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2 **Leaf isoprene emission as a trait that mediates the growth-defense tradeoff in the face of**
3 **climate stress**

4 Russell K. Monson¹, Sarathi M. Weraduwage^{2,3}, Maaria Rosenkranz⁴, Jörg-Peter Schnitzler⁴,
5 Thomas D. Sharkey^{2,3,5}

6 ¹ Department of Ecology and Evolutionary Biology, University of Colorado, Boulder, CO 80309
7 USA

8 ² Department of Biochemistry and Molecular Biology, MSU-DOE Plant Research Laboratory,
9 East Lansing, MI 48824 USA

10 ³ Department of Energy, Office of Science, Office of Biological and Environmental Research,
11 Great Lakes Bioenergy Research Center, East Lansing, MI 48824 USA

12 ⁴ Research Unit Environmental Simulation, Institute of Biochemical Plant Pathology, Helmholtz
13 Zentrum München, 85764 Neuherberg, Germany

14 ⁵ Plant Resilience Institute, Michigan State University, East Lansing, MI 48824 USA

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18 Corresponding author: Russell Monson; russell.monson@colorado.edu

19

20 **Abstract** (244 words)

21 Plant isoprene emissions are known to contribute to abiotic stress tolerance, especially during
22 episodes of high temperature and drought, and during cellular oxidative stress. Recent studies
23 have shown that genetic transformations to add or remove isoprene emissions cause a cascade of
24 cellular modifications that include known signaling pathways, and interact to remodel adaptive
25 growth-defense tradeoffs. The most compelling evidence for isoprene signaling is found in the
26 shikimate and phenylpropanoid pathways, which produce salicylic acid, alkaloids, tannins,
27 anthocyanins, flavonols and other flavonoids; all of which have roles in stress tolerance and plant
28 defense. Isoprene also influences key gene expression patterns in the terpenoid biosynthetic
29 pathways, and the jasmonic acid, gibberellic acid and cytokinin signaling networks that have
30 important roles in controlling inducible defense responses and influencing plant growth and
31 development, particularly following defoliation. In this synthesis paper, using past studies of
32 transgenic poplar, tobacco and *Arabidopsis*, we present the evidence for isoprene acting as a
33 metabolite that coordinates aspects of cellular signaling, resulting in enhanced chemical defense
34 during periods of climate stress, while minimizing costs to growth. This perspective represents a
35 major shift in our thinking away from direct effects of isoprene, for example, by changing
36 membrane properties or quenching ROS, to indirect effects, through changes in gene expression
37 and protein abundances. Recognition of isoprene's role in the growth-defense tradeoff provides
38 new perspectives on evolution of the trait, its contribution to plant adaptation and resilience, and
39 the ecological niches in which it is most effective.

40 **Keywords** ozone • green leaf volatiles • carotenoids • chlorophyll • thermotolerance • drought •
41 photosynthetic capacity • phytohormones • ROS • growth differentiation balance • lignin

42 **Introduction**

43 Biogenic isoprene (2-methyl-1,3-butadiene) is a light-dependent, volatile, hemiterpene
44 emitted from the chloroplasts of many plants, including numerous woody species, and even some
45 ferns and mosses (Harley et al. 1999, Hanson et al. 1999; Monson et al. 2013). The molecule has
46 drawn the attention of plant physiologists, ecologists and atmospheric chemists because: (1) it
47 appears to be an important trait that protects the photosynthetic apparatus of plants in the face of
48 climate stress (Sharkey et al. 2008; Loreto and Schnitzler 2010); (2) it has a role in structuring
49 tritrophic interactions among plants, herbivores and their parasites (Loivamäki et al. 2008); and
50 (3) it has a role in controlling the oxidative state of the troposphere (Monson and Holland 2001,
51 Monson 2002; Pike and Young 2009). One of the most debated issues concerning the topic of
52 isoprene emissions is why plants produce it (Sharkey and Singsaas 1995; Monson et al. 2013;
53 Sharkey 2013; Dani et al. 2014; Loreto and Fineschi 2015). In this synthesis, we take up this
54 issue with a focus on recent data showing that isoprene participates in several cellular signaling
55 networks and has a role in the coordination of growth-defense trait tradeoffs.

56 For nearly thirty years, evidence has accumulated that isoprene protects the photosynthetic
57 apparatus of plants from abiotic stress, such as that caused by high temperature and drought
58 (Sharkey and Singsaas 1995; Sharkey and Yeh 2001; Ryan et al. 2014; Fini et al. 2017; Taylor et
59 al. 2019). Many of these studies relied on the elimination of isoprene emission through the
60 introduction of chemical inhibitors or genetic modification, and they focused on the thermal
61 tolerance of photosynthesis (Sharkey et al. 2001; Velikova and Loreto 2005; Behnke et al. 2007;
62 Sasaki et al. 2007). Other research, using similar approaches, showed that isoprene is effective
63 against cellular oxidative stresses that occur during drought and high light episodes, or ozone
64 exposure (Loreto and Velikova 2001; Affek and Yakir 2002; Velikova et al. 2004; Vickers et al.

65 2009b; Behnke et al. 2009; 2010b; Pollastri et al. 2014). When considered as a whole, the past
66 body of research focuses on properties of the isoprene molecule in an isolated protective role;
67 one in which unsaturated hydrocarbon bonds stabilize protein-lipid and protein-protein
68 interactions in chloroplast thylakoids and/or react directly with reactive oxygen species (ROS)
69 (Sharkey et al. 2008; Siwko et al. 2007; Vickers et al. 2009a; Velikova et al. 2011; Parveen et al.
70 2019a).

71 Even in the face of these numerous reports, however, some past studies failed to observe an
72 effect of isoprene applied as a short-term, single treatment on thermotolerance in isolated leaf
73 discs (Logan and Monson 1999), or on the permeability and stability of isolated thylakoids and
74 liposome membranes (Logan et al. 1999). Recently, Harvey et al. (2015) concluded that the
75 partitioning of gaseous isoprene into phospholipid membranes, at realistic intra-leaf
76 concentrations, was two orders of magnitude lower than levels thought to be effective in thermal
77 protection. Furthermore, they found that even extremely high isoprene concentrations failed to
78 affect the viscosity of phosphatidylcholine liposome membranes. These observations provide
79 opposing evidence to the past theories of isoprene acting alone to stabilize membrane
80 hydrophobic interactions. At the same time, a different set of observations began to emerge that
81 further challenged the adequacy of the traditional theories. Using transgenic technologies, the
82 capacity for isoprene emissions was introduced into otherwise non-emitting species, generating
83 novel phenotypic effects that could not be explained by the conventional theories. For example,
84 increased growth rates occurred in some species (Loivamäki et al. 2007; Sasaki et al. 2007;
85 Vickers et al. 2009b; Zuo et al. 2019), but they decreased in others, even under non-stressful
86 conditions (Zuo et al. 2019). These studies suggested that a complete understanding of isoprene's
87 effects would require a broader theoretical scope. Despite these challenges, the overall body of

88 work from nearly three decades of research has provided unequivocal evidence, and general
89 acceptance, that isoprene emission does indeed represent a trait with positive adaptive value in
90 plants, especially with regard to enhancing photosynthesis in the face of stress. The newer
91 questions that have been raised, are: what is the broader adaptive scope of the trait and how does
92 stress tolerance fit into that broader scope?

93 In recent work, observations have been made that address these questions and provide new
94 hypotheses with regard to isoprene's mode of action. Rather than acting alone, isoprene is more
95 likely to interact with several other metabolites known to protect the photosynthetic apparatus
96 during stress (Loreto and Schnitzler 2010; Behnke et al. 2010a; Tattini et al. 2014; Velikova et
97 al. 2014; Brunetti et al. 2015; Tattini et al. 2015; Vanzo et al. 2016; Marino et al. 2017; Lantz et
98 al. 2019; Zuo et al. 2019; Parveen et al. 2019b; Monson et al. 2020; Liu et al. 2020). Much of the
99 new evidence has been obtained employing plants modified in their isoprene emission capability;
100 natural isoprene-emitters (IE) that were transformed to be non-emitters (NE) by knocking-down
101 expression of the isoprene synthase gene (*ISPS*) (Behnke et al. 2007), and conversely, NE plants
102 that were converted to IE plants by introducing *ISPS* (Loivamäki et al. 2007; Sasaki et al. 2007;
103 Vickers et al. 2009b; Zuo et al. 2019). A picture is emerging of isoprene acting within a broad
104 cellular network to both contribute to abiotic stress tolerance and organize aspects of growth-
105 defense trait tradeoffs. In this synthesis, we bring together these emerging concepts on cellular
106 signaling, growth and abiotic stress tolerance, and through an integration with known relations
107 between metabolite production and plant defense, provide a new hypothesis on isoprene's
108 potential role in mediating the growth-defense tradeoff. In essence, we make the case that
109 isoprene has evolved in certain plant lineages as a means to stage an effective form of chemical
110 defense, with minimal costs to growth, in the face of climate stress.

111 **The growth defense tradeoff**

112 Plant growth rate, whether considered in physiological or evolutionary terms, reflects
113 resource limitations and the combined contributions and interactions of multiple traits as they
114 acquire and use limited resources. Patterns in resource use and allocation can be described
115 according to economic tradeoffs (Mooney 1972; Chapin 1980; Bloom et al. 1985; Reich 2014);
116 whereby, utilization of limiting resources for one function must come at a cost to their utilization
117 for a competing function. The realization of common currencies and interdependencies, within
118 the context of plant resource use, lies at the foundation of growth-defense tradeoff theory – the
119 mandated relation by which allocation to growth in a resource-limited environment occurs at a
120 cost to defense, and vice versa (Stamp 2003; Schuman and Baldwin 2016; Züst and Agrawal
121 2017). Recognition of the utility of an economic framework, within which to describe growth,
122 defense and fitness, has led to a broad foundation on which to build concepts of adaptation,
123 trophic interactions and evolutionary compromises.

124 The principal ecological theories underlying growth-defense tradeoffs contain some aspect
125 of cost and benefit. Bryant et al. (1983) conducted studies in boreal ecosystems and observed
126 that plants native to environments with low nutrient availability, such as black spruce (*Picea*
127 *mariana*) and various graminoid species, had slow growth rates, utilized carbon-based defenses
128 and had high levels of defense at all life-cycle phases. Plants native to sites with higher nutrient
129 levels, such as quaking aspen (*Populus tremuloides*) and several dicot herbaceous species,
130 exhibited faster growth rates, had high levels of defense only as juveniles and, in many species,
131 relied more often on nitrogen-based defenses. These observations were synthesized to produce
132 the Carbon-Nutrient Balance (CNB) hypothesis, which has been shown to be most useful in
133 addressing physiological plasticity, and the movement of carbon or nitrogen in excess of that

134 required for growth, to defense (Lerdau and Coley 2002; Stamp 2003). Coley et al. (1985)
135 expanded the principals of the CNB hypothesis to account for genetically fixed G-D tradeoffs
136 and interspecific differences in adaptation. In formulating the Resource Availability Hypothesis
137 (RAH), they presented the case that the G-D tradeoff reflects selection across a broad range of
138 traits in plants native to environments with different levels of resource availability. According to
139 the RAH, slower growth rates, but higher levels of constitutive defense, occur in plants native to
140 resource-limited environments, compared to plants native to resource-rich environments, and
141 these patterns of trait covariance maximize fitness. In the Growth-Differentiation Balance
142 Hypothesis (GDBH), Herms and Mattson (1992) developed the physiological case for a
143 mutually-exclusive allocation of resources to one or the other function – growth versus
144 differentiation (including defense). The GDBH included the tenets of the CNB hypothesis, but
145 expanded the arguments to include ontogenetic constraints and it framed growth-differentiation
146 tradeoffs in terms of cellular processes. Herms and Mattson (1992) applied their physiological
147 hypotheses to an evolutionary optimality model in an effort to link the processes of cellular
148 tradeoffs with natural selection, and therefore integrate the RAH and GDBH. The Optimal
149 Defense Theory (ODT), which predates the tradeoff theories discussed above, emerged from
150 studies showing that plants deploy their chemical defenses in ways that maximize their
151 effectiveness against herbivores (McKey 1974; Rhoades 1979). These observations led to a
152 general theory that plants have evolved patterns of defense in direct proportion to levels of
153 herbivory and impacts on fitness. The ODT differs from the RAH and GDBH in that the
154 principal determinant in natural selection is the balance between demand and fitness cost, rather
155 than growth and fitness cost.

156 Tradeoff theories that focus on selection and genetic correlations among traits, such as the
157 RAH, GDBH and ODT, have generally worked well to predict patterns of growth and defense at
158 higher taxonomic levels (Endara and Coley 2011; Schuman and Baldwin 2016; Züst and
159 Agrawal 2017). However, they often fail in predicting tradeoffs within species (Agrawal 2020).
160 With regard to intraspecific processes, it is common to find increases in both growth and defense
161 as resource availability increases (van Nordwijk and de Jong 1986; Agrawal 2020). This is due
162 to phenotypic plasticity – those individuals within a species with greater access to limiting
163 resources will allocate greater amounts of those resources to both growth and defense, compared
164 to individuals with lesser access to resources. Thus, in comparing individuals within a species, a
165 transition from high-to-low resource availability may not result in the prioritization of defense
166 over growth, as predicted by the optimization theories. This crucial concept is key to
167 understanding the selective forces at play when a novel trait, such as isoprene emission, is
168 introduced into a species. If that trait facilitates greater resource acquisition, and has the potential
169 to direct the allocation of those resources to both growth and defense, then the selective value of
170 the trait to both processes could increase.

171 In the past decade, progress has been made, using multi-omic approaches, applied primarily
172 to *Arabidopsis*, within the context of growth-defense tradeoffs (D'Auria and Gershenzon 2005;
173 Campos et al. 2016; Züst and Agrawal 2017). At the molecular scale, growth-defense tradeoffs
174 are orchestrated through cellular signaling networks and enacted, to a large extent, through
175 interactions among genetic transcription factors (TFs) (Karasov et al. 2017). These interactions
176 control gene expression in the pathways of both constitutive and induced defenses, and are
177 coupled through pathway crosstalk to the production and sensitivities of growth regulators, such

178 as salicylic acid, jasmonic acid, gibberellins, cytokinins, ethylene and abscisic acid (Howe et al.
179 2018; Guo et al. 2018a; Koo et al. 2020).

180 Transcriptional control of gene expression determines the enzymatic potential for increases
181 or decreases in metabolite production (Schuman and Baldwin 2016; Züst and Agrawal 2017),
182 and along with enzyme kinetics and substrate availability establishes the antagonistic push or
183 pull 'forces' that determine whole-plant growth-defense allocation patterns. Transcriptional
184 control can also provide resilience in the face of allocation commitments in a variable
185 environment. In the theoretical condition of a zero-sum constraint, there is no margin for error in
186 the face of environmental fluctuation, and mistakes in allocation strategy can lead to reduced
187 fitness. Transcriptional control provides a means for selection to favor phenotypes that operate
188 conservatively, below the zero-sum constraint, and retain a resource margin that can be used for
189 adjustments in the growth-defense balance, leading to increased plant resilience in the face of
190 episodic stress (Guo et al 2018b).

191 **Isoprene as a broad modulator of gene expression and pathway interactions**

192 Much of the evidence for a broader cellular role for isoprene has come through the
193 application of multi-omic observations of isoprene-altered phenotypes, using poplar (primarily
194 the hybrid *Populus x canescens*), tobacco (*Nicotiana tabacum* L.) and *Arabidopsis thaliana*.
195 Each species offers advantages, and together they provide insight across a range of phenotypes
196 and growth forms reflecting different evolutionary histories. Poplar is a woody, IE species,
197 providing an opportunity to study molecular and biochemical interactions when isoprene is
198 present as a native trait. *Arabidopsis* and tobacco are herbaceous, NE species. While *Arabidopsis*
199 follows a monocarpic annual life cycle, tobacco is a polycarpic perennial plant in its native or
200 invasive forms (see Ren and Timko 2001; Jassbi et al. 2017), though it is often cultivated on an

201 annual rotation. The inclusion of *Arabidopsis* and tobacco as non-emitting species in the wild
202 type, allowed us to study how the introduction of isoprene interacts with 'naïve' metabolism,
203 similar to what happens following the initial evolutionary introduction of this trait into a
204 population. In the *Arabidopsis* system we used transcriptomic results from: (1) a fumigation
205 treatment intended to explore the direct, targeted role of isoprene as a signal, and (2) transgenic
206 transfer of *ISPS* to explore the role of isoprene as an integrated, permanent component of the
207 plant genotype. In the following sections, we consider patterns of cellular adjustment in key
208 pathways and signaling networks in all three of these species focusing on those most relevant to
209 the growth-defense tradeoff.

210 *Isoprene and the shikimate/phenylpropanoid pathways*

211 In early work, RNA interference (RNAi) was used to achieve translational inhibition of
212 *ISPS* in poplar trees (Behnke et al. 2007). Transcriptomic and metabolomic analyses showed that
213 RNAi silencing of *ISPS* also reduced gene expression in the shikimate and associated
214 phenylpropanoid pathways, indicating a crucial role for isoprene as a positive regulator of
215 pathway expression patterns (Behnke et al. 2010a). In a recent proteome study, also using RNAi
216 poplar lines, growth in an experimental plantation in Arizona confirmed that transgenic
217 suppression of *ISPS* caused reductions in multiple proteins associated with both pathways
218 (Monson et al. 2020). When combined, these studies showed that at least fifteen genes/proteins
219 in the shikimate and phenylpropanoid pathways were expressed at higher levels, and one protein
220 was expressed at a lower level in IE poplar trees, compared to transgenic NE poplar (Fig. 1).

221 There is uncertainty about the significance of changes in one isoform of cinnamyl alcohol
222 dehydrogenase (CAD9; Fig. 1), which participates in the final step in the synthesis of lignin
223 monomers, and was present at 50% lower levels in IE trees, in the study of Monson et al. (2020).

224 The gene, *CAD9*, belongs to a family of 15 genes in poplar, and specifically to a sub-group of
225 leaf *CAD* genes with primary roles in plant defense (Barakat et al. 2009). At this time, the
226 potential advantage of 50% less *CAD9* protein in the presence of isoprene, is not clear. One
227 hypothesis is that its expression of the gene for this protein is suppressed in the presence of
228 isoprene and that this adjustment balances resource flow from one type of less advantageous
229 defense metabolite (e.g., lignins) to a different, more effective, defense metabolite (e.g., phenolic
230 glycosides). Transcripts for a second member of the same gene family (*CAD1*) were observed to
231 be higher in IE poplar, compared to NE poplar in the study of Behnke et al. (2010a). *CAD1* is
232 associated with monolignol production in xylem tissues, where it forms a multi-protein complex
233 with cinnamoyl CoA-reductase (CCR) (Yan et al. 2019). In past studies using RNAi to minimize
234 CCR in poplar, a 50% reduction in lignin production, along with *CAD1*, was observed (Li et al.
235 2014), and temperature-sensitive dwarfed growth has been observed in alfalfa above 30 °C in
236 *CAD1* mutants (Zhao et al. 2013). Thus, we hypothesize that there is an upregulation of *CAD1* in
237 the presence of isoprene, and that this might improve growth, especially in warm habitats or in
238 the face of sustained high temperature stress.

239 Increases in phenylpropanoid pathway gene expression have also been observed in tobacco
240 and *Arabidopsis* (Tattini et al. 2014; Zuo et al. 2019). In wild-type *Arabidopsis* exposed to
241 isoprene (Harvey and Sharkey, 2016), expression of genes encoding the first six enzymes of the
242 phenylpropanoid pathway, along with the later-pathway enzyme, flavonol 3-O-glucosyl
243 transferase (3GT), were upregulated. The first step of the pathway is catalyzed by phenylalanine
244 ammonia-lyase (PAL), which is a family of four isoforms coded by genes, *PAL1 – PAL4* (Raes
245 et al. 2003). *PAL1* and *PAL2* encode the phenylpropanoid forms in *Arabidopsis* (Fraser and
246 Chapelle 2011), and these are two of the forms with increased expression in *Arabidopsis* exposed

247 to isoprene, along with *PAL4* (Harvey and Sharkey 2016); *PAL1* expression was also increased
248 in transformed IE tobacco (Zuo et al. 2019). In transformed IE tobacco and *Arabidopsis*, and
249 wild-type *Arabidopsis* fumigated with isoprene, expression of *calcium dependent protein kinase*
250 *1* (*CPKI*) that phosphorylates and activates PAL protein, increased, while in the two *Arabidopsis*
251 systems, the expression of the *Kelch repeat F-box 20 protein* (*KFB20*), which mediates PAL
252 degradation, decreased (Harvey and Sharkey 2016; Zuo et al. 2019). Thus, there is strong
253 validation of isoprene effects that enhance phenylpropanoid expression in species beyond poplar,
254 indicating generality in this signaling system.

255 *Isoprene and the terpenoid pathway*

256 Isoprene is produced from dimethylallyl diphosphate (DMADP), a product of the
257 methylerythritol 4-phosphate (MEP) pathway in chloroplasts. MEP pathway flux begins with the
258 substrates glyceraldehyde 3-phosphate (GAP) and pyruvate (Pyr) from the reductive pentose
259 phosphate (RPP) pathway (the photosynthetic Calvin-Benson cycle). The amount of DMADP
260 available for isoprene production is not only sensitive to upstream flux through the MEP
261 pathway, but also its downstream utilization for the production of higher terpenoids, such as the
262 carotenoids, gibberellic acid, cytokinins, tocopherols, the phytol tail of chlorophylls, abscisic
263 acid and an array of monoterpenes and sesquiterpenes.

264 The presence of isoprene in cells controls metabolite flows through alternative branches of
265 the terpenoid pathway, and there are clear interspecific differences. In IE poplar and tobacco, in
266 contrast to the situation for the shikimate and phenylpropanoid pathways, the presence of
267 isoprene caused reductions in multiple steps in the expression of terpene biosynthetic genes and
268 proteins (Fig. 2; Monson et al. 2020; Zuo et al. 2019). In *Arabidopsis* wild-type plants exposed to

269 isoprene, increases, rather than decreases in transcript abundances were observed, especially
270 concerning genes related to carotenoid biosynthesis.

271 One novel observation in poplar involves a set of genes encoding rubber elongation factors
272 (*REFs*; see #1 in Figure 2). The abundance of two different REF proteins was decreased in the
273 presence of isoprene (Monson et al. 2020). REF proteins are typically found in rubber-producing
274 plants, such as rubber tree (*Hevea brasiliensis*) and guayule (*Parthenium argentatum*) (Lau et al.
275 2016). Rubber elongation is catalyzed by membrane-bound complexes containing *cis*-
276 prenyltransferase (CPT) enzymes. At least five different CPT genes have been reported in
277 *Populus trichocarpa*, in which they have roles in the synthesis of tetra- (C₄₀), di- (C₂₀) and
278 sesquiterpenes (C₁₅). It is possible that the REF proteins detected in poplar are part of a larger
279 family of proteins with prenyltransferase roles (i.e., lipid-droplet associated proteins, LDAPs;
280 Gidda et al. 2013), including those associated with the cellular compartmentation and storage of
281 isoprenoid compounds. In leaves, LDAPs may be involved with cellular energy balance, cellular
282 signaling and plant stress responses (van der Schoot et al. 2011, Walther and Farese, 2012).
283 Thus, in poplar, isoprene may contribute to coordination between photosynthetic capacity and
284 lipid energy storage and/or lipid-based stress responses. The channeling of photosynthate to
285 growth during periods of high photosynthetic capacity (and high rates of isoprene synthesis), but
286 storage or stress tolerance during periods of low photosynthetic capacity (and low rates of
287 isoprene synthesis), would be consistent with observations of *REF* gene suppression in the
288 presence of isoprene. The situation appears to be different in wild-type *Arabidopsis*, as *REF* gene
289 expression is increased in response to fumigation with isoprene (Harvey and Sharkey, 2016).

290 In IE poplar, we observed a reduction in the amount of protein for tocopherol cyclase
291 (VTE1) (Monson et al. 2020; see #2 in Figure 2). The observation of reduced VTE1 levels is

292 consistent with past studies that showed reduced levels of the associated metabolite, α -
293 tocopherol, in IE poplar leaves (Behnke et al. 2009), and heat-stressed Holm oak (*Quercus ilex*)
294 leaves fumigated with isoprene (Peñuelas et al. 2005). α -tocopherol is an important antioxidant
295 that is active in reducing cellular ROS during physiological stress. In wild-type *Arabidopsis*
296 plants exposed to isoprene, *VTE1* gene transcripts were increased, as well as those for tocopherol
297 methyltransferase (*TMT*) (Harvey and Sharkey 2016), once again, showing the tendency for
298 interspecific differences in terpenoid pathway effects, especially between poplar and
299 *Arabidopsis*.

300 The proteins for z-carotene desaturase (*ZDS1*), carotene ϵ monooxygenase (*CYP97C1*), and
301 zeaxanthin epoxidase (*ZEP*), which catalyze key steps in the flow of GGPP toward carotenoid
302 and ABA biosynthesis, were at lower abundances in IE poplar lines, compared to NE lines
303 (Monson et al. 2020; see #3 in Fig. 2). These patterns were similar in the transcriptomic studies
304 in tobacco, in which transcript numbers for genes of several enzymes involved in the channeling
305 of GGPP to carotenoid synthesis were reduced, including those for phytoene synthase (*PSY*),
306 phytoene desaturase (*PSD*), z-carotene desaturase (*ZDS1*), carotenoid isomerase (*CRTISO*), β -
307 cyclase (*LCY1*) and zeaxanthin epoxidase (*ZEP*; Zuo et al. 2019). A few past studies on empty-
308 vector IE and transgenic NE poplar trees have shown variable results on leaf concentrations of
309 carotenoids and the non-stressed deepoxidation state. Behnke et al. (2010b) observed no
310 significant differences in carotenoid amount or epoxidation status in non-stressed IE and NE
311 poplars. However, in a different study, Behnke et al. (2009) observed reduced amounts of
312 zeaxanthin, but a higher deepoxidation ratio in IE poplars, which is consistent with the proteomic
313 analyses that show reduced expression in *ZDS1* (decreasing the pool of zeaxanthin) and *ZEP*
314 (increasing the deepoxidation ratio) (Fig. 2).

315 In contrast to the poplar and tobacco systems, transcripts of most carotenoid-related
316 enzymes were significantly increased in wild-type *Arabidopsis* exposed to isoprene (Harvey and
317 Sharkey 2016; Fig. 2). In addition to upregulation in many of the same proteins that were
318 described above, transcripts for the gene encoding β -carotene hydroxylase (*crtZ*) were increased
319 in wild-type *Arabidopsis* exposed to isoprene. The protein, *crtZ*, converts β -carotene to
320 zeaxanthin and can control the size of the xanthophyll cycle pool, and overexpression of *crtZ* in
321 *Arabidopsis* has been linked to high-light and high-temperature stress tolerance (Davison et al.
322 2002). Finally, expression of the gene for the enzyme violaxanthin deepoxidase (*VDE*), which
323 converts violaxanthin to zeaxanthin in the flexible, photoprotective part of the carotenoid cycle,
324 was increased in wild-type *Arabidopsis* exposed to isoprene (Harvey and Sharkey 2016; Zuo et
325 al. 2019).

326 A second component of the antioxidant system in plants, ascorbate, also appears to be
327 differentially expressed in response to isoprene in poplar versus *Arabidopsis*.
328 In poplar, the enzyme ascorbate oxidase (AAO), which leads to a lower ascorbate content in
329 leaves, is present at a 44% higher level in IE lines, compared to NE lines (Monson et al. 2020).
330 This observation is consistent with past observations showing lower ascorbate contents in non-
331 stressed IE poplar (Behnke et al 2009). In wild-type *Arabidopsis* exposed to isoprene (Harvey
332 and Sharkey 2016; Zuo et al. 2019), two genes for the protein GDP-galactose phosphorylase
333 (*VTC2* and *5*), which has been shown to be the only significant pathway in *Arabidopsis* that
334 produces ascorbate (Dowdle et al. 2007), exhibited increased expression.

335 In poplar, there were no isoprene-associated shifts in gene expression or protein contents
336 related to the synthesis of abscisic acid (ABA) (Behnke et al. 2010a; Monson et al. 2020).
337 However, in transformed *Arabidopsis* and wild-type *Arabidopsis* exposed to isoprene, expression

338 of the chloroplast gene that encodes an isoform of 9-cis-epoxycarotenoid dioxygenase (NCED),
339 which converts violaxanthin to ABA, was upregulated (Fig. 2). Two isoforms, NCED3 and
340 NCED5, have been shown to have important roles in plant growth and drought tolerance (Frey et
341 al. 2012). In both *Arabidopsis* systems and transgenic tobacco, transcription of *ATAF1*, a key TF
342 that positively regulates *NCED3* (Jensen et al. 2013), was also increased. This observation was
343 of particular interest for transgenic tobacco, given that the general expression of carotenoid genes
344 was downregulated by the introduction of isoprene in this species. The results suggest that
345 isoprene-mediated upregulation of the ABA component of the carotenoid pathway is controlled
346 independently from factors that control the remainder of the pathway.

347 Finally, in poplar, it was observed that isoprene synthesis reduces production of the enzyme
348 that converts *cis*-zeatin, the low-activity form of cytokinin, to a stabilized, metabolically-inert
349 pool (Monson et al. 2020). This action sustains a pool of *cis*-zeatin for isomerization to *trans*-
350 zeatin, the most active form of cytokinin (see #4 in Fig. 2). Signaling through *trans*-zeatin has
351 been implicated in controls over plant re-growth and shifts in carbohydrate source-sink balance
352 that enhance photosynthetic capacity following partial plant defoliation (Roitsch and Ehneß
353 2000; Glanz-Idan et al. 2020).

354 There was a clear pattern of species differences in the effects of isoprene on gene
355 expression in the terpenoid pathway. Poplar and tobacco tend to downregulate several pathway
356 steps in the presence of isoprene, whereas *Arabidopsis* upregulates them (in both the fumigation
357 and transgenic treatments). It is most likely that these interspecific differences are due to past
358 selection for different adaptive priorities in plant responses to abiotic stress. In poplar and
359 tobacco, selection to increase allocation to shikimate and phenylpropanoid production, at the
360 expense of carotenoid production, might have occurred in response to high levels of herbivory

361 during past selection episodes. [Although tobacco, *Nicotiana tabacum* L., is cultivated on an
362 annual rotation, it grows naturally as a perennial and was likely derived as a natural amphidiploid
363 hybrid with genetic contributions from three perennial ancestors (Ren and Timko 2001). The
364 native perennial nature of tobacco likely explains its tendency to allocate a relatively high
365 amount of resource to the production of defensive metabolites.] In contrast, in *Arabidopsis*, an
366 annual plant with relatively high growth rates and native affinities for open habitats with well-
367 drained soils, photoprotection, administered through an effective antioxidant system, might have
368 carried a higher selective value, at the expense of a well-provisioned chemical defense system.
369 Given these pre-existing differences in phenotype, the introduction of isoprene, either through
370 evolution as in the case of poplar, or through transgenic introduction as in tobacco and
371 *Arabidopsis*, would be differentially integrated into existing signaling systems. Thus, the pre-
372 existing metabolic phenotype might be as important as the properties of isoprene itself in
373 determining its role as a signal modulator.

374 *Isoprene and the oxylipin pathways*

375 The oxylipin pathway is initiated in the chloroplast and produces C-6 aldehydes, alcohols
376 and esters, known as green leaf volatiles (GLVs) (Hatanaka et al. 1987), and the jasmonic acid
377 (JA) pathway precursor, 13(S)-hydroperoxy-octadecatrienoic acid (13-HPOT). Chloroplast-
378 derived oxylipins are produced from C18-polyunsaturated fatty acids (Matsui 2006), which are
379 freed from membranes by a family of phospholipases in response to herbivory, pathogen
380 infection or abiotic stress (Ameye et al. 2018; Liu et al. 2020). Once freed, the fatty acids are
381 oxidized by a family of lipoxygenase (LOX) enzymes, which control the channeling of oxylipins
382 into several wound- and defense-associated pathways, some of which involve other organelles,
383 such as peroxisomes (Feussner and Wasternack 2002; Koo 2018).

384 The presence of isoprene increases expression in several *LOX* genes, and thus the potential
385 for GLV production, in all of the experimental systems used in this analysis (Behnke et al.
386 2010a; Harvey and Sharkey, 2016; Zuo et al. 2019; Monson et al. 2020) (Fig. 3). Beyond this
387 initial step, however, isoprene effects on oxylipin processes among species begin to diverge,
388 especially with respect to JA signaling.

389 The active form of JA occurs when it forms a macromolecular complex with the amino
390 acid, isoleucine (Ile). A principal receptor of the JA-Ile conjugate is the Coronatine Insensitive 1
391 (COI1) F-box protein, which is part of an E3 ubiquitin ligase complex. When JA-Ile reaches a
392 threshold level, COI1 associates with JA-Ile to form a multi-protein transcriptional modifier
393 (Ruan et al. 2019). The JA-Ile-COI1 complex targets a large family of proteins known as
394 Jasmonate ZIM (JAZ), which generally function as negative transcriptional regulators (Fig. 3). In
395 the absence of JA-Ile-COI1, JAZ proteins bind to TFs with a high degree of specificity, blocking
396 their ability to participate in the formation of transcription initiation complexes. The COI1-JAZ
397 interaction triggers ubiquitin-dependent degradation of the JAZ proteins, freeing the TFs and
398 activating transcription.

399 From the past studies in *Arabidopsis*, there are antagonistic interactions involving proteins
400 in the JAZ and DELLA families, which are associated with the JA and gibberellic acid (GA)
401 pathways, respectively. JA synthesis leads to an enhancement in DELLA transcription and
402 degradation of JAZ proteins, which leads to decreased growth and increased defense,
403 respectively – in essence, providing a molecular context for the growth-defense tradeoff; in this
404 case, favoring defense (Wild et al. 2012; Campos et al. 2016). GA synthesis leads to degradation
405 of DELLA proteins and an enhancement of GA-activator signaling, which enhances growth and,
406 at the same time, increases the number of free JAZ proteins, which suppress JA-associated

407 defense signaling (Campos et al. 2016). Once again, these molecular interactions enable the
408 growth-defense tradeoff; but, in this case, favoring growth. Antagonism between JA and GA
409 signaling in determining the growth-defense tradeoff is also influenced by extrinsic
410 environmental cues involving spectral shifts in incident light (Ballarè 2014), and it can be
411 uncoupled by phytochrome b gene mutations (Campos et al. 2016).

412 The expression of three genes important for JA synthesis were increased by isoprene in
413 *Arabidopsis* and tobacco. For example, in addition to LOX, 12-oxo-phytodienoic acid reductase
414 (*OPR*) was enhanced in transgenic IE *Arabidopsis* and IE tobacco, while 3-oxo-2-(2'-[Z]-
415 pentenyl) cyclopentane-1-octanoate CoA ligase (*OPC-8:CoA* ligase) increased in IE tobacco. In
416 addition, expression of the gene for MYB59 transcription factor was reduced in both the
417 fumigated and transgenic *Arabidopsis* systems in response to isoprene (Harvey and Sharkey
418 2016; Zuo et al. 2019). MYB59 suppresses expression of certain genes important for oxylipin
419 biosynthesis (e.g., the gene for allene oxide synthase, *CYP74A*, and *OPR3*), as well as the gene
420 for the MYC2 TF that regulates JA signaling (Boter et al. 2004). Expression of the gene for
421 jasmonate O-methyltransferase (*JMTI*) was increased in wild-type *Arabidopsis* exposed to
422 isoprene. JMT catalyzes the reaction leading to methyl jasmonate (Me-JA), a volatile
423 hydrocarbon that can function as a long distance chemical signal with significant effects on JA
424 related gene expression (Benevenuto et al. 2019). Both Me-JA and isoprene have been shown to
425 be produced in response to wounding (Loreto and Sharkey 1993; Lantz et al. 2019; Benevenuto
426 et al. 2019); a response that seemed out of place with the past literature emphasizing a role for
427 isoprene in abiotic stress tolerance, but that now makes more sense within the context of a multi-
428 pathway signaling network. In addition to an isoprene-induced increase in *JMTI* expression in
429 wild-type *Arabidopsis*, two negative regulators of *JMTI* expression (*BBD1* and *BBD2*; Seo et al.

430 2013) exhibited reduced transcript levels in transgenic IE *Arabidopsis* and wild-type *Arabidopsis*
431 exposed to isoprene (Zuo et al. 2019). In IE tobacco, there was evidence for reduced jasmonate-
432 amido synthase (*JARI*), expression which catalyzes the synthesis of JA-Ile. Thus, based on
433 transcriptomic data, it seems that isoprene favors production of volatile Me-JA over JA-Ile, at
434 least in *Arabidopsis* and tobacco. We are just beginning to sort the ways that isoprene interacts
435 with JA and its derivatives in determining growth-defense tradeoffs; though, it is clear that much
436 of the insight concerning molecular control over the tradeoff is to be found in JA and GA
437 pathway interactions (see Züst and Agrawal 2017).

438 Unlike the case for IE tobacco, in IE poplar lines Monson *et al.* (2020) observed a 40%
439 decrease of the JA pathway protein, 3-oxo-2-(2'-[Z]-pentenyl) cyclopentane-1-octanoate CoA
440 ligase (OPC-8:CoA ligase), compared to the transgenic NE lines (Fig. 3). Such a large
441 downregulation of this 'gatekeeper' enzyme would likely lead to substantial decreases in the
442 capacity for JA signaling. In poplar, there might be advantages to suppressing JA signaling in the
443 presence of isoprene. Two of the JA signaling processes that are activated when JAZ proteins are
444 degraded involve upregulation of the phenylpropanoid pathway (Zhou et al. 2017) and
445 downregulation of the salicylic acid (SA) pathway (Campos et al. 2014). In poplar, suppression
446 of JA-Ile formation might provide a means to sustain JAZ-mediated suppression of
447 phenylpropanoid biosynthesis, thus avoiding conflicts between JA- and isoprene-regulated
448 controls in the same pathway. The ability to isolate control over phenylpropanoid biosynthesis
449 from the several other growth and defense interactions affected by JA signaling might also be
450 especially important for poplar. For example, SA signaling has an important role in plant defense
451 that is distinct from, and in several cases antagonistic to, JA signaling (Durrant and Dong 2004).
452 A suppression of JA signaling allows for sustained JAZ-mediated suppression of those TFs that

453 might otherwise negatively influence SA signaling (Fig. 3). Sustained SA activity might be an
454 important source of defense signaling in poplars, especially concerning the production of
455 phenylpropanoid-associated compounds, such as proanthocyanidins (condensed tannins) and
456 their monomeric catechin constituents (Ullah et al. 2019).

457 In closing this section, it is also worth noting the potential interactions between isoprene
458 and oxylipin synthesis and signaling in nature, especially involving signals that affect MEP
459 pathway activity and, ultimately, the production of isoprene itself. It has been reported that
460 transcription of 1-deoxy-D-xylulose 5-phosphate synthase 2 (*DXS*), of the MEP pathway,
461 increases in response to exogenous JA and exposure to mechanical stress or wounding (Tretner
462 et al. 2008). A second study showed an increase in *DXS* expression, but a decrease in *ISPS*
463 expression and isoprene emission, in *Ficus septica* treated with JA (Parveen et al. 2019b).
464 Collectively, we can assume that isoprene and JA levels are modulated in native emitters (or
465 when plants are engineered to emit isoprene) to optimize responses to herbivory and other
466 stresses. Enhancement of *DXS* expression and suppression of *ISPS* expression by JA likely
467 enables MEP pathway metabolites to be diverted from isoprene production and channeled into
468 the synthesis of other MEP pathway-derived secondary metabolites, many of which are involved
469 in defense roles (e.g., the higher-order terpenes). Parveen et al. (2019b) also found cis-elements
470 on the *Ficus septica* *ISPS* promotor that makes it responsive to regulation by MYC2, a TF
471 protein under the control of Me-JA synthesis. This establishes the condition for JA-associated
472 feedback to isoprene signaling, and represents an example of potential crosstalk between JA and
473 isoprene pathway signaling. A recent analysis of sequence motifs in the promoter of *ISPS* from
474 poplar revealed several elements that are likely responsive to multiple signaling pathways,

475 including those associated with Me-JA, gibberellins, auxins, SA and ABA (see Fig. S3 in
476 Miloradovic van Doorn et al. 2020).

477 **Isoprene as a mediator of the growth-defense tradeoff in the face of climate stress**

478 Given the several studies that show isoprene effects on gene expression and cellular
479 signaling, there is need for a theory that provides a broader adaptive scope for the trait. The
480 multiple interactions among signaling pathways are complex and variable across species, which
481 make it difficult to identify the primary costs and benefits that have shaped the trait. However,
482 there are some clear influences that lead us to a starting point. In every species and experimental
483 system examined to date, the presence of isoprene causes large, positive effects on gene
484 expression in the shikimate and phenylpropanoid pathways. Phenolic compounds and
485 phenylpropanoids serve multiple adaptive roles in plants, including those associated with growth
486 (e.g., auxins), defense (alkaloids, salicylic acid, phenolic glycosides, tannins, flavonols), and
487 abiotic stress tolerance (anthocyanins, flavonoids). The universal trend toward isoprene-induced
488 upregulation of shikimate and phenylpropanoid production, shows the central importance of
489 these pathways to integrated, multi-trait adaptive responses that drive the growth-defense
490 tradeoff within the constraint of abiotic stress tolerance.

491 While there are clear interspecific similarities in the way that isoprene influences
492 expression in the shikimate and phenylpropanoid pathways, there are differences in its role
493 within the terpenoid and JA pathways. In a comparative analysis of isoprene-associated gene
494 expression in *Arabidopsis* and tobacco, Zuo et al. (2019) concluded that isoprene shifts
495 expression in favor of defense-associated metabolites in tobacco, but growth-associated
496 metabolites in *Arabidopsis*. In this study, we noted that poplar was observed to downregulate the
497 potential for carotenoid and α -tocopherol synthesis, as well as the prenyl-transferase activity

498 associated with REF proteins, while in *Arabidopsis* expression for all three were increased.
499 These differences are likely to reflect the different selection regimes experienced by the species.
500 Interestingly, *Arabidopsis* and tobacco were capable of responding to the introduction of
501 isoprene, despite no past history of selection for phenotypes expressing the trait. This indicates
502 that isoprene is a trait that, once evolved, can rapidly become incorporated into naïve
503 phenotypes. Many of the regulatory sequence motifs in the *ISPS* promoter of IE poplar are
504 shared with the promoter of *1,8-cineole synthase (AtTPS-Cin)* from NE *Arabidopsis*, the gene for
505 a terpene synthase that is closely-related to *ISPS* (see Fig. S3 in Miloradovic van Doorn et al.
506 2020). These include regulatory elements that respond to ABA, MeJA, SA and binding with the
507 MYB transcription factor family. Thus, at least one terpene synthase, unrelated to isoprene
508 biosynthesis, exists in *Arabidopsis* with pre-existing metabolic connections to several cellular
509 signaling pathways. These factors might facilitate rapid (generationally-speaking) directional
510 selection during the evolution of isoprene signaling within naïve lineages, and provide high
511 selective value to the trait early during its appearance.

512 The adaptive value of isoprene is likely also linked to its association with photosynthesis.
513 From the earliest days of isoprene research, it has been recognized that variances in emission rate
514 and photosynthesis are correlated across many determining variables (Sanadze et al. 1972;
515 Rasmussen and Jones 1973; Tingey et al. 1979). Subsequent research showed numerous
516 biochemical dependencies of isoprene synthesis on photosynthate and photoreductant (Sharkey
517 and Yeh 2001; Sharkey and Monson 2017; Sharkey et al. 2020). These dependencies provided
518 explanations for: (1) the positive correlations of isoprene emission and photosynthetic capacity
519 (Monson and Fall 1989; Loreto and Sharkey 1990), (2) the role of increased resources, such as
520 nitrogen, in supporting higher isoprene emission rates (Harley et al. 1994; Litvak et al. 1996;

521 Fernandez-Martinez et al. 2018), and (3) the observations that within a phylogenetic clade,
522 isoprene emissions are found in species with higher photosynthetic capacities, higher growth
523 rates, and niche affinities that favor sunny habitats (Harley et al. 1999; Dani et al. 2014; Loreto
524 and Fineschi 2015). The many metabolic and ecological associations between isoprene and
525 photosynthesis likely elevate the selection differential for isoprene. (Selection differential is the
526 difference between the average value to fitness of a quantitative trait in an entire population and
527 the average value in those individuals selected to reproduce and form the next generation). With
528 the new insight into isoprene's role in regulating gene expression, we can expand the scope of
529 isoprene's trait value to include plant defense. Metabolically, isoprene sits in a position wedged
530 between the processes that determine the supply of photosynthetic substrates and the demand of
531 secondary metabolite products – the two economic determinants underlying most contemporary
532 growth-defense theories.

533 Relying on these concepts, we offer a new formal theory for the adaptive role of isoprene
534 emission (Fig. 4). Within the context of traditional theories (e.g., the GDBH), the growth-defense
535 tradeoff can be represented as a negative correlation constrained at an upper limit by
536 photosynthetic capacity (e.g., Züst and Argawal 2017). The upper limit is reduced in the face of
537 stress, providing the realized limit for operation of the tradeoff (shown as the dashed line in Fig.
538 4). The observed tradeoff for non-isoprene emitting plants (NE) must occur within the space
539 below the realized limit. We propose that the evolution of isoprene emission (IE) will provide the
540 advantage of better tolerating the stresses that determine the realized limit, allowing plants to
541 accommodate allocations to defense with less cost to growth.

542 This view of isoprene emissions is compatible with previous observations and theories
543 proposing a positive effect on abiotic stress tolerance (e.g., Loreto and Schnitzler 2010). It is also

544 compatible with past reports that isoprene protects photosynthesis against the impact of oxidative
545 stress (e.g., Agati et al. 2013). The new theory that we offer only diverges from past perspectives
546 in positioning the adaptive role of isoprene as a direct contributor to multiple signaling networks,
547 affecting broad patterns of gene expression, rather than the role of isoprene as a single and
548 isolated actor. Further, our theory expands the role of isoprene from one of stress tolerance,
549 alone, to one of stress tolerance within the context of the growth-defense tradeoff.

550 The theory that we propose is also consistent with the perspective that the G-D tradeoff is
551 dependent on the overall amount of resource available for G-D support (van Noordwijk and de
552 Jong 1986; Agrawal 2020). The dependency of the G-D tradeoff, along with life-history
553 tradeoffs, on the total amount of available resource has been explained by the simple fact that as
554 organisms obtain more resources, either through changes in environmental conditions or the
555 evolution of novel traits, they often allocate more resource to both G and D (van Noordwijk and
556 de Jong 1986; Houle et al. 1991; Agrawal et al. 2010). From our theory, the evolution of
557 isoprene emission, and its positive influence on the expression of selected defense compounds
558 that also carry advantages in abiotic stress tolerance, will facilitate greater amounts of
559 photosynthate that will be available for allocation to both G and D, compared to non-emitting
560 phenotypes. This would drive positive selection for the trait in certain environments.

561 **The mechanism(s) of isoprene in cellular signaling networks**

562 Isoprene is relatively insoluble in water, and its solubility in lipids, while higher than that in
563 water, is less than often assumed (Appendix 2 in Harvey et al. 2015). As a result, isoprene exists
564 predominantly in the gas phase of the leaf and is quickly lost through diffusion. Isoprene's high
565 volatility is difficult to reconcile with conventional cellular signaling mechanisms. Other
566 signaling volatiles, such as Me-JA are converted to an active and relatively soluble form by

567 conjugation; for example, with isoleucine. To date, no conjugates involving isoprene have been
568 identified. The hydrophobic nature of isoprene, however, may confer signaling advantages,
569 including its ability to cross membranes and influence pathways that are distributed across
570 multiple organelles, and to access signaling components in the hydrophobic domain of
571 membrane lipid bilayers.

572 Conventional signal-receptor interactions are based on macromolecular shape
573 complementarity (Covarrubias et al. 2020). In the case of the plant immune signal, salicylic acid
574 (SA), SA-protein binding interactions, involving a set of cooperating transcriptional activators
575 (NPR1) and repressors (NPR3 and NPR4), occurs in a conventional manner and controls SA
576 signal reception (Wu et al. 2012; Ding et al. 2018). However, redox-driven interactions are also
577 common, especially in signaling networks based on reactive oxygen (ROS) and nitrogen species
578 (RNS) (Nathan 2003; Shetty et al. 2008; Vickers et al. 2009a). For ROS, signaling occurs
579 through the oxidation of key amino acids, such as cysteine, on the surface of transcription-
580 modifier proteins, leading to direct control over gene expression (Neill et al. 2002). In the case of
581 RNS, protein activities are modified by post-translational modification (PTM) involving nitric
582 oxide (NO), a process known as S-nitrosylation (Spadaro et al. 2010). S-nitrosylation most
583 commonly reduces enzyme activities in plants, leading to reduced rates of metabolite production
584 (Lindermayr et al. 2005). S-nitrosylation is known to regulate inducible responses to abiotic
585 stress (Corpas et al. 2011; Vanzo et al. 2014; Begara-Morales et al. 2018), and past studies in
586 poplar have shown that isoprene (through an unknown mechanism) modulates the H₂O₂-
587 dependent translational cascade and NO-dependent post-translational cascade (Vanzo et al
588 2016). Several past theories concerning the adaptive functions of isoprene have posited that
589 isoprene's potential role as an antioxidant molecule provides a mechanism for direct chemical

590 reactions that reduce ROS or RNS to a level that causes pathway modulation (Velikova et al.
591 2005; 2008; Vickers et al. 2009a; Behnke et al. 2010a). Such past theories of isoprene directly
592 affecting cellular redox signaling through its role as an antioxidant are too narrow to account for
593 the broad set of influences and diverse set of pathways indicated in the recent multi-omic
594 analyses.

595 Furthermore, there are reasons to question the feasibility of isoprene acting as an effective
596 antioxidant metabolite, even separate from its potential to directly trigger ROS and RNS
597 networks. Given that the concentration of isoprene is about 60 molecules per million (lipid
598 molecules), which is much lower than the cellular concentration of known antioxidant molecules
599 such as carotenoids (Harvey et al. 2015), it does not follow logically that isoprene can increase
600 the antioxidant margin of cellular protection to a significant advantage. Isoprene only contributes
601 ~0.1% the double bonds available for nucleophilic reactions, compared to carotenoids (Harvey et
602 al. 2015); and, while some past reports have included observations of isoprene's oxidation
603 products, such as methyl vinyl ketone and methacrolein, as leaf volatiles (Jardine et al. 2012),
604 these observations have not been widely confirmed. From these perspectives, isoprene does not
605 appear to be a plausible candidate to provide effective antioxidant protection, or to act as an
606 efficient redox signal compound, at least not with respect to direct alterations of ROS and RNS
607 concentrations. Harvey and Sharkey (2016) hypothesized that in *Arabidopsis* the presence of
608 isoprene causes an upregulation of key TFs involved in phenylpropanoid and carotenoid pathway
609 expression which, in turn, alter cellular ROS and RNS concentrations, as well as activating other
610 signal cascades. This perspective leads to the hypothesis of an indirect role between isoprene and
611 cellular oxidants (Fig. 5). The recent discovery of isoprene effects in roots, especially in the
612 vascular tissues, where ISPS is present but expressed at very low levels, supports a role for

613 indirect (through signaling) involvement of isoprene in ROS signaling (Miloradovic van Doorn
614 et al. 2020). The transgenic (RNAi) suppression of *ISPS* expression, even at the very low levels
615 present in roots, affected lateral root development in a manner consistent with ROS
616 accumulation. Furthermore, expression of *ISPS* in roots was increased in the presence of auxin,
617 suggesting a complex interaction with developmental processes; well beyond the more
618 commonly cited context of antioxidant activity. These observations are most compatible with the
619 model of isoprene acting as a signal molecule, not an antioxidant. The studies by Harvey and
620 Sharkey (2016) and Miloradovic van Doorn et al. (2020) provide good reason to recast the
621 discussion of isoprene effects, away from single molecule redox reactions, and toward one of
622 broad interactions involving numerous signaling pathways.

623 At this time, it is not clear as to how the dependency of isoprene emissions on diurnal and
624 seasonal environmental change fits into its role as a signaling metabolite. In some ways, isoprene
625 is similar to an inducible defense trait – its expression is promoted by high temperature, high
626 light, mechanical wounding and high nitrogen availability (Harley et al. 1999; Sharkey et al.
627 2008). Observations have also revealed higher emission rates during drought recovery periods
628 (Fortunati et al. 2008; Tattini et al. 2015; Velikova et al. 2016; 2018), and in the isoprene-
629 emitting species *Arundo donax* the increased emission rates were accompanied by increased
630 production of selected phenylpropanoid compounds (Velikova et al. 2016; Ahrar et al. 2017).
631 Generally, the seasonal environments that promote isoprene emission also provide reliable cues
632 of herbivory risk – warm, high light, mid-summer weather with high soil fertility, and/or
633 following the stressful conditions of a drought. The work by Tattini et al. (2015) showed that
634 metabolic adjustments that promote tolerance of abiotic stress extremes can also occur on the
635 scale of hours, and are timed for midday stress extremes; including isoprene emission. Thus, like

636 the cues for inducible defenses, high isoprene emission rates could condition plants to anticipate
637 future episodes of combined abiotic and biotic stress on quite short time scales, and thus provide
638 an effective form of phenotypic plasticity.

639 **Conclusions**

640 The explanation of the effects of isoprene on plants proposed here represents a major shift
641 in our thinking away from direct effects of isoprene, for example, by changing membrane
642 properties or quenching ROS, to indirect effects, through changes in gene expression and protein
643 abundances. The presence of isoprene affects a number of transcription factors important in
644 signaling processes involved in shikimate, phenylpropanoid, terpenoid and oxylipin synthesis,
645 and in the production of numerous compounds involved in plant growth, defense, and abiotic
646 stress tolerance. This suggests that isoprene can affect the outcome of several regulatory
647 cascades. We presume that in some conditions the altered regulatory landscape must be
648 deleterious, in order to account for the frequent losses of the isoprene emission trait. The
649 research challenges that lie ahead include developing an improved context for how the multiple
650 effects of isoprene are adjusted in the face of interactive influences involving growth, herbivory
651 and threats from extreme climate stress, to optimize fitness. Improved knowledge of how
652 environmental variation influences the processes that control trait tradeoffs, including those
653 influenced by isoprene, will not only lead to more accurate ecological theories concerning plant
654 adaptation, but will also facilitate better strategies for the development of sustainable agriculture.
655 With the rapidly expanding opportunities for genetic modification using CRISPR-Cas9
656 technology, the possibility exists to develop finely-targeted strategies to mediate the growth-
657 defense tradeoff to simultaneously improve yield through the combined effects of higher rates of

658 biomass increase and lower rates of biomass loss. Isoprene might have an important role in the
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660

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667

668 **Data accessibility** This is a synthesis study such that all data was drawn from the existing
669 publications cited in the text; including electronic supplementary materials.

670

671

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1032 **Figure Legends**

1033 **Figure 1.** Protein or transcriptome expression modulation in the shikimate (A) and
1034 phenylpropanoid (B) pathways in IE *Populus x canescens*, compared to NE transgenic lines.
1035 Bold blue or orange arrows indicate increased or decreased expression in the presence of
1036 isoprene, respectively. Protein labels in italics with asterisks indicate transcriptome data from
1037 Behnke et al. (2010a); otherwise data are from the proteome study of Monson et al. (2020). Fold-
1038 change (FC) multipliers are shown in parentheses of some steps and refer to isoprene-
1039 emitting/non-emitting (IE/NE) plants. Not all pathway steps are shown; only those with
1040 significant ($P < 0.05$; $n = 4$ replicate trees) changes in transcription or protein amount (bold
1041 arrows; $0.5 \leq FC \leq 1.5$) or steps that produce metabolites that are not directly regulated, but are
1042 of importance to defense, growth or stress tolerance as discussed in the main text (broken
1043 arrows). Metabolite boxes highlighted in green, red or yellow refer to those most influencing
1044 growth, defense or stress tolerance, respectively.

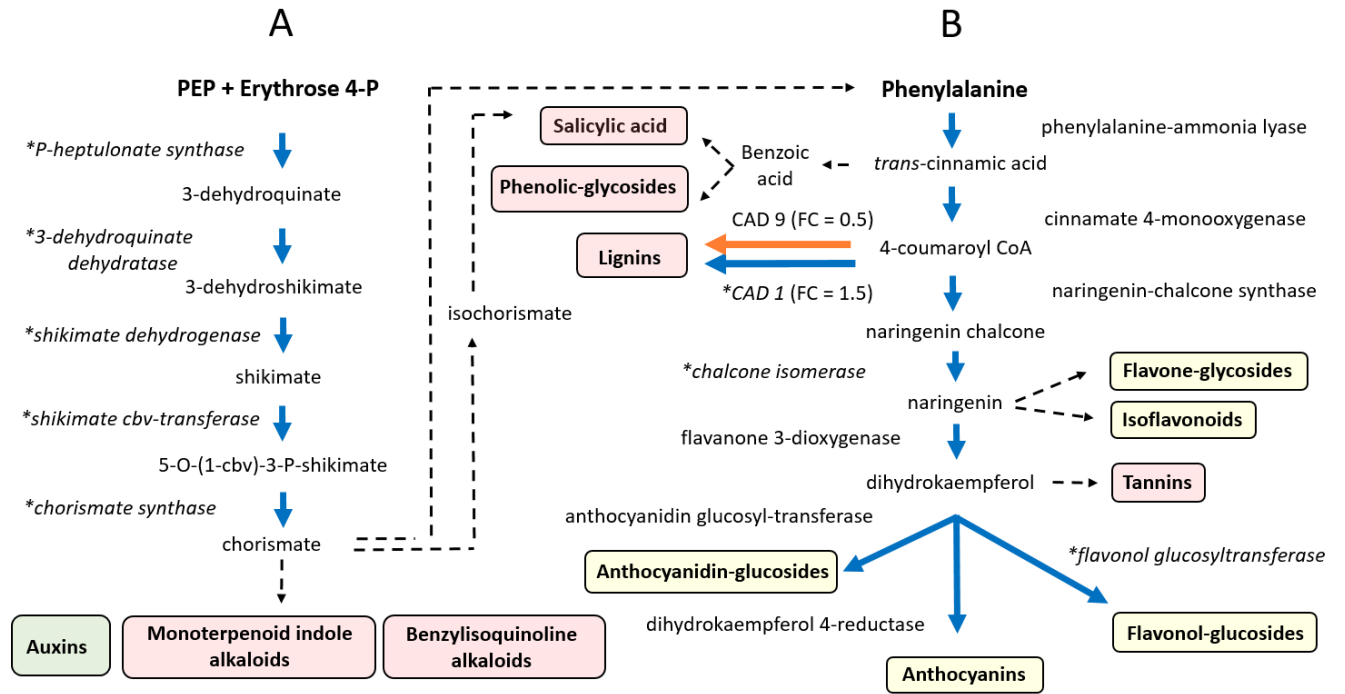
1045 **Figure 2.** Proteomic and transcriptomic adjustments in the pathways of terpenoid synthesis in
1046 response to presence or absence of isoprene in *Populus x canescens* and *Arabidopsis*. Orange and
1047 blue arrows/labels indicate decreased or increased protein contents or transcripts, respectively, in
1048 the presence of isoprene. Fold-change (FC) multipliers are shown in parentheses for some steps
1049 and refer to isoprene-emitting/non-emitting (IE/NE) plants. Not all pathway steps are shown;
1050 only those with significant ($P < 0.05$) changes in protein amount (bold arrows) or steps that
1051 produce metabolites of importance to defense, growth or stress tolerance (dashed arrows) as
1052 discussed in the main text. Metabolite boxes highlighted in green, or yellow refer to those most
1053 influencing growth or stress tolerance, respectively. **Inset:** Bold arrows represent changes in
1054 *Arabidopsis* determined as observations that showed a consistent directional change in isoprene-

1055 fumigated wild-type *Arabidopsis*). All values are derived from Monson et al. (2020) and Harvey
1056 and Sharkey (2016).

1057 **Figure 3.** Proteomic or transcriptomic adjustments in the oxylipin pathways in response to
1058 presence or absence of isoprene in *Populus x canescens*, *Nicotiana tabacum* (tobacco) and
1059 *Arabidopsis*. Blue and orange bold arrows indicate increased or decreased expression,
1060 respectively, in the presence of isoprene. Arrows with one or two asterisks represent changes in
1061 tobacco and/or *Arabidopsis* determined as observations that showed a consistent directional
1062 change in one or two, respectively, of the three experimental systems that were examined (wild-
1063 type, isoprene-fumigated *Arabidopsis* plants and transgenic IE *Arabidopsis* and tobacco). Not all
1064 pathway steps are shown; only those with significant ($P < 0.05$) changes in transcription or protein
1065 amount (bold arrows) or steps that produce metabolites of importance to defense, growth or
1066 stress tolerance as discussed in the main text. Data from Monson et al. (2020), Harvey and
1067 Sharkey (2016) and Zuo et al. (2019).

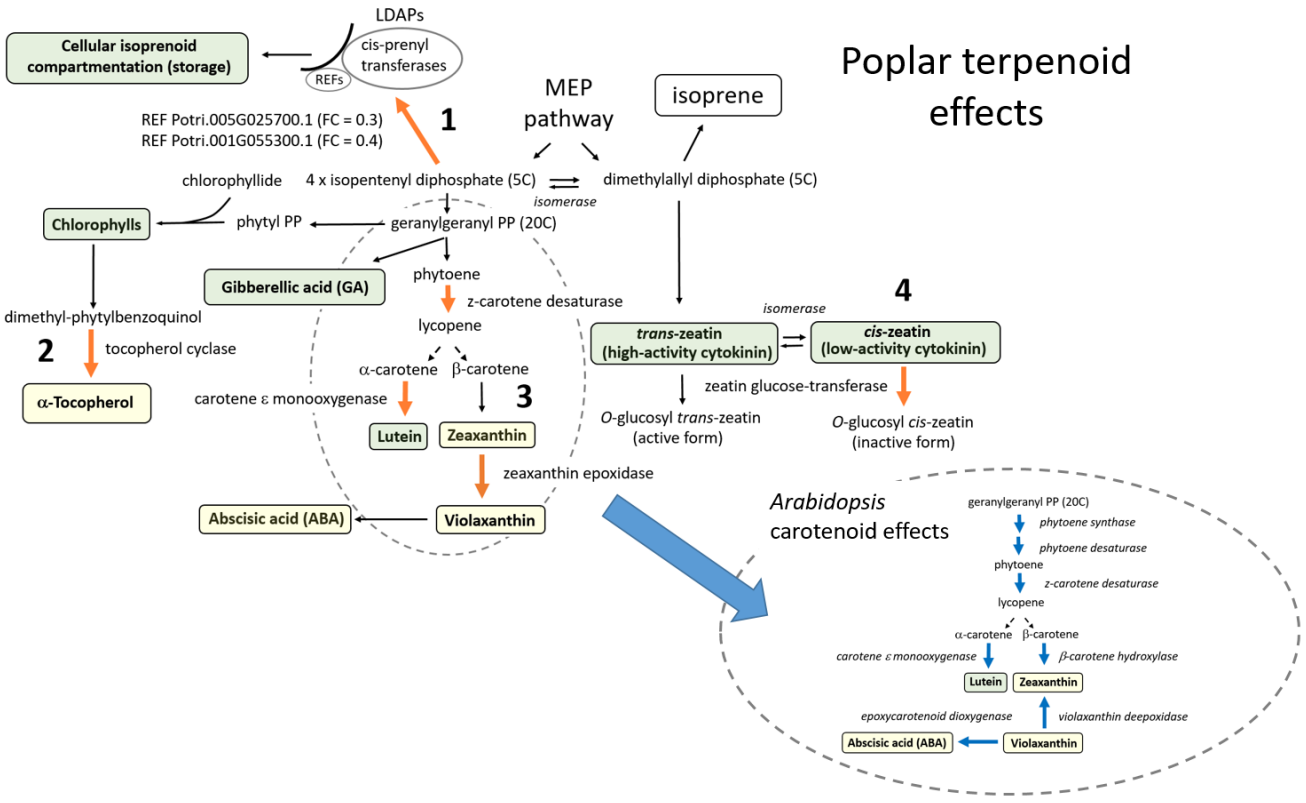
1068 **Figure 4.** Graph showing relationships among growth, defense and isoprene emissions in the
1069 presence of abiotic stress. From the perspective of plant carbon budgets, the growth-defense
1070 tradeoff can be described in relation to a limit set by the plant's maximum photosynthetic
1071 capacity. In the absence of isoprene emissions, environmental stresses will reduce the
1072 photosynthetic capacity to a hypothetical limit defined by the dashed line. The evolution of
1073 isoprene emissions in taxonomic lineages will improve the potential for allocation to defense
1074 with reduced cost to growth, compared to the case for non-emitters. Based on concepts originally
1075 presented in Stamp (2003) and Züst and Argawal (2017).

1076 **Figure 5.** Responses to stresses or isoprene involve perception (yellow chevrons) or signal
1077 transduction pathways (yellow arrows). Pathways stimulated (blue) or inhibited (orange) by
1078 stress are also shown, as is isoprene modulation of these pathways. Direct effects are shown as
1079 dark grey arrows. Four possible roles for isoprene in stress responses are depicted. Pathway 1
1080 depicts effects of isoprene on an isoprene-specific signal cascade that can affect growth and / or
1081 defense. Pathway 2 suggests isoprene can modify signal cascades, for example JA or GA
1082 signaling pathways. Pathways may be upregulated (blue) or downregulated (orange) by stress
1083 and isoprene could modulate the changes in pathway regulation caused by stress. By modulating
1084 signal transduction pathways, isoprene could prevent the buildup of ROS and NO rather than
1085 quenching as a mechanism for keeping their concentrations low. Pathway 3 (right side) is direct
1086 quenching of H₂O₂ and other ROS. This pathway was highly favored in earlier studies, but is
1087 now considered unlikely to account for the effects of isoprene on plants. Pathway 4 shows
1088 isoprene signaling causing changes in metabolites that can quench ROS and or NO. Disruption
1089 of transduction pathways initiated by H₂O₂ or NO (e.g., the hypersensitive response) are
1090 suppressed by isoprene effects and could indirectly affect programmed cell death (PCD; see
1091 Vickers et al. 2009a; Vanzo et al. 2016).
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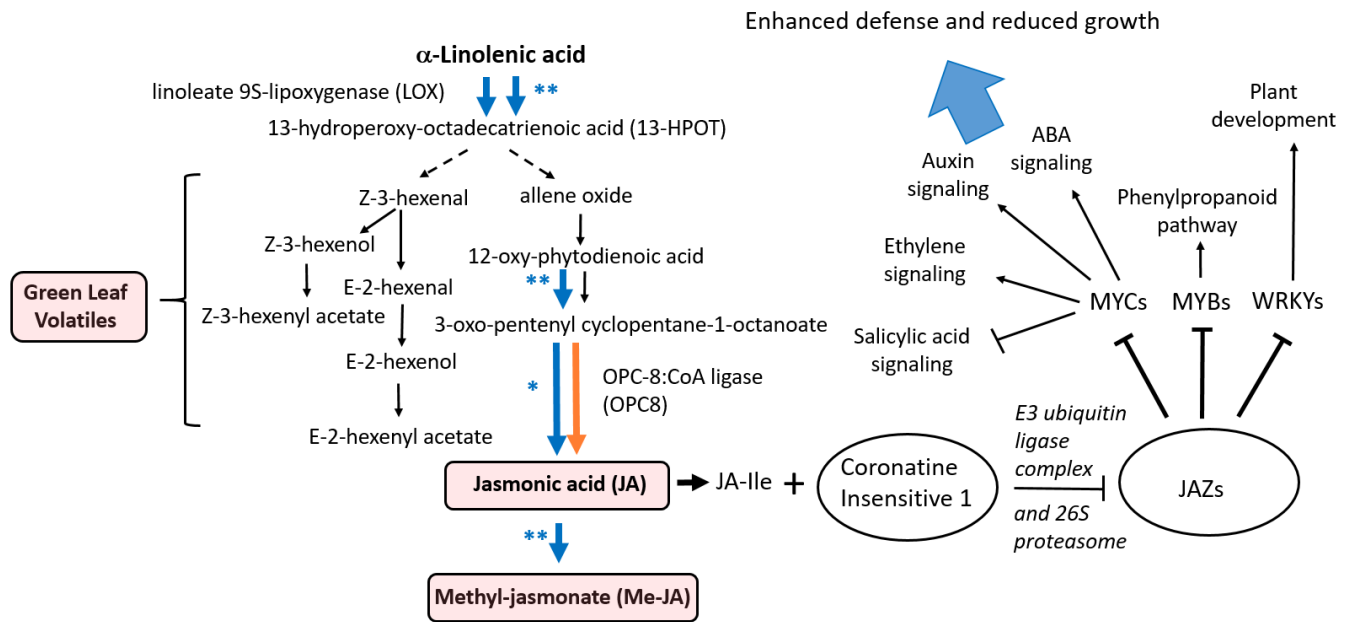
Figure 1



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Figure 2

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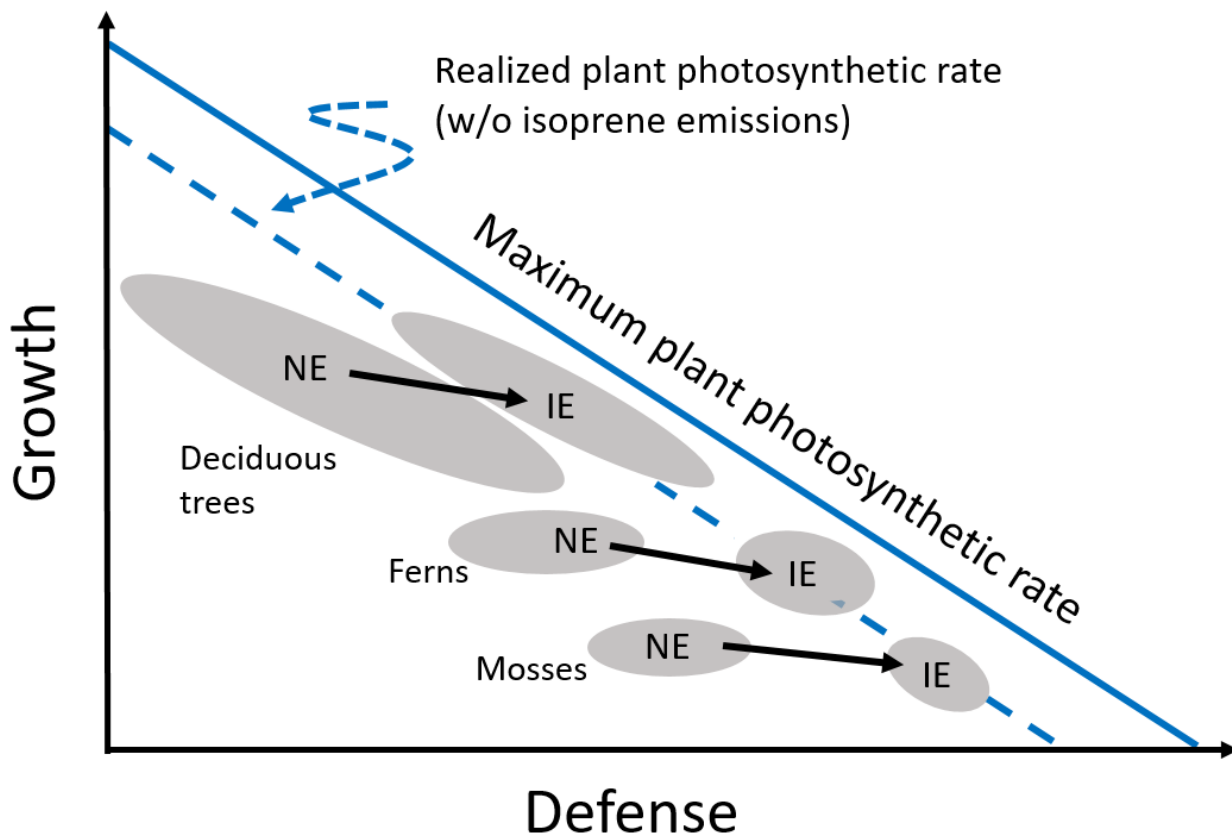
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1111 **Figure 3**

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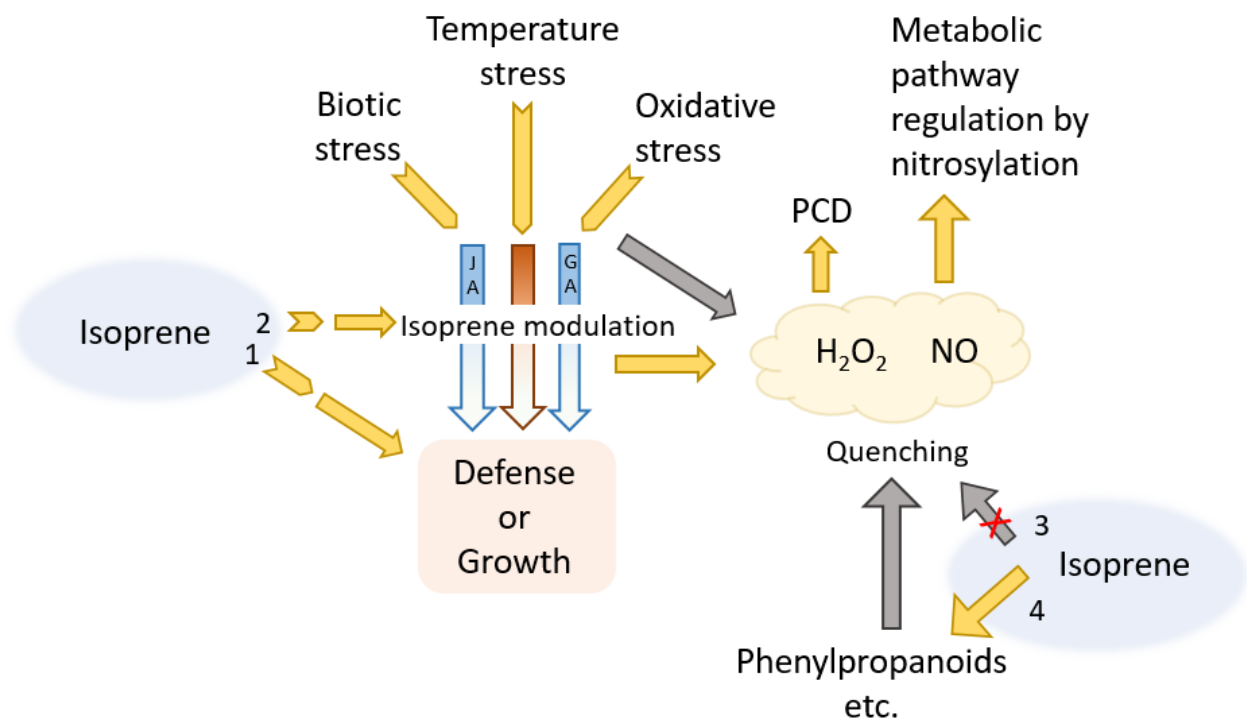
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1118 **Figure 4**

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Figure 5