprene. We propose that faster cell metabolism and faster senescence in IE leaves are driven by increased activity and recycling of CKs.

IE poplars showed a distinct within-plant distribution of biomass due to their apically dominant and branch-heavy phenotype (Fig. 1e, 1f, Table S3). Indeed, the *LOG* genes *PtLOG1* and *PtLOG3* (*LONELEY GUY* family), coding for cytokinin riboside 5'-monophosphate phosphoribohydrolase involved in the activation of CKs and regulation of shoot apical meristematic growth (Kurakawa *et al.*, 2007), were significantly overexpressed in the healthy mature leaves of apically dominant IE poplars (Fig. 3b). IE leaves were richer than NE leaves in tZR (trans-zeatin riboside; *Arabidopsis*), tZOG (trans-zeatin O glucoside; poplar, Fig. 2d), and even cytosolic tRNA-derived cZR (cis-zeatin riboside; *Arabidopsis*, Table S2; also see Notes S1). Enlargement and thinning of IE leaves in *Arabidopsis* (Fig. S2d), and less consistently also in IE poplar, were likely triggered by altered zeatin ratios, which can affect leaf size (Osugi *et al.*, 2017). It remains to be verified if leaf CK enrichment in IE leaves also affected the ratio between CK and auxins, the latter being a key regulator of apical dominance. Since iPR is transported in the phloem from leaves to all other meristems (Hirose et al. 2007), excess foliar iPR in IE leaves may cause faster tissue differentiation and expansion in all the organs of IE plants.

IE *Arabidopsis* flowered significantly earlier compared to NE control under both low-light and high-light (Fig. 2c, Fig. S2e). Since high exogenous or endogenous CKs promote flowering and CK depletion delays flowering in annuals (Choudhury *et al.*, 1993; Corbesier *et al.*, 2003), CK enrichment and heightened degradation of CKs in IE *Arabidopsis* leaves likely contributed to early-flowering. Isoprene had either negative or no influence on genes involved in gibberellin (GA) signaling (Fig. 3a). Thus, CKs and not GAs likely induce early flowering in IE *Arabidopsis*, and perhaps also in plants fumigated with isoprene (Terry *et al.*, 1995). *COL9* (*CONSTANS-LIKE 9*) and *ELF4* (*ETS-related transcription factor Elf-4*) are negative regulators of flowering time (Doyle *et al.*, 2002; Cheng and Wang, 2005; Kim *et al.*, 2005), and their significant downregulation in IE *Arabidopsis* under high-light may have contributed to longer hypocotyls (Zuo *et al.*, 2019) and early flowering. The poplar *COL9* and *ELF3* transcripts were significantly enriched in NE leaves (Fig. 3b), which was consistent with their suppression in IE *Arabidopsis* leaves. Hence, isoprene and CK-mediated changes in the circadian and photoperiodic signaling pathways likely contributed to early flowering and early senescence in IE plants. Given that isoprene interacts with stress-responsive pathways (Zuo *et al.*, 2019; Frank *et al.*, 2021), early-flowering in IE *Arabidopsis* could be partly also an evolutionary consequence of upregulated stress signaling.

Isoprene associated CK enrichment induces stress and defense response genes, with limited benefits under stress-free conditions

The impact of isoprene on stress response, expression of genes involved in stress mitigation and less clearly also on growth are acknowledged in different experimental contexts recently (Behnke et al., 2010; Zuo et al., 2019; Monson et al., 2020; Xu et al., 2020; Frank et al., 2021). However, it is unlikely that isoprene played either thermo or photoprotective role in Arabidopsis grown at 18 ± 2 °C under low-light intensity (100 µmol photons m⁻² s⁻¹) since the photochemical parameters ETR and Y(II) of IE and NE Arabidopsis were similar (Fig. S3). Photosynthetic capacities of young and mature NE and IE poplar leaves were also comparable even at relatively hot ambient temperatures >35 °C (Fig. 1g). Thus, the difference in the senescence course of IE and NE leaves was not caused by extrinsic stresses, where isoprene is known to be beneficial (Way et al., 2011; Vickers et al., 2011). It was not tested if changes in foliar hormonal balance due to isoprene suppression caused any 'intrinsic stress' that involved metabolically damaging changes in free radicle and redox status. Any intrinsic stress in NE leaves could have led to them senescing sooner than IE leaves, which was not the case. In Arabidopsis, thinning of IE leaves under low-light may have led to lower P_n , but not when light-intensity was increased (200 µmol photons m⁻² s⁻¹) while keeping all other conditions identical. However, one of the IE lines (ISPS-C4) showed the highest isoprene emission rate and P_n under high-light (Fig. 1c, 2a) where expression of PSI and PSII light harvesting complexes was less suppressed relative to NE leaves (Fig. S4), indicating photoprotective effects of isoprene. A sweeping downregulation of heat shock proteins (HSPs, Fig. 3b), differential expression of NAC and WRKY transcription factors, and ethylene response factors (ERFs) involved in abiotic stress responses (Figs. S5-S7) in NE poplar leaves broadly agreed with previous studies showing interactions between isoprene and stress-response pathways (Behnke et al., 2010; Zuo et al., 2019; Xu et al., 2020). Noting that CKs themselves can impart photoprotective benefits, and since stress-induced leaf senescence involves CK depletion and senescence is delayed by CK enrichment (Rivero et al., 2007; Cortleven & Schmülling, 2015), we attribute known tolerance for transient heat and high-light stresses in IE leaves also to localised enhancement of foliar CK synthesis and activity. The concentration of active CK-free base (iP) increased by ten times in high-light in all Arabidopsis lines (Fig. 2c), indicating stimulation of CK-riboside to active CK-freebase conversion by light, which was most pronounced and significant in IE leaves (Fig. 2c). Such high levels of iP (and no change in iPR; Fig. 2b) may have partly caused downregulation of leaf AtIPT genes (CK synthesis) in IE leaves under high-light (Fig. 3a). However, as in low-light, transcripts of AtCKX genes (CK degradation) were enriched in early-senescing IE leaves under high-light (Fig. 3a). Besides, sustained CK activity under prolonged stress can be detrimental (Nishiyama et al., 2011), and this, among other costs associated with fast-growth, can also make IE plants more susceptible to premature mortality under severe stresses (Fig. 4).

Early leaf senescence in isoprene-emitting poplars is an innate behaviour not driven by faster growth rate

Plant growth and reproductive strategies i.e., growing fast or slow and flowering early or late, influence the timing and pace of leaf senescence at least in annuals, and the same is evident in IE Arabidopsis. Accelerated development and early flowering in IE Arabidopsis was associated with lighter above-ground vegetative body mass and faster body (leaf) mass loss during senescence (Fig. S1d). It is difficult to verify the same in longlived trees. However, body size maturation and attainment of reproductive maturity may determine when and how whole-plant senescence proceeds even in perennials (Dani & Kodandaramiah, 2019). Although the total biomass of IE and NE poplars was comparable in our study (also in Way et al., 2011; Behnke et al., 2012; Monson et al., 2020), the early-rising and apically dominant (light-competitive) IE trees were expected to attain adulthood sooner and to undergo actuarial and demographic senescence earlier than closely related NE species. However, timing and pace of autumn leaf senescence, despite being a strongly heritable character preserved over generations in deciduous perennials, is sensitive to year-on-year variation in seasonal growth and environmental stress factors (Panchen et al., 2015; Zohner & Renner, 2017; Michelson et al., 2018). The role of fast or slow growth strategy on the timing and pace of autumn leaf senescence has not been explicitly tested in poplars. Timing of bud-burst and leaf flushing correlate strongly with growth rate, growth duration and biomass accumulation in poplars (Yu et al., 2001; Muller et al., 2013). Spring-sprouting occurred at the same time in all four popular lines, which ruled out spring-phenology influence on senescence regimes of IE and NE poplars. However, the impact of growth rate on leaf senescence was less clear in poplars and varied within plants. Leaf lifespan was shorter when the length of sylleptic branches was longer in NE poplars (Fig. S8c), suggesting that modular nutrient remobilization and relative strength of local sinks (adjacent to senescing nodal leaves) was a key factor that determined which leaf senesced first and at what rate (Table S3, Notes S2; Sprugel et al., 1991). We note that IE leaves may have abscised early by the virtue of early senescence and P_n could not be monitored at high frequency to examine if IE leaves senesced faster and not just sooner. Instead, we tested whether the observed early senescence (early decline in age-specific P_n ; Fig. 1g) in IE leaves on the main stem occurred as a consequence of faster apical growth and metabolism (proxied by chloroplast status) of IE poplars relative to NE poplars.

We imposed "no new growth" condition on both IE and NE poplars where apical and axillary sprouts were pruned once and repeatedly until the end of the experiment in December. Decapitation and "imposed nogrowth" treatment extended leaf lifespan by more than a month in all poplars confirming the importance of the strength of growing sinks in pacing leaf senescence. However, in apically intact populars, an early decline in P_n was observed even in late-emerging apical IE autumn leaves relative to equivalent NE leaves, both presumably not supplying remobilized nutrients to any significant growing sink late in the season (Fig. S8a). Even under "no-growth" scenario, P_n declined significantly sooner in senescing IE leaves than in senescing NE poplar leaves (Fig. S8d), and the early-senescing phenotype of IE leaves was preserved (Fig. S9). These observations show that early- or late-senescing tendency of leaves is governed by isoprene-induced reprogramming of biological processes that are innate and distinct from those affecting seasonal sink constraints and growth rates. We attribute the earlier senescence primarily to enhanced CK metabolism and CK-targeted transcriptional regulation in presence of isoprene in unstressed condition, and other isoprene-associated changes that are detectable (although likely always true) under stressful conditions. The question, what about leaf senescence in those deciduous and evergreen trees that emit tiny amounts of isoprene and deciduous trees that do not emit isoprene at all, does arise. These trends need to be further tested in other deciduous isoprene-emitting tree genera that are typically faster-growing and more diverse than related non-emitting genera (Dani et al., 2014a). At this stage, isoprene emission seems to act as a pacemaker of leaf senescence when it occurs at measurable, thus significantly high rates. However, absolute rates of emission may be less important than emission relative to rate of photosynthesis and other demands in chloroplasts. There are limitations to studying seasonal senescence in juvenile poplars, which have different body architecture and growth constraints compared to adult trees. Nonetheless, the leaf level metabolic changes due to isoprene are likely under identical genetic and physiological constraints in both juveniles and adult trees in most species (including poplars), except perhaps in cases where tree maturation also alters leaf structure. Tissue-specific metabolic constraints are as important as temperature, photoperiod, nutrient availability and organismal age in determining the timing and pace of seasonal (and whole plant) senescence.

Global warming and CO₂ fertilization by the end of this century are expected to either delay autumn senescence due to extended periods of favorable temperatures or cause early autumn senescence due to quicker growth and water-limited conditions (Richardson *et al.*, 2010; Chen *et al.*, 2020; Zani *et al.*, 2020). More isoprene could be released if growing seasons get longer and/or warmer in the future (Peñuelas *et al.*, 2009; Sharkey & Monson, 2014). Our result, that isoprene-emission changes CK homeostasis and induces early leaf senescence

irrespective of growth rates and sink constraints, becomes an important factor that influences the course of autumn leaf senescence and limits the release of carbon as isoprene from deciduous forests.

Conclusion

Isoprene-emitting plants are more tolerant to transient abiotic stresses than non-emitters, but an overarching explanation for the relevance of constitutively emitted isoprene in unstressed plants was missing. We show that isoprene emission from unstressed leaves significantly enhances leaf CK synthesis and induces a remarkably consistent pattern of CK-associated transcriptional regulation in both a herbaceous and a tree model species. Early leaf senescence in presence of isoprene and associated changes in CK metabolism suggests that even a constitutive plant volatile like isoprene has a non-trivial stimulatory impact on development, reproduction, and senescence. These results unveil a hitherto unknown significance for constitutive isoprene emission that (a) makes emitting-species more competitive by hastening their development, and more critically (b) curtails leaf and plant lifespan by accelerating ageing and senescence (Fig. 4). Faster development, early maturation, early senescence and shortening of generation time due to isoprene emission likely confers a major evolutionary advantage in emitting plants by not only imparting transient stress-resilience (well recognized) but more fundamentally by facilitating stress-escape (shorter life cycle), faster species diversification and potential colonization of new favorable habitats.

The consequences of endogenous isoprene on plant development and phenotype appears remarkably similar to more-widely studied stimulation of plant growth by various exogenous microbial volatiles (Junker & Tholl, 2013; Sharifi & Ryu, 2018). A key distinction is that continuous endogenous emission (and not exogenous exposure) incurs metabolic costs to the emitter, which may directly affect organellar carbon and energy status and hormone synthesis (our study) and also hormone-mediated signalling (various studies). We anticipate that other endogenous volatiles, which are so far purely implicated in stress mitigation and inter-organismal communication, will be involved in interactions with canonical hormone-mediated plant developmental pathways (Dani & Loreto, *invited/under review*). Future experiments involving mutants of CK synthesis and signalling will shed more light on specific mechanisms of interference from isoprene in CK-mediated changes in unstressed leaf and plant metabolism.

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Supporting Information:

- Fig. S1 Chloroplast energy status of poplar leaves during senescence
- Fig. S2 Quantum yield of photosystem II in Arabidopsis
- Fig. S3 Growth and flowering in isoprene-emitting *Arabidopsis*
- Fig. S4 Isoprene and the expression of photosystem genes in Arabidopsis
- Fig. S5 Isoprene and the expression of poplar NAC transcription factors
- Fig. S6 Isoprene and the expression of poplar WRKY transcription factors
- Fig. S7 Isoprene and the expression of poplar ERFs
- **Fig. S8** Net photosynthesis (P_n) in apically intact and "no growth" poplars
- Fig. S9 Leaf photographs showing early senescence in apically intact and "no-growth" poplars
- **Table S1** Parameters of LC-MS/MS analysis of cytokinins
- **Table S2** Cytokinin supplementary information including cis zeatin-riboside (cZR) for *Arabidopsis*
- **Table S3** Poplar above-ground phenomic data
- Notes S1 Supplementary note associated with Table S2
- Notes S2 Supplementary note associated with Table S3

Acknowledgments

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Author contributions: KGSD conceived the hypothesis, visualized, designed, led the study, and conducted the experiments with FL. TDS and J-PS provided the transgenic *Arabidopsis* and poplar lines, respectively. S Pollastri helped raising the *Arabidopsis* seed stock, along with FL, helped with logistics. MR extracted and quantified cytokinins. S Pinosio assembled the RNA-seq raw reads and the transcript abundance data. KGSD monitored leaves from emergence until senescence, quantified leaf photosynthesis, chloroplast energy status, leaf and plant phenotype, analysed hormonal and differential gene expression data. KGSD wrote and revised the manuscript with inputs from FL and other authors.

Data availability:

All data are available in the main text or the supplementary materials. The RNA seq data that support the findings of this study are deposited to Sequence Read Archive of NCBI and the concerned accession numbers are PRJNA694035 and PRJNA690701 and the URLs are

- (i) https://www.ncbi.nlm.nih.gov/Traces/study/?acc=PRJNA694035
- (ii) https://www.ncbi.nlm.nih.gov/Traces/study/?acc=PRJNA690701

Figure legends

Fig. 1. Growth rate and leaf senescence course in (a-d) isoprene-emitting (white columns) and non-emitting (grey-columns) *Arabidopsis* and (e-i) isoprene-emitting (white columns) and isoprene-suppressed (grey-columns) poplar. For *Arabidopsis*, the responses of plants grown under low-light (100 μ mol photons m⁻² s⁻¹) and under high-light (200 μ mol photons m⁻² s⁻¹) are compared. (a) Growth rate for the first 28 days after germination (DAG), (b) Photographs showing above-ground plant phenotype, (c) Plant age-specific net photosynthesis (P_n) in fully mature and senescing *Arabidopsis* leaves, (d) Representative chlorophyll fluorescence images showing the quantum yield of photosystem II (Y(II)) and early senescence in isoprene-emitting leaves of *Arabidopsis* (72 DAG). For poplars, (e) Apical growth rate (rate of change in the main stem height) in mid-summer (July), late-summer (Aug-Sept), and early-autumn (Oct), (f) Plant phenotype in 10-month-old saplings. (g) Leaf age-specific net photosynthesis (P_n) and (h) Maximum carboxylation rate by Rubisco (V_{cmax}) for leaves at 30, 60, 90 and 120 days after emergence (DAE), (i) Chlorophyll fluorescence-based estimation of maximum photochemical yield of photosystem II (Y(II)) for summer-leaves of age 60 and 90 DAE. Six to ten individual plants per line (true biological replicates) were sampled for both *Arabidopsis* and poplar. See methods for details of technical replicates. The box for each plant type in each plot includes the mean (circle), median (horizontal line), and the box spans lower and upper quartiles. The whiskers span the full data range. Tukey's test, $\alpha = 0.05$. Means that are

significantly different do not share alphabetical letter codes. WT: Wild Type (white bars); EV-B3 (*Arabidopsis*) and EV (poplar): Empty Vector control (grey bars).

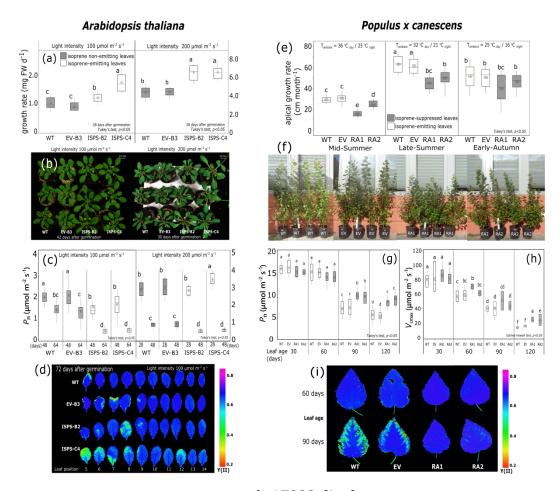
ISPS-B2 and ISPS-C4: transgenic isoprene-emitting *Arabidopsis* lines (white bars); RA1 and RA2: transgenic isoprene-suppressed (nearly non-emitting) popular lines (grey bars).

Fig. 2. Isoprene emission rate and cytokinin concentration (CKs) in Arabidopsis (left panels) and poplar leaves (right panels). Measurements from both low (100 µmol photons m⁻² s⁻¹) and high-light (200 µmol photons m⁻² s⁻¹) treatments are shown for Arabidopsis. (a) isoprene emission rate (b) isopentenyladenine riboside (iPR) (c) isopentenyladenine (iP), (d) trans zeatin riboside (tZR) for Arabidopsis and trans zeatin-o-glucoside (tZOG) for poplars, (e) trans zeatin (tZ). Leaves from at least four individual plants per line (biological replicates) were pooled to get one true replicate and four such pooled replicates per line were analysed in Arabidopsis, which meant that >10 individuals were sampled per line. Arabidopsis leaves sampled were 36 to 48 days old (low-light), 28 to 36 days old (high-light), all before flowering, and were maintained at 20 °C during sampling. In poplars, cut parts of two to three leaves per individual (per line) were pooled to get one biological replicate and four such true biological replicates (individuals) were sampled. Poplar leaves were 45 to 60 days old, and were maintained at a light intensity of 1500 µmol photons m⁻² s⁻¹ before and during sampling. The horizonal line within the box corresponds to the median and the box marks the lower and upper quartiles. Circles with + symbol mark the mean. Statistical significance of differences in isoprene emission rate were ascertained by one-way ANOVA followed by a post-hoc Tukey's test. For CKs, both Kruskal-Wallis (KW) H test for medians and ANOVA followed by Games-Howell (GH) test for equality of means were applied (all α =0.05). Means that are significantly different do not share alphabetical letter codes. WT: Wild Type (white bars); EV-B3 (Arabidopsis) and EV (poplar): Empty Vector control (grey bars). ISPS-B2 and ISPS-C4: transgenic isoprene-emitting Arabidopsis lines (white bars); RA1 and RA2: transgenic isoprene-suppressed (nearly non-emitting) popular lines (grey bars).

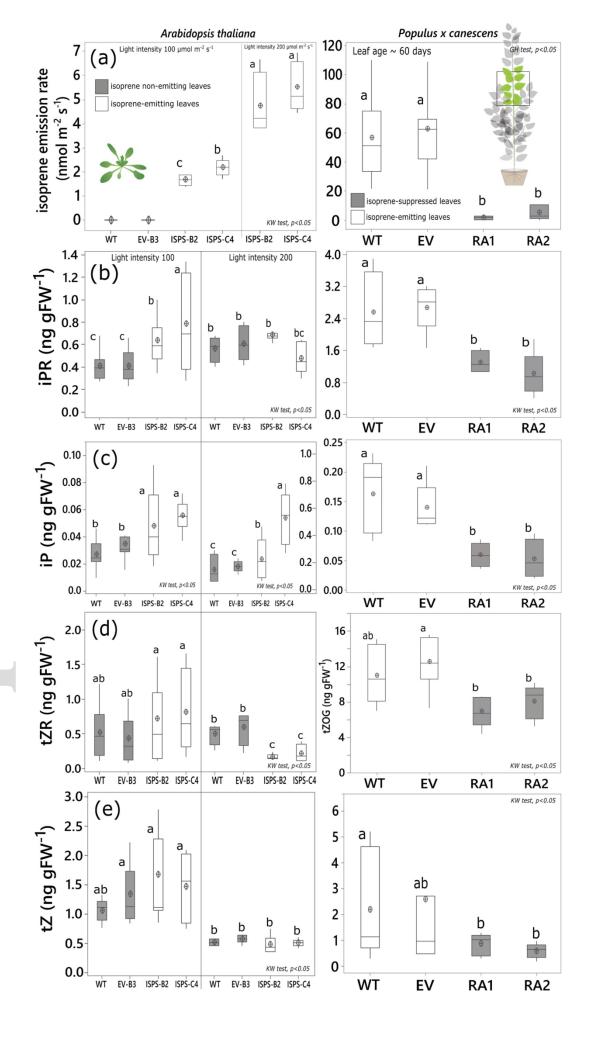
Fig. 3. Impact of isoprene emission in *Arabidopsis* (a), and isoprene suppression in poplar (b) on the differential expression of genes involved in cytokinin synthesis, cytokinin degradation, cytokinin signaling, photosynthesis, chloroplast metabolism, non-cytokinin hormone signaling, senescence associated transcription factors and flowering. Most significant $\log_2[\text{fold change}]$ in transcript abundance during pairwise comparisons are shown. Darker the blue, more depleted are the transcripts and similarly brighter the red, more enriched are the transcripts. Wherever $\log_2[\text{fold change}]$ is >+1.5 and <-1.5, the corresponding p_{adj} is <0.001. Wherever the fold change is less prominent but significant, the corresponding p_{adj} is often <0.05. Non-significant changes are represented by white blanks without values. In panel (a), the change in transcript abundance is due to the presence of isoprene in isoprene-emitting (IE) *Arabidopsis* leaves relative to non-emitting (NE) control except in column 1 where the effect of HL in NE leaves is indicated. The gene IDs correspond to the latest annotation of *Arabidopsis thaliana* genome (TAIR10). In panel (b), the change in transcript abundance is shown for healthy (60 days after leaf emergence) and senescing (90 DAE) leaves of isoprene-suppressed (nearly non-emitting) RA1 relative to IE control (WT). Darker blue represents depletion and brighter red represents

enrichment in NE leaves at 60 and 90 DAE. In the 3rd column of panel (b) comparison between senescing IE and healthy IE leaves shows changes in transcript abundance during natural senescence in IE control leaves. The gene IDs and chromosomal loci correspond to the latest annotation of the whole genome of *Populus trichocarpa* at NCBI.

Fig. 4. Illustration of metabolic and evolutionary consequences of isoprene emission capacity in plants. The statements in red are the most significant novel insights obtained from the current study, and the remaining facts (some but not all verified by other published studies) are also summarized.



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POPTR_007G121200 POPTR_006G233800

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(-)							
(a)	No isoprene control high-light	Isoprene-emitting high-light	Effec	t of isoprene	-4.00 -0.20 0.20	4.00	
GENE ID	(relative to low-light)		low-lig	ht high light	Arabidopsis thaliana		
AT3G63110	1.8	-0.8		-2.4	IPT3 / Isopentenyltransferase 3	Cytokinin synthesis	
AT3G23630	2.5	-0.7		-2.0	IPT7 / Isopentenyltransferase 7	Cytokiriiri syritilesis	
AT2G41510	-0.6	0.9		1.6	CKX1 / CK oxidase/dehydrogenase 1	Cytokinin degradation	
AT3G63440	-0.9	0.4		1.7	CKX6 / CK oxidase/dehydrogenase 6	Cytokiriiri degradatiori	
AT3G16857			0.3	0.8	ARR1 / Response regulator 1		
AT2G25180	0.3	0.3	0.4	0.7	ARR12 / Response regulator 12		
AT1G74890	-2.2	0.6		2.0	ARR15 / Response regulator 15		
AT2G40670	-0.7			1.1	ARR16 / Response regulator 16	Cytokinin signaling	
AT4G23750	2.2	-0.8		-2.6	CRF2 / Cytokinin response factor 2		
AT5G53290	2.4	-0.8		-2.9	CRF3 / Cytokinin response factor 3		
AT2G46310	1.0			-1.6	CRF5 / Cytokinin response factor 5		
AT5G38410	-4.0	-2.2		1.3	RuBisCO SSU	Photosynthesis / Glycolysis	
AT1G12900	-2.5	-1.9		0.3	GAPDH / Glyceraldehyde 3-phosphate dehydrogenase	1 Hotosynthesis / Olycolysis	
AT1G64860	-1.2	-0.6		1.0	Sigma factor A		
AT5G24120	-1.0	0.6		1.9	Sigma factor E	Chloroplast metabolism	
AT2G16070			0.4	1.0	Plastid division 2	Omoropidot motaboliom	
AT5G44190	-1.3	0.6		2.4	GOLDEN2-like 2		
AT1G63970	-1.0	0.5		1.0	2C-methyl-D-erythritol 2,4-cyclodiphosphate synthase	MEP pathway	
AT4G34350	-2.1	-0.5		1.3	4-hydroxy-3-methylbut-2-enyl diphosphate reductase	mer paamay	
AT5G55250	-1.1 -1.8	0.7 -1 .9		2.6	IAA carboxylmethyltransferase 1	Auxin synthesis and signaling	
AT3G23050	-1.6 -1.5	-1.9			Auxin-responsive protein IAA7		
AT1G30040	-1.5 -4.1	-1.4 -2.5	-0.9		Gibberellin 2-oxidase	CA degradation	
AT5G15230	-3.4	-3.8	-0.9	-0.5	GASA4 / Gibberellin-regulated protein 4	GA degradation	
AT1G74670 AT1G22690	2.0	-0.4		-3.0	GASA6 / Gibberellin-regulated protein 6 GASA9 / Gibberellin-regulated protein 9 (GASA9)	GA signaling	
AT 1G22690 AT5G14920	-0.5	-2.7	-0.6	-2.8	GASA14 / Gibberellin-regulated protein 14		
AT5G67030	-1.1	-2.1	-0.0	1.6	Zeaxanthin epoxidase	ABA synthesis	
AT5G39610	-1.2	0.6		2.7	NAC domain containing protein 92	ADA SYTILICSIS	
AT1G69490	-2.6	0.7		2.7	NAC-like, activated by AP3/PI		
AT3G29035	1.3	2.0	0.6	1.9	NAC domain containing protein 3	Senescence associated	
AT1G66330	-0.5		0.4	1.1	Senescence-associated protein AAF/ ethylene signalin	a	
AT3G07650	3.3	-1.6		-4.8	COL9 / CONSTANS-like 9	•	
AT2G40080	3.3	-0.6		-2.9	ELF4 / EARLYFLOWERING 4	Circadian clock/ Flowering	
log,[Fold Change]							
	oprene suppressed	Isoprene suppressed		fect of senescence	4.00	4.00	
(D) '	A1 healthy leaves 60 DAE	RA1 senescing leave 90 DAE		isoprene-emitting NT leaves 90 DAE			
(re	ative to WT 60 DAE)	(relative to WT 90 DA		ative to WT 60 DA	_{E)} Populus x canescens		
GENE ID	-6.1	-3.6		-1.1	Isoprene synthase		
POPTR 0002s02560g		2.0		-3.2	LOG1 / CK riboside 5'-mpp ribohydrolase 1	Out-libration in the	
POPTR_0006s12920g		2.3		-3.5	LOG3 / CK riboside 5'-mpp ribohydrolase 3	Cytokinin synthesis	
POPTR 0002s031900		2.5		-4.1	CKX5 / Cytokinin oxidase/dehydrogenase 5	0.11111111111	
POPTR 0006s23760g		3.4		-3.8	CKX7 / Cytokinin oxidase/dehydrogenase 7	Cytokinin degradation	
POPTR 008G193000	-0.9	1.5		-2.9	APRR2 / Response regulator		
POPTR 001G027000		0.7		-1.5	APRR5 / Response regulator		
POPTR 016G038000	-1.3	2.3		-2.0	APRR7 / Response regulator	Cytokinin signaling	
POPTR 008G181000	-0.5	1.8		-1.9	APRR14 / Response regulator		
POPTR 018G091401	-1.7	1.8		-2.5	RuBisCO SSU		
POPTR 016G079600	-1.9	3.3		-3.7	Carbonic anhydrase	Photosynthesis / Glycolysis	
POPTR 008G179300	-3.0	1.9		0.5	Glyceraldehyde 3-phosphate dehydrogenase (GAPDH)	,	
POPTR 012G031100	-1.9	-0.5		0.4		Oblamata at market "	
POPTR 015G022100		-1.1		1.0	SIGMA factors	Chloroplast metabolism	
POPTR 001G023500	1.5	2.8		1.5	SCARECROW-LIKE3 (GRAS domain)	CA simulian	
POPTR_010G110700	0.5	2.2		-2.3	DELLA (GRAS domain)	GA signaling	
POPTR_007G121200	2.3	-1.0		1.2	COL9 / CONSTANS-like 9	Circodian alask/ Flavorin-	
POPTR 006G233800	3.5	-1.4		1.3	ELES / EARLY ELOWERING 3	Circadian clock/ Flowering	

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ELF3 / EARLY FLOWERING 3
Heat shock proteins (HSP20, HSP70, HSP90, N=33)

-1.4 1.9

Absence or suppression of isoprene

Decreased levels of leaf cytokinins

Heightened activity of heat-shock transcriptional elements even under stress-free conditions

Significant changes in photoperiodic and circadian clock elements, NAC, WRKY elements and Ethylene response factors

> Slow growth, weak apical dominance, Long sylleptic branching (perennials)

Delayed leaf senescence

Delayed body maturation Delayed flowering (annuals) Deceleration of organismal senescence

Compromised performance under transient stresses Greater survival probability under long-term stresses

In presence of isoprene

Enhanced synthesis and degradation of cytokinins Enhanced cytokinin signalling and associated transcriptomic changes under stress-free conditions

Increased rate of chloroplast metabolic activities

Attenuation of photoinhibition (in shade leaves)

Strong apical dominance (perennials) Short sylleptic branches

Early and faster leaf senescece independent of growth rate and biomass (sink) constraints

Early flowering and organismal senescence (annuals)

Greater tolerance to shoter-term abiotic stresses Greater mortality under severe and sustained stresses

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