

REVIEW

View Article Online
View Journal | View Issue



Cite this: *Nat. Prod. Rep.*, 2023, 40, 1901

Chemistry, biosynthesis and biology of floral volatiles: roles in pollination and other functions

Stefan Dötterl *^a and Jonathan Gershenzon *^b

Covering: 2010 to 2023

Floral volatiles are a chemically diverse group of plant metabolites that serve multiple functions. Their composition is shaped by environmental, ecological and evolutionary factors. This review will summarize recent advances in floral scent research from chemical, molecular and ecological perspectives. It will focus on the major chemical classes of floral volatiles, on notable new structures, and on recent discoveries regarding the biosynthesis and the regulation of volatile emission. Special attention will be devoted to the various functions of floral volatiles, not only as attractants for different types of pollinators, but also as defenses of flowers against enemies. We will also summarize recent findings on how floral volatiles are affected by abiotic stressors, such as increased temperatures and drought, and by other organisms, such as herbivores and flower-dwelling microbes. Finally, this review will indicate current research gaps, such as the very limited knowledge of the isomeric pattern of chiral compounds and its importance in interspecific interactions.

Received 8th May 2023

DOI: 10.1039/d3np00024a

rsc.li/npr

1	Introduction	6.3	Flies
2	Reports of notable new floral volatiles	6.3.1	Mutualistic pollination systems
3	Biosynthesis of floral volatiles	6.3.2	Deceptive pollination systems
3.1	Fatty acid derivatives	6.4	Bees
3.2	Terpenoids	6.4.1	Generalist bees
3.3	Phenylpropanoids and benzenoids	6.4.2	Nocturnal bees
3.4	Other amino acid derivatives	6.4.3	Pollen-specialist (oligolectic) bees
4	Regulation of floral volatile emission	6.4.4	Male euglossine bees
4.1	Transcription factors	6.5	Wasps
4.2	Other factors regulating biosynthesis	6.6	Other pollinators
4.3	Factors regulating storage, transport and emission	7	Functions of floral volatiles other than pollinator attraction
5	Effects of abiotic and biotic factors on floral scent emissions	7.1	Repelling florivores
5.1	Air pollution	7.1.1	Defense against pathogenic microorganisms
5.2	Temperature and drought	7.1.2	Attraction of enemies of plant pests
5.3	Nutrient availability and soil type	8	Chirality in flower scents and its importance for inter-specific interactions
5.4	Bacteria and fungi	9	Conclusions
5.5	Herbivores	10	Author contributions
6	Floral volatiles as attractants for flower visitors	11	Conflicts of interest
6.1	Beetles	12	Acknowledgements
6.2	Moths, butterflies	13	References
6.2.1	Hawkmoths		
6.2.2	Settling moths		
6.2.3	Butterflies		

^aDepartment of Environment & Biodiversity, Paris Lodron University Salzburg, Hellbrunnerstr 34, 5020 Salzburg, Austria. E-mail: stefan.doetterl@plus.ac.at

^bDepartment of Biochemistry, Max Planck Institute for Chemical Ecology, Hans-Knöll-Strasse 8, 07745 Jena, Germany. E-mail: gershenzon@ice.mpg.de

1 Introduction

Floral volatiles are key mediators of plant–pollinator interactions.^{1–6} Considering that flowering plants (angiosperms) are by far the most diverse group of terrestrial plants and that 85% of these plants are pollinated by animals,⁷



among them various crops,⁸ pollinator attraction by olfactory means is a highly important process in terrestrial ecosystems and for crop production. About 1700 floral volatiles had already been described more than a decade ago,⁴ but many new compounds have been discovered since then,^{e.g.,9–12} making floral scent a high dimensional trait with many possible ways to vary among species. Indeed, floral scents are highly variable among species, either due to differences in the compounds emitted or due to differences in the absolute or relative amounts of single scent compounds. Thus, it is not surprising that floral scents are involved in shaping plant–pollinator networks.^{13–15} Quite recently, it has been demonstrated that floral volatiles serve additional functions, such as defending flowers against enemies (e.g., pathogens)¹⁶ and attracting predators of plant pests.¹⁷ It was also shown that they are attractive to florivores and their occurrence is influenced by biotic and abiotic stressors at the organ, plant and population levels. As our knowledge of floral volatile biosynthesis has increased, especially on the regulation of biosynthesis, we have learned more about the mechanisms underlying these patterns. This review summarizes advances in floral scent research since 2010 from chemical, molecular and ecological perspectives, excluding recent findings on the olfactory communication between orchids and their pollinators, given that this topic is considered by a recent review.¹⁸ It will also indicate current research gaps, such as the very limited knowledge of the occurrence and ratio of stereoisomers of chiral compounds and its importance in interspecific interactions.

2 Reports of notable new floral volatiles

Floral volatiles have long been classified into a few basic groups according to their chemistry and probable

biosynthetic origin: fatty acid derivatives, terpenoids, phenylpropanoids/benzenoids and other amino acid derivatives. During the period of this review, specific compounds from each of these groups were identified as floral volatiles for the first time. Representatives of these compounds are summarized in Table 1 and Fig. 1–5, and the biological roles in pollination of several of these compounds are discussed later in the chapter. Compounds isolated from orchids are also covered by Perkins *et al.*,¹⁸ but are included here for completeness.

The length of this list of new floral volatiles suggests that many more substances still remain to be discovered in the blends emitted from flowers. This is not simply because of advances in analytical instrumentation that provide ever increasing sensitivity. Some of the new volatiles are compounds that had been missed in earlier surveys that relied only on headspace trapping methods, rather than also carrying out solvent extraction from floral tissues.¹⁹ Floral volatiles of low volatility, such as the C₂₀ diterpene alcohol, copalol,²⁰ are especially difficult to detect by headspace analysis. However, most of the new volatiles reported are constituents of floral blends of species that had not been previously examined.

The discovery of new floral volatiles in new species is consistent with the concept that plant scents have been generally selected to be unique to attract specific pollinators and ensure a high degree of intraspecific gene flow during pollen transfer. The release of unique compounds has been described as creating a private channel between the plant and pollinator species,^{5,19} with volatiles detected by the intended receiver, but not by other potential flower visitors. On the other hand, specificity can also be achieved by the emission of blends of widespread compounds in unique proportions, and new floral blends dominated by previously characterized widespread compounds are continually reported.



Stefan Dötterl studied biology at the University of Bayreuth in Germany, where he also obtained his PhD degree (2004) and worked on his habilitation (submitted 2010). In 2012, he became Full Professor at the Paris Lodron University in Salzburg, Austria. Since more than 20 years, he is working on the ecology of evolution of floral scents and of plant–pollinator interactions. Together with his

team and collaboration partners, he identified floral scents, among them several novel natural products, from plants of various lineages. Also, he identified pollinator attractants for various pollinator groups, among them bees, beetles, flies, lepidopterans and mammals.



Jonathan Gershenzon has been fascinated with the volatile natural products of plants for much of his career. He completed his undergraduate studies at the University of California at Santa Cruz with Jean Langenheim, obtained a PhD at the University of Texas at Austin with Tom J Mabry, carried out post-doctoral research at Washington State University with Rodney Croteau, and then joined the Max Planck

Institute for Chemical Ecology in Jena, Germany. With numerous coworkers and collaborators he has been investigating the biochemistry and ecological function of floral and vegetative plant volatile for a number of years, specializing in the terpene and nitrogen-containing volatiles of mints, Arabidopsis, maize, poplars and conifers.



Table 1 New and notable floral volatiles 2010–2022 (see also Fig. 1–5)

Compound number and name	Plant species	Family	Reference
Fatty acid derivatives			
(1) (6Z,9Z)-1,6,9-tricosatriene	<i>Pterostylis orbiculata</i>	Orchidaceae	21
(2) (<i>E</i>)-2-octen-1-yl acetate	<i>Ceropegia sandersonii</i>	Apocynaceae	22
(3) isojasmol	<i>Thaumatococcus mello-baretoanum</i> , <i>Xanthosoma hylaeae</i> , <i>Ludovia lancifolia</i>	Araceae, Cyclanthaceae	11, 12 and 23
(4) isojasmyl acetate	<i>Xanthosoma hylaeae</i>	Araceae	11
(5) dehydrojasmon	<i>Thaumatococcus mello-baretoanum</i>	Araceae	11
(6) <i>trans</i> -3-methyldodecano-4-lactone	<i>Passiflora chocoensis</i> , <i>Kefersteinia aurorae</i> , <i>Scuticaria salesiana</i> , <i>Lycaste brevispatha</i>	Passifloraceae, Orchidaceae	24
(7) <i>trans</i> -(<i>Z</i>)-3-methyldodec-7-eno-4-lactone	<i>Kefersteinia</i> , <i>Scuticaria salesiana</i> , <i>Lycaste brevispatha</i> , <i>Chaubardiella dalessandroi</i> , <i>Mormodes romanii</i>	Orchidaceae	24
(8) <i>trans</i> -(<i>Z</i>)-3-methyldodec-6-eno-4-lactone	<i>Kefersteinia pellita</i> , <i>Chaubardiella dalessandroi</i> , <i>Mormodes romanii</i>	Orchidaceae	24
(9) <i>cis</i> -(<i>Z</i>)-3-methyldodec-6-eno-4-lactone	<i>Kefersteinia pellita</i> , <i>Chaubardiella dalessandroi</i> , <i>Mormodes romanii</i>	Orchidaceae	24
(10) (<i>Z</i>)-deca-7,9-dieno-5-lactone	<i>Gardenia brighamii</i>	Rubiaceae	24
(11) (16 <i>S</i> ,9 <i>Z</i>)-16-ethylhexadec-9-enolide	<i>Disa forficaria</i>	Orchidaceae	25
(12) (<i>Z</i>)-3-isopropylpent-3-en-1-ol	<i>Syngonium hastiferum</i>	Araceae	26
(13) (<i>S</i>)-2-(tetrahydrofuran-2-yl)acetic acid	<i>Cryptostylis ovata</i>	Orchidaceae	27
(14) methyl-(<i>S</i>)-2-(tetrahydrofuran-2-yl)acetate	<i>Cryptostylis ovata</i>	Orchidaceae	27
(15) ethyl-(<i>S</i>)-2-(tetrahydrofuran-2-yl)acetate	<i>Cryptostylis ovata</i>	Orchidaceae	27
(16) 2-ethyl-5-propylcyclohexan-1,3-dione (chiloglottone 1)	<i>Chiloglottis</i>	Orchidaceae	28
(17) 2-ethyl-5-pentylcyclohexan-1,3-dione (chiloglottone 2)	<i>Chiloglottis</i>	Orchidaceae	28
(18) 2-butyl-5-methylcyclohexan-1,3-dione (chiloglottone 3)	<i>Chiloglottis</i>	Orchidaceae	28
(19) 5-allyl-2-ethylcyclohexan-1,3-dione (chiloglottone 4)	<i>Chiloglottis</i>	Orchidaceae	28
(20) 2-butyl-5-propylcyclohexan-1,3-dione (chiloglottone 5)	<i>Chiloglottis</i>	Orchidaceae	28
(21) 2-hexyl-5-methylcyclohexan-1,3-dione (chiloglottone 6)	<i>Chiloglottis</i>	Orchidaceae	28
Terpenoids			
(22) (<i>Z</i>)-2-allylidene-6-methylhept-5-en-1-ol: (<i>Z</i>)-filamentol	<i>Yucca</i>	Asparagaceae	29
(23) (<i>Z</i>)-2-allylidene-6-methylhept-5-en-1-ol: (<i>Z</i>)-filamentol	<i>Yucca</i>	Asparagaceae	29
(24) (<i>Z,R</i>)-3-allylidene-5-(2-methylprop-1-enyl) dihydro(3 <i>H</i>)furan-2-one: (<i>Z</i>)-filamentolide	<i>Yucca</i>	Asparagaceae	29
(25) (<i>E</i>)-2-allylidene-6-methylhept-5-en-1-ol: (<i>E</i>)-filamentol	<i>Yucca</i>	Asparagaceae	29
(26) (<i>E</i>)-2-allylidene-6-methylhept-5-en-1-ol: (<i>E</i>)-filamentol	<i>Yucca</i>	Asparagaceae	29
(27) (<i>E,R</i>)-3-allylidene-5-(2-methylprop-1-enyl) dihydro(3 <i>H</i>)furan-2-one: (<i>E</i>)-filamentolide	<i>Yucca</i>	Asparagaceae	29
(28) 2-(4-methylpent-3-en-1-yl)cyclopent-3-enone: filamentone	<i>Yucca</i>	Asparagaceae	29
(29) (<i>E</i>)-4,8-dimethylnona-1,3,7-trien-5-yl acetate	<i>Philodendron squamiferum</i>	Araceae	11
(30) 4-hydroxy-3-methyl-6 <i>S</i> -(pentan-2 <i>S</i> -yl)-5,6-dihydro-2 <i>H</i> -pyran-2-one (drakolide)	<i>Drakaea micrantha</i>	Orchidaceae	9
(31) (+)-copalol	<i>Cryptanthus burle-marxii</i>	Bromeliaceae	20
(32) (β)-safranin	<i>Anthurium salvadorens</i>	Araceae	24
(33) safranin acetate	<i>Anthurium salvadorens</i>	Araceae	24
Phenylpropanoids/benzenoids			
(34) 2-methoxy-4-vinyl phenol	<i>Dichaea pendula</i>	Orchidaceae	30
(35) 2-methoxy-6-methylacetophenone	<i>Colocasia gigantea</i> , <i>Cyphomandra divaricata</i> , <i>Pterygodium</i>	Araceae, Solanaceae, Orchidaceae	24
(36) (<i>Z</i>)-hept-4-en-2-yl salicylate	<i>Saraca asoca</i>	Fabaceae	24
(37) 3-acetyloxy-4-phenylbutan-2-one	<i>Ceropegia stenantha</i>	Apocynaceae	31
(38) 3-acetyloxy-1-phenylbutan-2-one	<i>Ceropegia stenantha</i>	Apocynaceae	31



Table 1 (Contd.)

Compound number and name	Plant species	Family	Reference
Other amino acid derivatives			
(39) 2-hydroxymethyl-3,5,6-trimethylpyrazine	<i>Drakaea livida</i>	Orchidaceae	32
(40) 2-hydroxymethyl-3,5-dimethyl-6-ethylpyrazine	<i>Drakaea micrantha</i>	Orchidaceae	9
(41) 2-hydroxymethyl-3-(3-methylbutyl)-5-methylpyrazine	<i>Drakaea livida</i>	Orchidaceae	33
(42) (3,5,6-trimethylpyrazin-2-yl) methyl 3-methylbutanoate	<i>Drakaea livida</i>	Orchidaceae	32
(43) (3,6-dimethylpyrazin-2-yl) methyl 3-methylbutanoate	<i>Drakaea livida</i>	Orchidaceae	32
(44) (3,5,6-trimethylpyrazin-2-yl) methyl (2 <i>S</i>)-methylbutanoate	<i>Drakaea livida</i>	Orchidaceae	32
(45) 2-(3-methylbutyl)-3,5,6-trimethylpyrazine	<i>Drakaea livida</i>	Orchidaceae	32
(46) (<i>E</i>)- <i>N</i> -(2-methylbutyl)-1-(pyridin-3-yl)methanimine	<i>Pyrus communis</i>	Rosaceae	10
(47) (<i>E</i>)- <i>N</i> -(3-methylbutyl)-1-(pyridin-3-yl)methanimine	<i>Pyrus communis</i>	Rosaceae	10
(48) 4-methyl-5-vinylthiazole	<i>Annona</i> , <i>Caladium bicolor</i>	Annonaceae, Araceae	34
(49) 2-(methylthio)-4-(hydroxy)phenol	<i>Caladenia crebra</i>	Orchidaceae	35
(50) 2-(methylthio)-4-(hydroxymethyl)phenol	<i>Caladenia crebra</i>	Orchidaceae	35
(51) 2-(methylthio)-4-(formyl)phenol	<i>Caladenia crebra</i>	Orchidaceae	35
(52) 2,3-dihydroxypropyl isovalerate	<i>Luisia teres</i>	Orchidaceae	36
Miscellaneous			
(53) 1,2-diacetin	Multiple	Multiple	19
(54) 1,3-diacetin	Multiple	Multiple	19
(55) 2,3-butanediol acetate	<i>Arum palaestinum</i>	Araceae	37
(56) acetoin acetate	<i>Arum palaestinum</i>	Araceae	37

3 Biosynthesis of floral volatiles

Recent progress in understanding the biosynthesis of floral volatiles has been dominated by reports of new genes and enzymes participating in volatile formation. Advances in DNA sequencing technology and the availability of genome and transcriptome data for many plant species have now made it possible to readily identify genes in floral volatile formation. Differential gene expression studies based on RNA-Seq allow researchers to focus on genes expressed specifically in floral organs and at the precise time of volatile emission. In addition, co-expression surveys can help to identify biosynthetic genes as long as one gene of a pathway is already known, since the genes of a single pathway are usually co-regulated. Meanwhile, increasing knowledge of the enzymes participating in floral scent formation has provided more target gene families to search for. Once candidate genes are chosen, advances in heterologous expression of the encoded enzymes both in model plants and in microbial systems for *in vivo* characterization have made it much easier to determine their biochemical function.

3.1 Fatty acid derivatives

The methylation of carboxylic acids is a common step in the biosynthesis of floral volatiles. These reactions are catalyzed by SABATH methyltransferases,³⁸ which carry out the *O*-methylation of small carboxylic acids to form common floral volatiles,

such as the benzenoids methyl salicylate and methyl benzoate. The methyl group itself is derived from *S*-adenosyl methionine (SAM). Recently, a SABATH *O*-methyltransferase was characterized that forms methyl hexanoate (57), the major constituent of floral scent of the water lily *Victoria cruziana* (Fig. 6).³⁹ In addition, a jasmonic acid *O*-methyltransferase was identified from *Cymbidium faberi* that produces methyl jasmonate (58), the major floral scent component of this orchid.⁴⁰

3.2 Terpenoids

The multitude of terpene carbon skeletons found in floral scents can nearly all be attributed to catalysis by terpene synthases. This large enzyme family converts acyclic prenyl diphosphates *via* carbocation-dependent reactions into an enormous variety of cyclic and acyclic products that are often released directly as floral volatiles or otherwise metabolized further. The great diversity of floral terpenes arises because each plant typically has a sizable group of different terpene synthases,⁴¹ plus some terpene synthases produce multiple products from a single substrate.⁴² Many new terpene synthases involved in floral scent formation have been characterized in the period under review, especially in horticulturally or economically important plants such as sweet osmanthus (*Osmanthus fragrans*, Oleaceae),⁴³ and species of the orchid genera *Dendrobium*^{44,45} and *Phalaenopsis* (moth orchids).⁴⁶ Although too numerous to mention individually, most of these



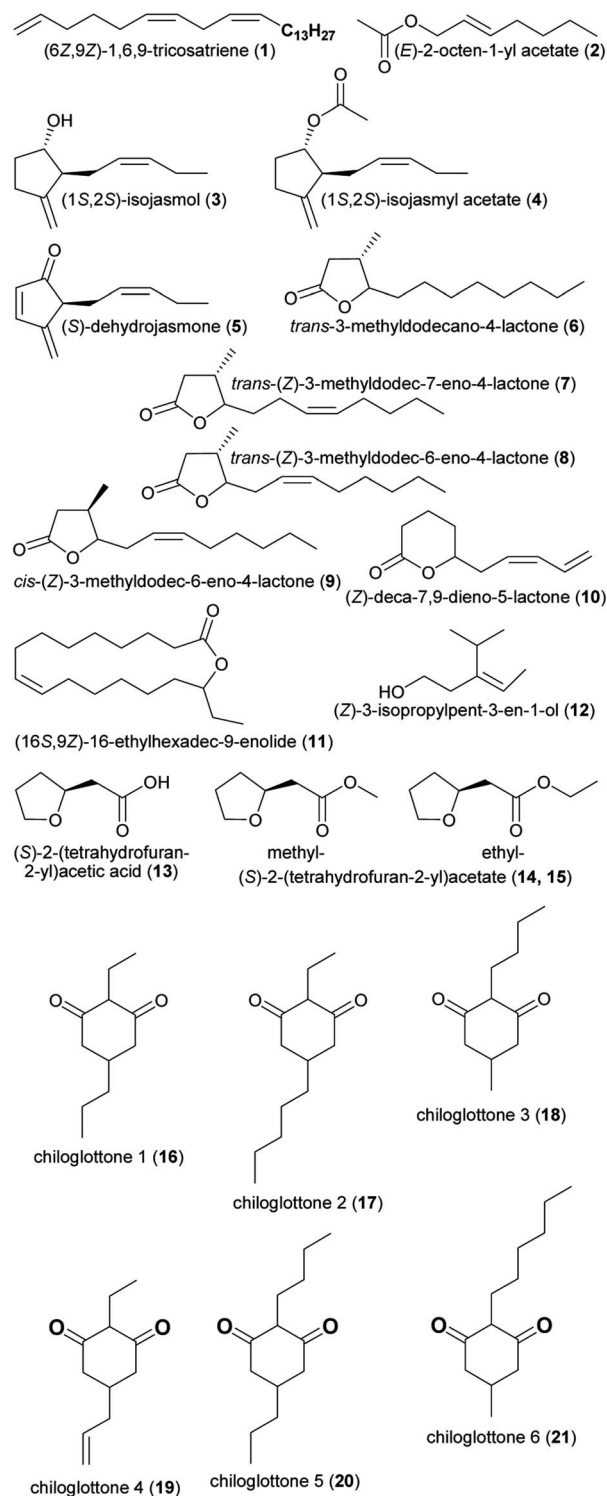


Fig. 1 Fatty acid derivatives listed in Table 1.

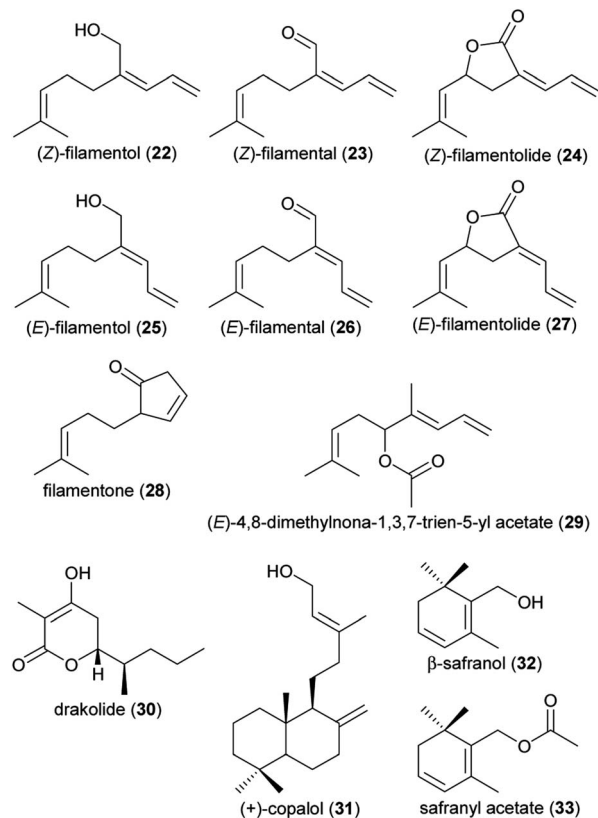


Fig. 2 Terpenoids listed in Table 1.

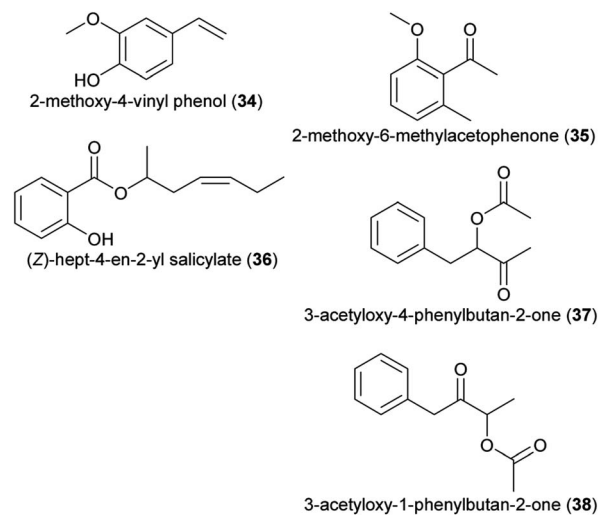


Fig. 3 Phenylpropanoids/benzenoids listed in Table 1.

terpene synthases make widespread floral volatiles for which terpene synthases are already known, like the monoterpenes geraniol (59),^{46,47} linalool (60 and/or 61),^{43,46,48–50} and (*E*)-β-ocimene (62)^{43,49} (Fig. 7), or the sesquiterpenes (*E,E*)-α-farnesene (63),⁴³ (*E*)-β-caryophyllene (64)⁴⁴ and (*E*)-α-bergamotene (65)⁵¹ (Fig. 8). These enzymes share their basic properties with other

terpene synthases making the same products, but often have divergent sequences. Terpene synthases making products such as geraniol, linalool and (*E*)-β-ocimene have originated independently many times over the course of plant evolution.⁵² However, other new terpene synthases make more unusual products, such as the sesquiterpene valerianol (66)⁵³ or unusual blends of products, such as the sesquiterpenes β-ylangene (67),





Fig. 4 Other amino acid derivatives listed in Table 1.



Fig. 5 Miscellaneous compounds listed in Table 1.

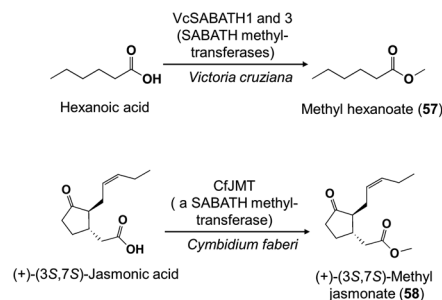


Fig. 6 Methylation of carboxylic acids in biosynthesis of volatile fatty acid derivatives in flowers.



Fig. 7 Examples of new monoterpene synthases from flowers.



Fig. 8 Examples of new sesquiterpene synthases from flowers.

β -copaene (68) and β -cubebene (69)⁵¹ (Fig. 9) or others.^{48–50} One group of monoterpene synthases from the flowers of *Nicotiana* species all make a similar mixture of seven products dominated by 1,8-cineole (70) with some α -terpineol (71) and lesser amounts of α -pinene (72), β -pinene (73), sabinene (74), limonene (75) and β -myrcene (76) (Fig. 10).⁵⁴ Interestingly, terpene synthase differences are at least in part responsible for the divergent pollination syndromes of two *Mimulus* (Phrymaceae) species: the hummingbird-pollinated *M. cardinalis* and the bumblebee-pollinated *M. lewisii*. The latter makes several monoterpenes, (E)- β -ocimene, limonene and myrcene, not produced by the former. These differences can be ascribed to two terpene synthases, an ocimene synthase that is functional





Fig. 9 Examples of new single and multi-product sesquiterpene synthases from flowers.

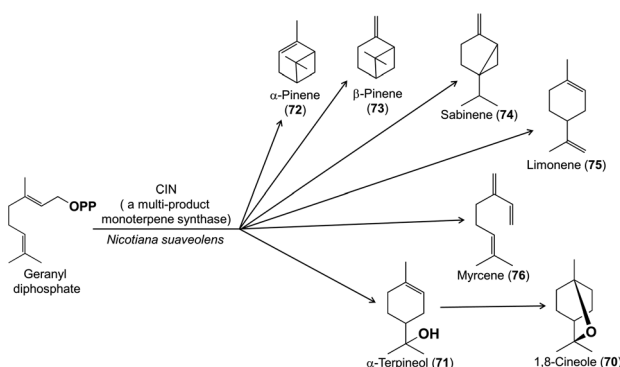


Fig. 10 A multi-product monoterpene synthase that synthesizes a mixture known as the "cineole cassette". 1,8-Cineole is formed via an enzyme-bound α -terpineol intermediate.

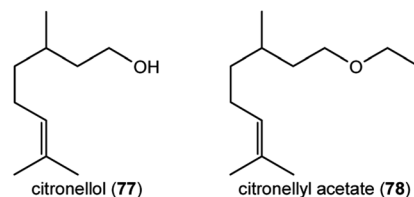
in *M. lewisii*, but not in *M. cardinalis*, and a limonene–myrcene synthase that is expressed at higher levels in *M. lewisii* than in *M. cardinalis*.⁵⁵

One of the most exciting advances of recent research on floral volatile biosynthesis is the discovery of a new pathway for the formation of the common monoterpene alcohol geraniol (59) in rose (*Rosa x hybrida*) flowers.⁵⁶ Geraniol in flowers has been previously shown to be made *via* terpene synthases,^{46,47} whose mechanisms involve simple cleavage of the diphosphate moiety from the precursor geranyl diphosphate with the resulting carbocation being quenched by water to form geraniol. Rose flowers were also assumed to have a geraniol-forming terpene synthase, but geranyl diphosphate was instead found to be hydrolyzed in a completely unexpected way:



Fig. 11 Formation of geraniol in roses *via* a new pathway involving a Nudix hydrolase instead of a terpene synthase.

a diphosphohydrolase from the Nudix family cleaves one phosphate moiety to give geranyl monophosphate, which is then further hydrolyzed to geraniol by a second phosphatase (Fig. 11). The role of this Nudix hydrolase was first suggested by QTL (Quantitative Trait Loci) mapping in crosses between rose genotypes with different scents, and confirmed by RNA interference of the corresponding gene, which reduced emission of geraniol.⁵⁶ More recently, a Nudix hydrolase from another rose species, *Rosa rugosa*, was identified, and expression of the encoding gene was shown to correlate with the formation of geraniol and derivatives, such as citronellol (77) and citronellyl acetate (78) as in *R. x hybrida*.⁵⁷



When terpene synthase products in flowers are further metabolized, the reactions involved are frequently oxidations carried out by cytochrome P450s, the largest family of enzymes in plant metabolism. A series of fascinating studies has described the role of cytochrome P450s in the oxidation of terpenes in *Arabidopsis thaliana* flowers. Although this species is largely self-pollinated, the flowers still emit a diverse blend of volatiles, especially terpenes. Their biosynthesis starts from geranyl diphosphate with flower-localized terpene synthases (TPS10 and 14) that produce the alcohols (–)– and (+)–linalool (60, 61), respectively. Next, these are converted by two flower-localized P450s (CYP71B31 and CYP76C3) to products with additional hydroxyl groups or epoxide and aldehyde functions that accumulate in the flowers (Fig. 12).⁵⁸ A third floral P450 in this species (CYP76C1) converts the linalools into an even larger range of products with alcohol, aldehyde, carboxylic acid and cyclic ether functions.⁵⁹ These extensively oxidized products act as herbivore deterrents, and thus the P450s can be considered as enzymes that transform linalool, frequently reported as a pollinator attractant, to herbivore defense compounds (but see below) in *A. thaliana*, a self-compatible plant that is not



Fig. 12 Biosynthesis of oxidized monoterpene volatiles in *Arabidopsis thaliana* flowers that function as herbivore deterrents. Oxidations are catalyzed by cytochrome P450 enzymes and only their most abundant products are shown.

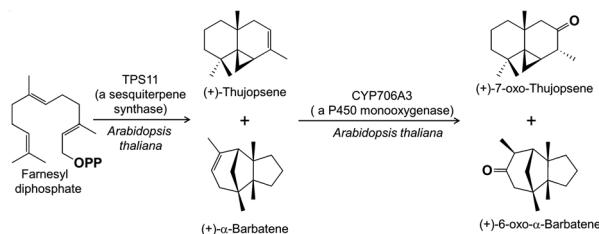


Fig. 13 Biosynthesis of oxidized sesquiterpene volatiles in *Arabidopsis thaliana* flowers that function as herbivore deterrents.

dependent on pollination for seed set. A parallel process involves sesquiterpenes in *A. thaliana* flowers with the olefins produced by terpene synthase TPS11 being oxidized to alcohols and ketones by yet another P450 enzyme (CYP706A3) (Fig. 13).¹⁶ In this case, the oxidized products again accumulate in flowers and serve as anti-herbivore defenses, in contrast to the more volatile sesquiterpene olefin substrates, which are emitted as attractants.

Cytochrome P450 enzymes have also been implicated in the formation of two irregular terpenes, the C_{16} (*E,E*)-4,8,12-trimethyltrideca-1,3,7,11-tetraene (TMTT) (79) and the C_{11} (*E*)-4,8-dimethyl-1,3,7-nonatriene (DMNT) (80), which are frequent constituents of floral scent, especially in night-blooming white flowers. TMTT and DMNT are also formed in the vegetative parts of plants after herbivore damage. Biosynthetic studies on these two compounds carried out with damaged *A. thaliana* and maize foliage demonstrated that the C_{20} alcohol geranylgeranyl, a terpene synthase product, was cleaved to form TMTT, and the C_{15} nerolidol cleaved to form DMNT, respectively (Fig. 14A).^{60,61} The enzymes involved in both plants are cytochrome P450s (CYP82G1 in *A. thaliana*, CYP92C5 and C6 in

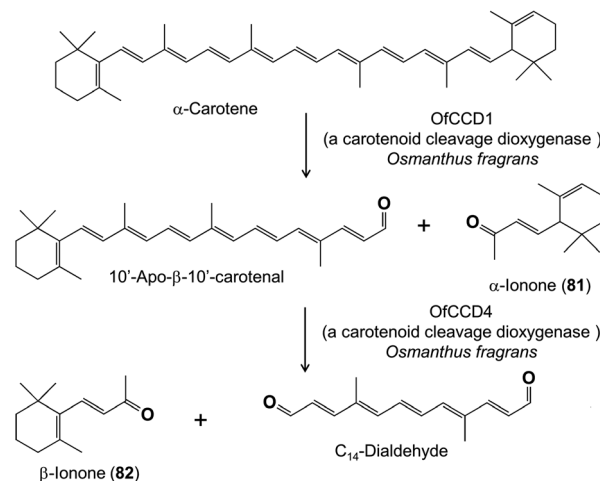


Fig. 15 Cleavage of a carotenoid to volatile apocarotenoids by carotenoid cleavage dioxygenase enzymes.

maize), but are from different subfamilies suggesting independent evolution of the pathway in different plant lineages. Given that the *A. thaliana* CYP82G1 is expressed after herbivore damage in leaves as well as constitutively in flowers, TMTT and DMNT seem very likely to be formed by similar pathways in flowers. *Arabidopsis thaliana* also possess a second completely different DMNT pathway in roots involving the cleavage of a triterpene (C_{30}) precursor by yet another cytochrome P450 (CYP705A1) to form the C_{11} product (Fig. 14B).⁶²

Another group of floral terpenes formed by cleavage reactions are those derived from carotenoids by catalysis of carotenoid cleavage dioxygenases (CCDs) instead of P450s. Here there are reports of the floral volatiles α - and β -ionone (81, 82) being produced from α - and β -carotene in *Osmanthus fragrans* by a CCD1 enzyme,⁶³ while β -ionone is produced by a CCD4 enzyme from β -carotene and lutein in *Crocus sativus*, the source of saffron⁶⁴ (Fig. 15).

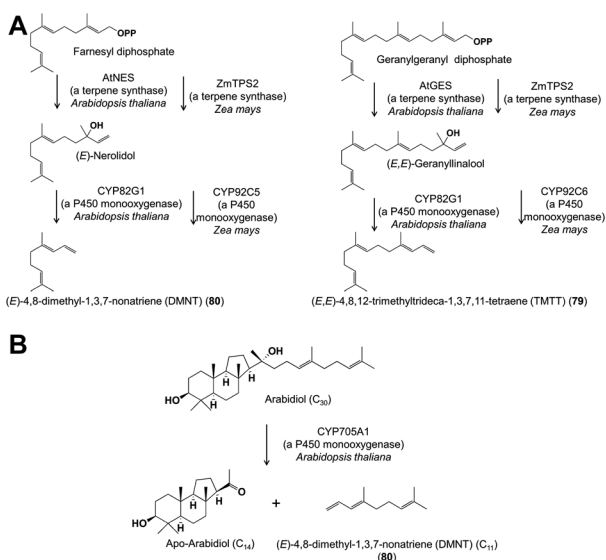


Fig. 14 (A) Biosynthesis of DMNT (80) and TMTT (79), two homoterpenes that are often constituents of floral volatile blends. (B) Biosynthesis of the volatile DMNT by a second pathway in the roots of *Arabidopsis thaliana*.

3.3 Phenylpropanoids and benzenoids

The biosynthesis of most phenylpropanoid and benzenoid floral volatiles from intermediates of primary metabolism typically requires longer pathways than those to terpenoid floral volatiles. One of the exceptions is the pathway leading from phenylalanine to the widely-encountered phenylacetaldehyde (83). This is a two-step decarboxylation-amine oxidation reaction catalyzed by phenylacetaldehyde synthase in a pyridoxal 5'-phosphate-dependent manner (Fig. 16). Described originally from petunia flowers,⁶⁵ this enzyme of the aromatic amino acid decarboxylase family has also been reported from the flowers of *A. thaliana* and rose (*R. x hybrida*).^{66,67}

Phenylacetaldehyde is next subject to reduction to form 2-phenylethanol (84), a well-known component of rose scent. The responsible enzyme, a member of the short chain dehydrogenase/reductase family previously characterized from *R. x hybrida* 'Hoh-Jun' flowers⁶⁸ and tomato fruit, was more recently characterized in detail from *R. x damascena* flowers (Fig. 17).⁶⁹ A QTL study employing a *R. x hybrida* mapping



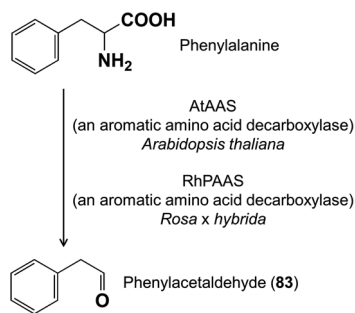


Fig. 16 Biosynthesis of phenylacetaldehyde from phenylalanine occurs via a two-step reaction catalyzed by phenylacetaldehyde synthase, a member of the aromatic amino acid decarboxylase family.

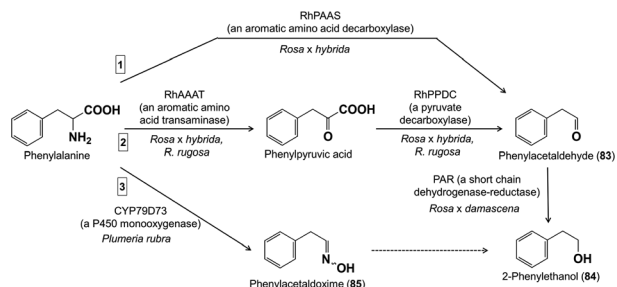


Fig. 17 2-Phenylethanol is formed by three different biosynthetic routes, and more than one may operate in a single plant species depending on the environmental conditions. The steps between phenylacetaldoxime and 2-phenylethanol are not known. Other floral volatiles are also formed by different pathways in different species or by similar pathways employing different enzymes.⁵²

population segregating for 2-phenylethanol emission implicated an allele of phenylacetaldehyde synthase as being critical for 2-phenylethanol formation.⁷⁰ Curiously, expression of the encoding gene was found to be active already in the flower bud stage causing 2-phenylethanol to accumulate in the petals as a glycoside. A UDP-glucosyltransferase responsible for glucosylation of 2-phenylethanol (and benzyl alcohol) was recently identified in petunia.⁷¹

A second pathway to 2-phenylethanol in roses (*R. x hybrida*) also starts from phenylalanine. This aromatic amino acid is first deaminated to phenylpyruvate by an aminotransferase, then decarboxylated to phenylacetaldehyde by a phenylpyruvate decarboxylase, and finally reduced as above to 2-phenylethanol (Fig. 17).⁷² Studies with isotopically-labeled phenylalanine demonstrated that the route *via* phenylpyruvate operates especially in summer due to higher temperatures. The authors suggest that the greater efficiency of the pathway *via* phenylpyruvate helps maintain 2-phenylethanol formation in summer despite reduced petal growth. Both pathways also seem to operate in *R. rugosa* flowers since the respective genes were expressed and overexpression of either the phenylacetaldehyde synthase gene or the phenylalanine aminotransferase gene increased 2-phenylethanol formation.⁷³

And, there is even a third pathway for 2-phenylethanol biosynthesis in flowers! In *Plumeria rubra* (frangipani,

Apocynaceae), phenylalanine is oxidized by a cytochrome P450 of the CYP79 family (CYP79D73) to phenylacetaldoxime (85), which was shown to be an intermediate in 2-phenylethanol formation by *in vivo* feeding (Fig. 17).⁷⁴ Expression of the CYP79D73 gene in *Nicotiana benthamiana* also resulted in 2-phenylethanol emission, but the biosynthetic steps from phenylacetaldoxime to 2-phenylethanol are not yet known. The evolution of three different pathways to 2-phenylethanol in flowers might suggest that it is easy to evolve a pathway to this compound or that it has a strong innate attractiveness for pollinating insects (see below). Multiple pathways for 2-phenylethanol formation are also known in poplar leaves.⁷⁵

Along with 2-phenylethanol, the volatile 1-phenylethanol (86) also occurs in flowers, such as those of *Camellia sinensis* (tea). Isotopic tracer experiments demonstrated that 1-phenylethanol is derived from phenylalanine *via* the intermediate acetophenone (Fig. 18).⁷⁶ Several genes and enzymes involved in the conversion of acetophenone to 1-phenylethanol have been identified in tea from the short chain dehydrogenase/reductase family.^{77–79} Interestingly, 1-phenylethanol is present as both (*R*)- and (*S*)-enantiomers and separate enzymes are responsible for making each enantiomer.

Another well-known group of phenylpropanoid volatiles are the phenylpropenes, including eugenol, chavicol and estragole, found in flowers and the foliage of certain herbs. Their biosynthesis leads from phenylalanine to the phenylpropanoids, which are then acetylated to serve as substrates for an NADPH-dependent reductase that forms the final phenylpropene by acetate elimination and reduction (Fig. 19).⁸⁰ Since this original report, genes participating in other steps in phenylpropene formation have been identified, such as in *O*-methylation of caffeoyl-CoA to feruloyl-CoA in petunia,⁸¹ the acetylation of coniferyl alcohol in *Prunus mume*,⁸² and the conversion of coniferyl acetate to eugenol (87) in roses.⁸³

The formation of benzenoid volatiles requires the shortening of the C₃ side chain of their phenylpropanoid precursors. Our understanding of this biosynthetic process has increased significantly in recent years thanks largely to work in petunia flowers as well as *A. thaliana*, but curiously several pathways may co-occur (reviewed in⁸⁴). Evidence is strongest for the operation of a β -oxidation pathway from cinnamic to benzoic acid that removes a C₂ fragment from the C₃ side chain in a manner analogous to the β -oxidation of fatty acids. The reaction sequence includes hydration of the C=C bond of the side chain, oxidation of the resulting alcohol to give a β -keto

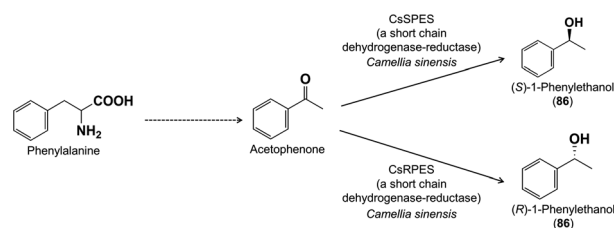


Fig. 18 The formation of 1-phenylethanol. Only the last step of the pathway has been elucidated in plants.





Fig. 19 The biosynthesis of the phenylpropene volatile eugenol (87) has been previously elucidated. Shown are new reports of enzymes involved in the biosynthetic pathway.

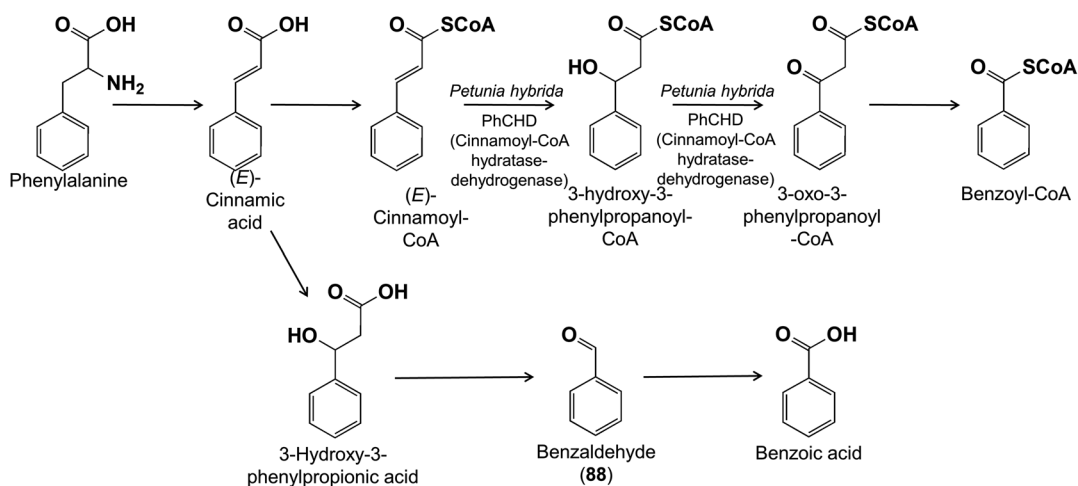


Fig. 20 Pathways for the formation of benzenoid flower volatiles. The β -oxidation pathway (upper row) proceeds via CoA-bound intermediates in the peroxisome. The most recently characterized enzyme is a cinnamoyl-CoA hydratase-dehydrogenase, which catalyzes consecutive steps. A non-oxidative pathway (lower row) with free or CoA-bound intermediates occurs in the cytosol.

function, and a reverse aldol cleavage to release the C_2 fragment as acetyl-CoA (Fig. 20). These steps proceed via CoA-bound intermediates in the peroxisome. The most recently characterized enzyme was the cinnamoyl-CoA hydratase-dehydrogenase, which catalyzes the consecutive hydration and oxidation steps, thereby filling the last gap in this pathway.⁸⁵

A non-oxidative pathway for benzenoid formation is also known that follows a similar chemical logic, but the reverse aldol cleavage leads to benzaldehyde (88) rather than benzoyl-CoA, which is then oxidized to benzoic acid (Fig. 20). The non-oxidative sequence occurs in the cytosol with free or CoA-bound intermediates. According to flux data from petunia flowers, both the β -oxidative and non-oxidative pathways participate in the formation of benzenoid volatiles with their contributions depending on light conditions and time of day.⁸⁴

A benzenoid with two *O*-methyl groups is veratrole (1,2-dimethoxybenzene) (89) found in the flowers of *Silene latifolia*

(Caryophyllaceae). Feeding of potential precursors established that this benzenoid is derived from phenylalanine via (*E*)-cinnamic acid, benzoic acid and salicylic acid.⁸⁶ Decarboxylation results in catechol,⁸⁷ which is then *O*-methylated sequentially to give 1,2-dimethoxybenzene (Fig. 21).^{86,87} The activities of the two *O*-methylation steps have been demonstrated,^{86,87} and one of the genes identified.^{86,88}

An *O*-methyltransferase has also been identified from the flowers of loquat (*Eriobotrya japonica*, Rosaceae) that reacts preferentially with phenylpropanoids and benzenoids containing *para*-OH and *meta*-OCH₃ functions *in vitro* and methylates the free OH to add a *para*-OCH₃ group to this ring system.⁸⁹ Loquat flowers also possess a carboxylate-methylating activity from the SABATH family that converts *p*-methoxybenzoic acid to its corresponding methyl ester (Fig. 22).⁹⁰ The enzyme has broad specificity and also methylates benzoic acid and jasmonic acid. Meanwhile, a BAHD acyltransferase was identified in *Jasminum*





Fig. 21 The biosynthetic pathway to veratrole (1,2-dimethoxybenzene). While the gene and enzyme have only been characterized for the last step, the roles of other intermediates have been established by administration of isotopically-labeled precursors.



Fig. 22 Other biosynthetic conversions of benzenoid floral volatiles.

sambac flowers that reacts with benzyl alcohol and acetyl-CoA to form benzyl acetate (90) (Fig. 22).⁹¹

A more elaborate benzenoid is benzylacetone (91) from the flowers of *Nicotiana attenuata*. Transcript analysis of RNA isolated from single corolla cells and gene co-expression studies helped suggest a biosynthetic pathway that could be confirmed by silencing candidate genes.⁹² Based in part on previous studies on benzylacetone⁹³ and other phenylbutanoids,⁹⁴ the pathway was found to proceed from phenylalanine via (*E*)-cinnamic acid and (*E*)-cinnamoyl-CoA. Next comes a single carbon extension: a condensation reaction of (*E*)-cinnamoyl-CoA and malonyl-CoA catalyzed by a polyketide synthase with an additional loss of CO₂ (besides the typical loss of one molecule of CO₂ from malonate in polyketide synthase catalysis). This is followed by a final reduction to give benzylacetone (Fig. 23).



Fig. 23 The biosynthesis of the floral volatile benzylacetone.



Fig. 24 The biosynthesis of several nitrogen-containing floral volatiles derived from phenylalanine. The formation of 2-phenylacetone nitrile has only been demonstrated in vegetative tissue to date. The pathway to 2-phenylnitroethane has not yet been elucidated.

3.4 Other amino acid derivatives

Relative to other classes of floral volatiles, few reports have appeared on the biosynthesis of amino acid derivatives. Most of these feature aldoximes, well known from the floral scent of species of *Oenothera* (Onagraceae)⁹⁵ and *Angraecum sesquipedale* (Orchidaceae),⁹⁶ and as herbivore-induced volatiles from vegetative plant parts and biosynthetic intermediates in the formation of anti-herbivore defenses such as cyanogenic glycosides. Aldoximes are all derived from amino acids under the catalysis of the CYP79 family of cytochrome P450 enzymes.^{97,98} The most common substrates are the branched chain amino acids, valine, leucine and isoleucine, as well as phenylalanine and tyrosine, while both (*E*)- and (*Z*)-configurations are present in the products. Once formed, the isomers of phenylacetaldoxime (85) are potential precursors of other nitrogen-containing floral volatiles, including 2-phenylacetone nitrile (benzyl nitrile, benzyl cyanide) (92) and 2-phenylnitroethane (93), based on feeding experiments in *Plumeria rubra* flowers (Fig. 24).⁷⁴ The conversion of phenylacetaldoxime to 2-phenylacetone nitrile is catalyzed by another family of P450 enzymes, the CYP71s, though this has only been demonstrated in connection with vegetative volatiles⁹⁹ and cyanogenic glycoside formation,⁹⁸ and not yet in flowers. Floral phenylacetaldoximes are also precursors of 2-phenylethanol and other benzenoid compounds in flowers, as described above in 3.3, but the reactions involved have not yet been elucidated.⁹⁶



Intriguing reports describe an amino acid-derived floral volatile that also has a role in the biosynthesis of a floral pigment. The volatile indole (94), derived from the tryptophan biosynthetic pathway, is released directly from the flowers of certain poppy species,¹⁰⁰ and also appears to combine with flavonoids to form the unique pigments called nudicaulins¹⁰¹ via reactions that still need investigation.



4 Regulation of floral volatile emission

In the period under review, nearly as many new findings have been published on the regulation of floral volatile biosynthesis as on the biosynthetic pathways themselves. Knowledge about regulation has given new insights into how floral volatile pathways operate in the plant as well as providing additional tools to manipulate volatile emission.

4.1 Transcription factors

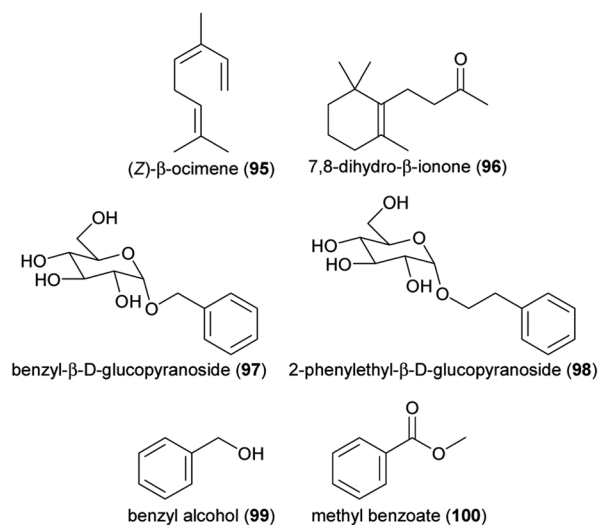
The majority of the regulatory studies have focused on transcription factors, which have been found to be critical in controlling the time and place of volatile biosynthesis and hence emission in flowers. Transcription factors limit synthesis to particular floral organs, stages of development and times of day. The first transcription factor identified as controlling floral volatile emission was *ODORANT1* (*ODO1*) from petunia (*Petunia x hybrida*), which stimulates the formation of the phenylpropanoid and benzenoid compounds that dominate petunia scent.¹⁰² First discovered by comparing the transcripts of scented and non-scented petunia lines, this R2R3-MYB family transcription factor was also identified as a regulatory element in a quantitative genetics study involving crosses between two petunia species with big difference in their floral volatile spectra.¹⁰³ After *ODO1* was reported, a second MYB transcription factor, *EMISSION OF BENZENOID II* (*EOBII*), was observed to stimulate volatile biosynthesis in petunia and its expression was also correlated with the peak of emission.¹⁰⁴ Research revealed that *EOBII* stimulates *ODO1* expression by binding to the *ODO1* promoter.¹⁰⁵ Further complexity comes from a third MYB transcription factor in petunia flowers, *EOB1*, which is upstream of *EOBII* and *ODO1* and activates both of them, while *ODO1* inhibits *EOB1* expression.¹⁰⁶ These transcription factors activate genes of phenylpropanoid and benzenoid biosynthesis as well as earlier metabolic steps, such as the shikimate pathway, which produces phenylalanine, the precursor to nearly all petunia flower volatiles.^{104,106,107} Still other MYB transcription factors seem to be involved in stimulating volatile formation in petunia flowers that target genes other than biosynthetic genes.¹⁰⁸ Petunia transcription factors have also been discovered that negatively regulate floral emission. A member of the ethylene response factor (ERF) family binds to *EOBI* and so inhibits its binding to promoters of floral biosynthetic genes.¹⁰⁹ Hence ethylene downregulates volatile formation in petunia. Other hormones that suppress volatile formation include gibberellins and auxin. Gibberellin signaling is blocked by a *DELLA* transcription factor that activates *EOBII*,¹¹⁰ while auxin suppresses volatile emission by decreasing phenylalanine levels.¹¹¹

MYB transcription factors regulate floral volatile formation in other plant species as well. A MYB called *PAP1*, known to activate phenylpropanoid biosynthesis in *A. thaliana*, increases phenylpropanoid volatile formation in roses.¹¹² Terpenoid volatiles are also influenced by MYB transcription factors in plants, such as roses,¹¹² *Freesia x hybrida* and *A. thaliana*.¹¹³ For

example, several different MYB factors regulate the formation of terpene synthase products in *F. x hybrida* and *A. thaliana* by binding directly to the promoters of terpene synthases. An antagonistic effect on volatile formation is caused by the basic helix-loop-helix (bHLH) transcription factor *MYC2*, which blocks MYB induction of monoterpene synthases, but stimulates expression of sesquiterpene synthases.¹¹³ The expression of *MYC2* in turn is induced by both jasmonic acid and gibberellin signaling pathways.¹¹⁴ In the orchid *Dendrobium officinale*, a jasmonate-responsive bHLH transcription factor was also reported to regulate the expression of a monoterpene synthase gene, but here the bHLH transcription factor promoted the expression of this gene, which encodes linalool (**60** and **61**) formation, instead of blocking it.¹¹⁵

An increasingly popular species for floral volatile studies is the white ginger lily (*Hedychium coronarium*), which produces a range of terpenoid and benzenoid volatiles. MYB transcription factors also play a role in regulating volatile emission here.¹¹⁶ For example, *H. coronarium* MYB factors were identified that bind to the promoter regions of floral biosynthetic genes after their expression is activated by auxin.¹¹⁷ Another *H. coronarium* transcription factor binds to the promoter region of an (*E*)- β -ocimene synthase,¹¹⁸ which produces a major floral volatile of *H. coronarium*.

Terpenoids, such as linalool (**60** and **61**), (*Z*)- β -ocimene (**95**) and β -ionone (**82**), are dominant floral volatiles in another species well-investigated for floral volatiles, sweet osmanthus (*Osmanthus fragrans*). Here, genomic and transcriptomic studies suggested that R2R3-MYB and WRKY transcription factors play a role in controlling volatile emission.^{119,120} MYB transcription factors were reported to be involved in regulating terpene emission in a cultivar of lily (*Lilium* sp.) as well, with one MYB demonstrated to bind to the promoter region of an ocimene synthase.¹²¹ Lilies also contain a MYB similar to petunia *ODO1* that upregulates genes involved in volatile phenylpropanoid and benzenoid biosynthesis and the shikimate pathway.¹²²



4.2 Other factors regulating biosynthesis

The pattern of gene expression is usually assumed to exert a dominant influence on the quantity and composition of floral volatiles formed. Support for this assertion comes not only from the many transcript studies and the impact of transcription factors, but also from the results of artificial selection experiments on the floral volatiles of the fast-cycling *Brassica rapa* plants.¹²³ After only three generations of selection for increased emission, elevated amounts of phenylpropanoid and benzenoid volatiles were evident and these were shown to arise from elevated expression of genes of the phenylpropanoid pathway and earlier stages in biosynthesis, such as the shikimate pathway, which supplies the precursor phenylalanine. This amino acid is often a limiting factor in phenylpropanoid and benzenoid volatile formation as emphasized by construction of transgenic petunia lines with increased phenylalanine supply due to overexpression of a bacterial shikimate pathway enzyme that is insensitive to feedback inhibition from phenylalanine.¹²⁴ These lines not only accumulated higher amounts of phenylalanine but also emitted higher amounts of volatiles than wild-type lines. Moreover, direct feeding of phenylalanine to cut flowers also increased the emission of phenylalanine-derived volatiles.¹²⁵ The importance of primary metabolite precursors in regulating flower volatile formation was also highlighted by research on a *Lilium* cultivar where glucose and fructose application increased the expression of genes for terpenoid volatile formation and terpenoid emission.¹²⁶ There was a regulatory context in this study too, since sugar kinases found to be highly expressed in flowers increased volatile emission when overexpressed and decreased emission when silenced.

Gene expression is also controlled by epigenetic influences, including DNA methylation, histone modification and chromatin remodeling. Using the petunia model system, research has found that histone acetylation during flower anthesis facilitated the activation of specific genes for the biosynthesis of the phenylpropanoid and benzenoid volatiles of petunia flowers.¹²⁷ Included in this list are genes encoding the main steps of the phenylpropanoid pathway plus those for the shikimate pathway. At the same time, genes encoding enzymes in pathways that compete for phenylalanine, such as those involved in the formation of flavonoids and lignin, were repressed. Long, non-coding RNAs may also influence gene expression at transcriptional and post-transcriptional stages. A recent investigation on roses implicated several long, non-coding RNAs in controlling floral volatile release from *Rosa x hybrida*.¹²⁸ Silencing of one such sequence increased the emission of the apocarotenoid, 7,8-dihydro- β -ionone (**96**).

4.3 Factors regulating storage, transport and emission

Once synthesized, floral volatiles have been generally assumed to be emitted from floral tissues into the atmosphere by simple diffusion through cell membranes. Thanks to recent breakthroughs in our knowledge, however, it is now clear that storage and transport can also regulate emission, while simple diffusion plays only a minor role. Based on the chemical properties of typical floral volatiles, Natalia Dudareva and coworkers¹²⁹

calculated that simple physical diffusion would require such high concentrations to build up in volatile-producing cells that these would become toxic, suggesting emission must occur instead *via* biological carriers. The Dudareva laboratory did indeed discover such a carrier, identifying a transport protein located in the plasma membranes of petunia flower cells that exports volatiles.¹³⁰ As a member of the ATP-binding cassette (ABC) transporter family closely related to previously described wax transporters, this protein was found to be specific for benzenoids. It was present almost exclusively in the cells of open flowers, and expression of the corresponding gene was controlled like so many other volatile biosynthetic genes in petunia by the transcription factor *ODORANT1*. Interestingly, silencing of the transporter not only reduced emission and increased accumulation of volatiles in flowers, but also caused signs of cell damage due to the build-up of volatiles to toxic levels in living cells.

After volatiles traverse the plasma membrane, the cuticle of floral organs also acts as a barrier to emission. The cuticle, however, has an unexpectedly complex role in the emission process serving also as a sink for volatiles prior to emission that protects cells from toxicity. Curiously, when cuticle thickness in petunia was reduced by silencing a petal wax transporter, the resulting flowers had lower rather than greater volatile emission rates since the reduction in sink size exerted negative feedback on volatile biosynthesis.¹³¹ Cuticular resistance was greater for petunia volatiles of lower volatility, such as benzyl alcohol (**99**) and eugenol (**87**), which have large storage pools relative to their emission rates. In contrast, higher volatility compounds, such as methyl benzoate (**100**) and benzaldehyde (**88**), have smaller storage pools relative to emission rate and emission more closely matches their rate of biosynthesis. The existence of additional control points between floral volatile synthesis and emission was suspected for many years based on the isolation of glycosylated volatiles from various plant organs. From petunia flowers, for example, glycosylated phenylpropanoids and benzenoids, including benzyl glucoside (**97**) and 2-phenylethyl glucoside (**98**) have been reported.¹³² Time course studies and isotopic labeling demonstrated that these glycosides form dynamic pools, and so likely represent storage forms of volatiles with glycosylation and glycoside hydrolysis helping to modulate release rates.

Dudareva and coworkers also discovered an unexpected new dimension in volatile storage and transport between the organs of petunia flowers.¹³³ A blend of sesquiterpene volatiles produced by the petal tube were found to be released inside the bud and to accumulate on the thick waxy cuticle of the stigma prior to floral opening. The compounds, germacrene D (**101**), bicyclgermacrene (**102**), and germacrene D-4-ol (**103**), the major products of a single sesquiterpene synthase expressed in the petal tube, were all mobilized to the stigma where they were shown to promote the development of this organ and enhance subsequent seed yield compared to plants in which this terpene synthase was silenced by RNA interference. These sesquiterpenes may also act to reduce the pathogen load on stigmas. Hence, greater understanding of floral volatile transport and storage can furnish significant new insights on the roles of



floral volatiles in the plant. These phenomena can also explain some of the inconsistencies between the results of headspace sampling and solvent extraction, and provide reasons for why expression of biosynthetic genes is sometimes poorly correlated with emission.



5 Effects of abiotic and biotic factors on floral scent emissions

Though floral scents are regarded as being species-specific,⁴ they also vary among individuals within species.¹³⁴ Some of this variation has a genetic basis,¹³⁵ while other variation is due to phenotypic plasticity and is shaped by the environment.¹³⁶ Both biotic and abiotic factors affect floral scents,¹³⁷ and here, we, focus on recent studies that demonstrated effects of air pollutants, increased temperature and drought, soil nutrient availability, bacterial, fungi and herbivores on floral scent emissions and on the chemical communication with pollinators.

5.1 Air pollution

Air pollutants result in lower abundances of pollinators and smaller numbers of flower visits, with negative effects on the seed set of plants.¹³⁸ One mechanistic explanation for this finding is that air pollutants affect floral scents, the perception of floral scents by pollinators, and thus, the olfactory communication between plants and pollinators. Such effects have been studied over the last decade by fumigating growing plants and pollinators with pollutants (e.g., ozone, diesel exhaust), or by applying the pollutants to scents after their release by the flowers.¹³⁹

Elevated ozone exposure during growth can affect the emission of some floral volatiles, as shown for a few plant species. After 35 days of elevated ozone exposure, roses (*Rosa* spp.) released higher amounts of sesquiterpenes, while monoterpene emissions were not affected.¹⁴⁰ Following 90 days of exposure, the total amount of scent decreased, mainly due to decreased emissions of monoterpenes. There was no effect on total sesquiterpene emissions following this long-time exposure, despite an increase in the sesquiterpene guaiene (**104**).¹⁴⁰ Sautier and Blande¹⁴¹ exposed four Brassicaceae species to elevated ozone levels. None of them showed changes in total scent emissions; however, the emission rates of single compounds changed in three of the four species. The exposure to elevated ozone levels resulted in higher amounts of α -phellandrene (**105**) in *Brassica nigra* (black mustard), decreased emission or even a loss of *p*-cymene (**106**) in *Sinapis alba* (white mustard), and a decreased emission of α -pinene (**72**) and losses of β -thujone (**107**) and camphor (**108**) in *Brassica napus* (oilseed rape). Elevated ozone exposure did not affect the emission of single compounds in *S. arvensis* (wild mustard).



A recent study showed that increased CO₂ levels (800 ppm, predicted as the atmospheric concentration for 2100) also result in altered floral scents.¹⁴² CO₂ fertilization decreased total scent in *Campanula rotundifolia* (Scottish bluebell) and increased total scent in *Phacelia hastata* (silverleaf phacelia). Furthermore, scent composition was altered following CO₂ enrichment. In *C. rotundifolia* 6-methyl-5-hepten-2-one (**109**), β -pinene (**73**) and β -myrcene (**76**), and in *P. hastata* linalool (**60** and/or **61**), limonene (**75**) and (*Z*)-3-hexenyl acetate (**110**) were emitted at higher rates from CO₂-fertilized plants compared to control plants. In two other study plants (*Heterotheca villosa*, the hairy goldenaster and *Potentilla recta*, the sulphur cinquefoil), CO₂ enrichment affected neither total scent emission nor scent composition.



In the examples above, plants were exposed for some time to air pollutants, while floral volatiles were collected at ambient conditions. Other studies, however, exposed the released floral volatiles directly to air pollutants. Volatiles are exposed to many airborne compounds after their release from the flower, among them pollutants in the atmosphere. Several air constituents (e.g., ozone, NO₃ radicals) are known to directly interact with floral volatiles and change the composition of scent blends.^{143–148} Recent advances have been made in quantifying the effects of ozone and diesel exhaust on floral volatiles and how this influences pollinator behavior. Studies on four Brassicaceae species demonstrated that ozone partly degrades some of the floral scent compounds released by the flowers, while others were not affected by ozone.^{141,149,150} In *Brassica nigra*, for example, strong negative effects were recorded for *p*-anisaldehyde (**111**), phenol (**112**) and *p*-cymene (**106**), as well as for total monoterpenes. Overall, increased levels of ozone affected the relative scent composition in this species, resulting in a decreased attractiveness of floral scent to bumblebee pollinators.¹⁴⁹ Negative effects of increased ozone levels on the attraction of pollinators were also shown for the tobacco hawkmoth *Manduca sexta* and one of its preferred host plants, the jasmine tobacco *Nicotiana glauca*.¹⁵¹ Here, increased ozone levels again affected flower scent composition resulting in drastically reduced amounts of linalool (**60** and/or **61**), (*E*)-nerolidol (**113**), and 3-methylbutylaldehyde (**114**), but in increased



amounts of decanal (**115**). These changes in scent composition reduced the time hawkmoths spent at the ozone-altered scent sources in wind tunnel bioassays. However, moths were able to learn the ozone-altered blend, which might reduce the negative impact increased ozone levels have on the communication in this pollination system.¹⁵¹ Ozone also affected the flower–florivore interaction between *Cucurbita foetidissima* (buffalo gourd) and the striped cucumber beetle (*Acalymma vittatum*). The floral scents, which were not chemically analyzed in detail, were less attractive to the beetle when they were exposed to increased ozone levels.¹⁵²



Diesel exhaust (mainly due to nitrogen oxides) affected different well known floral scent compounds differently, based on experiments with two synthetic blends consisting overall of eleven terpenoids and five aromatic compounds.^{153,154} Exhaust had negative effects on seven terpenoids and two aromatic compounds. The compounds α -terpinene (**116**), (*E,E*)- α -farnesene (**63**), β -ocimene (**62**) and (*E*)- β -caryophyllene (**64**) were most severely affected, as they became undetectable when exposed to diesel exhaust. **64** was completely transformed into its *cis*-isomer. The amount of only one compound [*p*-cymene (**106**)] increased during the experiments, whereas the other six compounds [*e.g.*, α -pinene (**72**), benzaldehyde (**88**)] were not affected by diesel exhaust. The alterations of the two diesel exhaust blends reduced the ability of Western honeybees (*Apis mellifera*; henceforth honeybee) to remember the original blend to which they had been trained.^{153,154}

The chemical degradation of floral volatiles depends on their reactivity with respect to the individual pollutants (*e.g.*, ozone, NO₃ radicals).¹⁴⁶ Some of the above described results can indeed be explained by the reactivity of the volatiles (*e.g.* strong ozone effects on linalool), but others not (strong ozone effects on *p*-cymene), and more research is needed to better understand the observed effects of air pollutants on floral volatiles. Furthermore, it would be interesting to know whether selection is currently occurring on pollinators to rely more on compounds less reactive during the Anthropocene.



Studies on how the perception, learning and memory of flower volatiles is affected when insects are exposed to air pollutants were also performed using ozone. Exposure of honeybees, bumblebees (*Bombus terrestris*) and fig wasps (*Blastophaga psenes*) to increased levels of ozone had variable effects on the perception of volatiles in the olfactory circuitry. The strength of the physiological response to a specific compound either decreased [*e.g.*, linalool (**60** and/or **61**) in fig wasps, benzaldehyde (**88**) in bumblebees], increased [*e.g.* benzyl alcohol (**99**) after short-time ozone exposure of fig wasps], or was not affected [*e.g.*, linalool (**60** and/or **61**) and 2-phenylethanol (**84**) in honeybees; **99** after long-time ozone exposure of fig wasps].^{155–157} In honeybees, ozone exposure not only influenced the perception of floral volatiles, but also resulted in less efficient olfactory learning.¹⁵⁵

5.2 Temperature and drought

Due to climate change, global mean surface temperatures and the rate as well as the intensity of droughts are increasing across many regions worldwide,¹⁵⁸ effecting plant reproduction¹⁵⁹ and floral scents (reviewed in Borghi *et al.*,¹⁵⁹ Gérard *et al.*¹⁶⁰). Recent studies confirmed previous results and consistently found that total scent emissions in various plant species are temperature dependent, with maximum emissions typically between 25 °C and 30 °C.^{161–165} However, in some species flowering during summer in Mediterranean climates, scent maxima were reached at temperatures as high as 40 °C.¹⁶³ In addition to the variation in total scent, temperature also effected scent composition. Both the number of compounds released^{163,165} and the absolute as well as relative amounts of single compounds varied among different temperatures.^{161–163,165–167} The finding that different compounds respond differently to temperature could be traced back to the activity of enzymes involved in the biosynthesis of the compounds in studies on the jasmine *Jasminum auriculatum* and on *Petunia x hybrida*.^{108,161} In *J. auriculatum*, for example, the enzyme responsible for the production of benzyl acetate (**90**) had a higher activity at 25 °C than at 30 °C, whereas the enzyme responsible for linalool (**60** and/or **61**) production had a high activity at both temperatures. These resulted in **90** emissions that peak at 25 °C, whereas high amounts of linalool (**60** and/or **61**) were emitted both at 25 °C and 30 °C.¹⁶¹

In most studied species, drought affected total scent emissions and/or scent composition. Drought increased the total amount of floral scent in *Ipomopsis aggregata*,¹⁶⁸ *Campanula rotundifolia* and *Potentilla recta*,^{142,169} but had no effects on total amount of scent in *Phacelia hastata*,^{142,169} *Fagopyrum esculentum* (buckwheat)¹⁷⁰ and *Matthiola livida*.¹⁷¹ Regarding scent composition, different compounds were responsible in the various study species for differences between drought and control plants.^{142,168,169,171,172} For example, in *Ipomopsis aggregata*, 1,3-octadiene (**117**) and benzyl alcohol (**99**) were released in higher amounts in wetter than drier conditions, whereas α -pinene (**72**), (*E*)- β -ocimene (**62**) and (*E,E*)- α -farnesene (**63**) were released in highest amounts in drier conditions.¹⁶⁸





In *Phacelia hastata*, β -myrcene (76), limonene (75), and 1,8-cineole (70) were emitted at higher rates in control than drought treatments, whereas 6-methyl-5-hepten-2-one (109) and (Z)-3-hexenyl acetate (110) were higher in the drought treatment.¹⁶⁹ In buckwheat, drought increased emissions of (Z)-3-hexenol (118), isobutyraldehyde (119), 2-methylbutanal (120) and 3-methylbutanal (121).¹⁷⁰ Drought was shown to affect (positive or negative) pollinator visitation (e.g., *C. rotundifolia*,¹⁶⁹ *Salvia rosmarinus*,¹⁷² *F. esculentum*¹⁷⁰), but if changes in floral scents contributed to this finding has not been tested so far. In some study species, drought neither affected total scent emission nor scent composition (*Sinapis arvensis*,¹⁷³ *Arabis alpina*,¹⁷⁴ *Heterotheca villosa*^{142,169}). The molecular mechanism of how drought affects the biosynthesis and emission of floral volatiles has not yet been studied in any species, so. It is difficult to explain why certain plants respond to drought differently and why different volatiles are affected.

5.3 Nutrient availability and soil type

Research findings on the effects of nutrient availability and soil type on floral scent emission have become available only in the last few years. In *Lithophragma bolanderi*, nutrient availability (N, P, K) did not have effects on scent emission,¹⁷⁵ whereas in *Petunia hybrida*, only eugenol (87), out of the 10 volatiles analyzed, increased with increasing nitrogen availability.¹⁷⁶ In *Arabis alpina*, plants with high nutrient availability (N, P, K, Mg, micronutrients) produce more scents, with scent composition not being influenced by nutrient availability.¹⁷⁴ A very recent study on *Brassica rapa* did not find an effect of soil type (limestone *versus* tuff) on floral scent emissions.¹⁷⁷

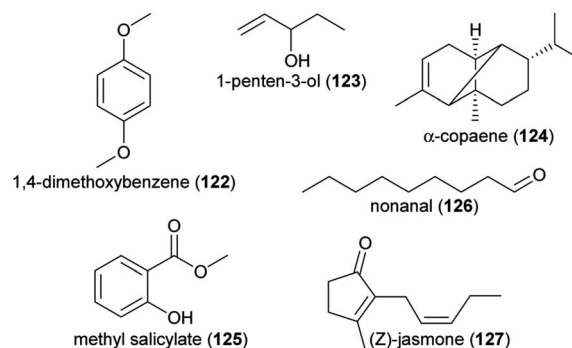
5.4 Bacteria and fungi

In the last decade, there were major advances in our understanding of how microorganisms influence floral scents and the chemical communication between flowers and their visitors. Microorganisms considered were arbuscular mycorrhizal fungi, plant endophytes, such as bacterial pathogens, and especially bacteria and yeasts colonizing flowers.

In *Polemonium viscosum*, resource allocation to scent production *vs.* to mycorrhizae was shown to be a tradeoff, and when fungicide was used to eliminate mycorrhizae as partners, scent production increased (number of compounds emitted and total scent).¹⁷⁸ In strawberry cultivars, mycorrhizal fungi did not have effects on floral scents.¹⁷⁹

The bacterial plant pathogen *Erwinia tracheiphila* that causes a fatal wilt disease in wild gourd (*Cucurbita pepo* ssp. *texana*), lowered total scent emissions, especially due to an attenuated release of linalool (60 and/or 61), 1,4-dimethoxybenzene (122), and a nonatriene.¹⁸⁰ *Erwinia amylovora*, causative agent of the fire blight disease, changed scent emission of apple flowers, which made flowers less attractive to honeybee pollinators.¹⁸¹

Compounds characteristic for apple flowers inoculated with *Erwinia amylovora* were, among others, 1-penten-3-ol (123), α -copaene (124) and methyl salicylate (125), whereas healthy flowers were characterized by benzaldehyde (88), benzyl alcohol (99), 2-phenylacetonitrile (92), nonanal (126) and (Z)-jasmone (127). Among other potential reasons, the observed changes in scent emission following pathogen attack might be due to the *E. amylovora*-induced compounds representing defenses of the plant against the pathogen or the use of some but not other volatiles as a carbon source by the pathogen.¹⁸²



Flowers are colonized by a high number of bacteria and fungi,^{183,184} and several recent review articles summarized how flower visitors disperse microbes among flowers, which effects flower microbes have on plant fitness, and how flower microbes affect floral phenotypes, such as floral scents, and flower-animal interactions.^{182,185–189} There is evidence that microbes dwelling in nectar and also microbes colonizing flower or other plant surfaces change the olfactory display of flowers, sometimes with consequences for the behavior of flower visitors. A high number of volatiles is released from nectar dwelling bacteria and fungi, among them various aliphatic alcohols and esters, such as ethanol (128), 3-methyl-1-butanol (129), acetoin (130), ethyl acetate (131), and 3-methylbutyl acetate (132), but also compounds from other chemical classes, such as the aromatic compound 2-phenylethanol (84) and sulfur-bearing methanethiol (133).^{190–194} Many of these compounds were found to elicit physiological responses in the antennae of pollinators,^{191,193} and scents released from several nectar dwelling microorganisms were found to be neutral, attractive or repellent to bee pollinators^{191–193,195,196} or to other flower visitors, such as parasitoids.^{194,197} Thus, compounds released from nectar-dwelling microorganisms mediate foraging behavior of flower visitors. As shown for *Aphidius ervi*, an aphid parasitoid that uses nectar as food source, feeding on nectar that contains specific nectar-dwelling microorganisms has negative effects on survival.¹⁹⁴



When compared to nectar-dwelling microorganisms, there is much less known about plant-surface inhabiting microorganisms. Helletsgrubber *et al.*¹⁹⁸ inoculated *Brassica rapa* plants with bacterial strains naturally growing on leaves and flowers of this plant species and compared the floral scents of these inoculated plants with the floral scent of plants grown under sterile conditions. They found that the scents strongly differed between the treatments, especially due to higher emission rates of acetoin (**130**) and 2,3-butanediol (**134**) in inoculated plants. Peñuelas *et al.*¹⁹⁹ fumigated *Sambucus nigra* plants with antibiotics and found that fumigation did not affect terpenes in the floral tissue, but negatively affected floral terpene emissions. They concluded that phyllospheric microorganisms contribute to floral terpene emissions and possibly affect the olfactory attraction of thrips pollinators.

5.5 Herbivores

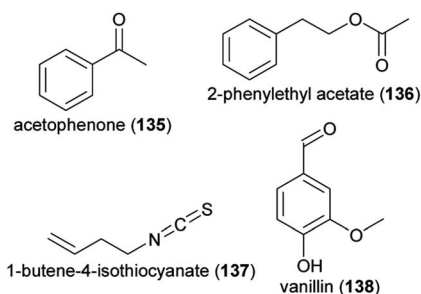
The first studies shedding light on the effects of herbivory/florivory on floral scents and pollinator behavior, were published less than two decades ago, reviewed by^{137,200–204} Meanwhile, there are some studies available that link herbivory on leaves and flowers to floral chemistry and pollinator attraction. They have found that herbivory/florivory often, but not always,^{169,205} has effects on floral scent emissions,^{177,206} and that different herbivores can have different effects on floral scent emissions.^{207–209} Further, herbivory-induced changes in scent emissions often translate to differences in flower visitor behaviors^{169,171,203,207,210–213} and plant reproduction,²¹⁴ but see²¹¹ and herbivores exert selective pressures on plants and affect the evolution of floral scents.^{215,216}

The results of Kessler *et al.*²¹³ on herbivore-induced changes in the volatile emission of *Nicotiana attenuata*, a night-flowering tobacco, are especially remarkable. This species normally releases high amounts of scent, mainly benzylacetone (**91**), at night to attract nocturnal hawkmoth pollinators, such as *Manduca quinquemaculata* and *M. sexta*. Interestingly, these moths not only visit the flowers for nectaring, but females also lay eggs on the leaves, which serve as food for the larvae. When plants are attacked by hawkmoth larvae, they open their flowers in the morning instead of dawn and emit strongly reduced amounts of benzylacetone, likely to avoid further herbivory by hawkmoth larvae. Such plants are then less frequently pollinated by hawkmoths at night, but mainly by hummingbirds during daytime.

In *Datura wrightii*, feeding of *Manduca sexta* larvae on leaves did not affect total flower scent emissions, even though the emission rates of methyl benzoate (**100**) and geraniol (**59**) were increased.²⁰⁶ Other compounds, such as the ones responsible for attraction of *Manduca sexta* to flowers [benzaldehyde (**88**), benzyl alcohol (**99**), and linalool (**60** and/or **61**)] were not affected by herbivory.

In *Brassica rapa*, leaf herbivory by both *Pieris brassicae* and *Spodoptera littoralis* larvae resulted in reduced scent emission and changed scent floral scent compositions.²⁰⁹ Herbivory by both lepidopteran species reduced amounts of 6-methyl-5-hepten-2-one (**109**), phenylacetaldehyde (**83**), acetophenone (**135**), 2-phenylethanol (**84**), 2-phenylethyl acetate (**136**) and decanal (**115**).

Herbivory by *P. brassicae* additionally reduced *p*-anisaldehyde (**111**), whereas this compound increased following herbivory by *S. littoralis*. Benzaldehyde (**88**) was (negatively) affected only by *P. brassicae*. Herbivory by aphids on this plant species also affected floral scent emissions.¹⁷⁷ The amounts of **88**, **111**, methyl benzoate (**100**) and methyl salicylate (**125**) were decreased, whereas 1-butene-4-isothiocyanate (**137**) and (*Z*)-3-hexenyl acetate (**110**) increased. For *B. rapa*, it was also shown that herbivory not only has direct effects on floral scent emissions, but also influences the evolution of floral scents.^{215,216} In the absence of herbivores, such as *Pieris brassicae* larvae, but not when herbivores were present, higher amounts of **88**, 2-phenylacetone nitrile (**92**), **111**, **83**, and **100** evolved within a few generations when plants were under selection of bee pollinators. This suggests that there is a trade-off between the attraction of pollinators and the deterrence of herbivores.



In white mustard, *Sinapis alba*, herbivory by two aphid species (*Lipaphis erysimi*, *Myzus persicae*) but not by the moth *Plutella xylostella* resulted in decreased scent emissions.²⁰⁷ Aphid herbivory resulted in reduced emissions of benzaldehyde (**88**), benzyl alcohol (**99**), *p*-anisaldehyde (**111**), and methyl salicylate (**125**). In contrast, **99** and vanillin (**138**) increased following attack by *Plutella xylostella*. Attack by aphids did not affect pollinator attraction, whereas an aphid predator, the ladybird *Coccinella septempunctata*, preferred flower scent of undamaged plants over that of aphid damaged plants, and the aphid parasitoid *Diaeretiella rapae* preferred flower scent of aphid-damaged plants over that of undamaged plants.²⁰⁷

6 Floral volatiles as attractants for flower visitors

In the last decade, there were major advances in our understanding of the floral volatile constituents involved in the communication between plants and their pollinators. For some pollinator groups (*e.g.*, rodents, Cyclocephalini scarab beetles), the first scent attractants were identified and, for other groups (*e.g.* flies, bees, moths, and butterflies), new attractants were added to already known ones. In moths, it was demonstrated that olfactory receptors on the tip of the proboscis affect foraging behaviors. We focus here on chemicals that were shown to be biologically active in pollinators and florivores, *i.e.* elicit behavioral or at least physiological (*e.g.* stimulate the olfactory circuitry) responses in flower visitors. We especially consider pollinator and flower visitor groups where most advances were made recently, and exclude examples involving orchids, given that orchid volatiles and their importance

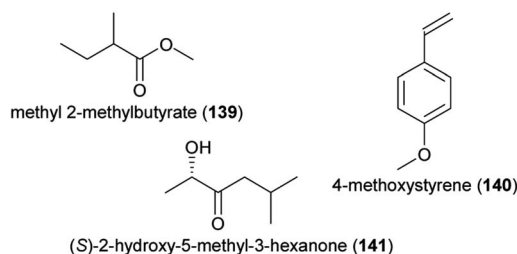


in attracting pollinators were considered in a recent review.¹⁸ First, we will summarize recent studies on beetle attractants, then focus on moths and butterflies, flies, bees and wasps, and finally some groups of flower visitors with only small numbers of species known to interact with flowers.

6.1 Beetles

Beetles are the second largest group of pollinator species (following lepidopterans)²¹⁷ and they are important pollinators of cycads²¹⁸ as well as of various angiosperms, among them species of early diverging lineages.²¹⁹ Some beetles are involved in highly specialized pollination systems (e.g. cyclocephalines, weevils), others are generalist pollinators (e.g., oedemeridae), and some beetles are major florivores (e.g., some cyclocephalines). For all of them, attractive floral volatiles were identified in the last decade.

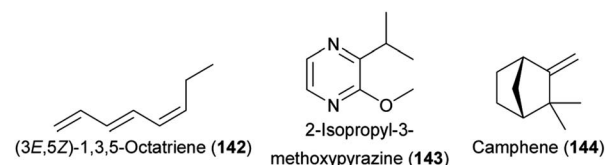
Cyclocephaline scarab beetles are a species-rich (c. 500 species) tribe of rhinoceros beetles and some species in this tribe are highly specific pollinators of various neotropical plant species.^{220,221} Plants pollinated by these beetles have a nocturnal anthesis and thermogenesis, and release extremely strong scents.²²⁰ The chemistry of plants pollinated by cyclocephaline beetles is highly variable, as are the compounds involved in the attraction of the beetles, and several new and uncommon floral scents have been identified while studying the chemical communication in these pollination systems. In 2012, the first attractants of cyclocephaline pollinators were identified. Methyl 2-methylbutyrate (**139**) strongly dominated the flower scent of *Magnolia ovata* (Magnoliaceae) and was attractive to the single pollinator species of this plant, *Cyclocephala literata*.²²² 4-Methyl-5-vinylthiazole (**48**), a new floral compound that strongly dominated (97–100%) the scents of *Annona* species (Annonaceae) and was a major compound in *Caladium bicolor* (Araceae), successfully attracted their pollinators (*Cyclocephala atricapilla*, *C. celata*, *C. vestita*).³⁴ Finally, 4-methoxystyrene (**140**) from *Philodendron form selloum* attracted its single pollinator species, *Erioscelis emarginata*.²²³ The compound (*Z*)-jasmonate (**127**) increased the attractiveness of this methoxylated aromatic compound. In 2013, Maia *et al.*²²⁴ identified (*S*)-2-hydroxy-5-methyl-3-hexanone (**141**), a rare floral scent, and 7,8-dihydro- β -ionone (**96**), a more widespread floral scent, as main compounds in the scent of *Taccarum ulei* (Araceae) and demonstrated that only **141** was attractive to the beetle pollinators (*Cyclocephala cearae*, *Cyclocephala celata*).



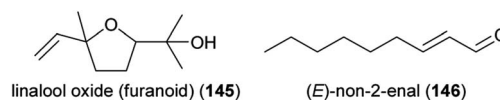
In 2014, Pereira *et al.*²²⁵ identified 7,8-dihydro- β -ionone (**96**) as a strongly dominating compound and methyl jasmonate (**58**) as a minor compound in the scent of *Philodendron adamantinum* (Araceae) and evidenced that both compounds alone,

but especially in a binary mixture, elicit behavioral responses in *Erioscelis emarginata*, the single beetle pollinator. Just recently, Maia *et al.*^{11,226} and Stamm *et al.*²²⁷ identified (*S*)-dehydrojasmonone (**5**) as a novel natural product and (1*S*,2*S*)-isojasmol (**3**) as a rare floral scent both from the scent of *Thaumatococcus melleo-barretoanum* (Araceae), as attractants for *C. gravis* and *C. amblyopsis*, respectively.

Key advances were recently made not only in beetle pollination systems that involve Cyclocephalini, but also in other systems with beetles as the only pollinators or as co-pollinators. The African cycad *Encephalartos villosus*, involved in a highly specialized brood site mutualism mainly with *Erotylidae* sp. nov. and *Portheles* sp. weevils, released the (3*E*)-isomer of 1,3-octadiene (**117**) and (3*E*,5*Z*)-1,3,5-octatriene (**142**) as well as 2-isopropyl-3-methoxypyrazine (**143**) that all elicited physiological responses in the antennae of the beetles.²²⁸ Field experiments proved that the two tested compounds **117** and **143** also elicited behavioral responses. In the related cycad *E. ghellinckii*, *Metacucujus goodie* (Boganidae) beetle pollinators were also attracted by the (3*E*)-isomer of **117**, whereas *Erotylidae* sp. nov. beetles were attracted by camphene (**144**).²²⁹ The weevil *Rhopalotria furfuracea*, a pollinator of the New World cycad *Zamia furfuracea*, was shown to be attracted by cone humidity.²³⁰



Atrichelaphinis tigrina, which is the most common cetonine beetle pollinator of sugarbushes (*Protea*), responded to 15 compounds in electroantennographic analyses, and five thereof were used for behavioral field tests: benzaldehyde (**88**), methyl benzoate (**100**), methyl salicylate (**125**), linalool (**60** and/or **61**), linalool oxide (furanoid) (**145**).²³¹ Only **100** and linalool elicited behavioral responses in the *Atrichelaphinis tigrina* beetles. Antennae of Oedemeridae beetles (*Oedemera* spp.), generalist pollinators of various plants, responded to several volatiles released from strawberry flowers: nonanal (**126**), decanal (**115**), (*E*)-non-2-enal (**146**), linalool (**60** and/or **61**) and 1,4-dimethoxybenzene (**122**).²³²



Rove beetle (Staphylinidae) pollinators were attracted by a blend of indole (**94**), skatole (**147**), *p*-cresol (**148**), 7,8-dihydro- β -ionone (**96**), geranylacetone (**149**) and β -ionone (**82**) to brood-site deceptive *Typhonium eliosurum*.²³³ Antennae of the rove beetle *Pelecomalium testaceum*, main pollinator of the aroid *Lysichiton americanus* (Western skunk cabbage) responded to (*E*)-4-nonene (**150**), (*E*)-5-undecene (**151**) and **94**, whereas the two hydrocarbons together as well as indole successfully attracted the beetle pollinators in field experiments.²³⁴





The compounds α -pinene (72) (*S*:*R* 16:1), sabinene (74), myrcene (76), limonene (75) (*S*:*R* 1:3), terpinolene (152) and (*S*)-bornyl acetate (153), which are released from flowers of the wild carrot *Daucus carota* (Apiaceae), elicited physiological responses in antennae of *Acanthoscelides obtectus* (Coleoptera: Bruchidae), a global pest of dry beans that visits various plants for their nectar and pollen, among them the wild carrot.²³⁵ A synthetic mixture of these compounds was also attractive to this beetle species in lab bioassays, though less than the natural scent of the wild carrot. Another pest of Fabaceae, the bean seed beetle *Bruchus rufimanus*, was shown to respond to floral volatiles of its principal host plant, the field bean *Vicia faba*, in physiological and behavioral experiments,²³⁶ though the beetle did not necessarily pollinate the flowers. Nine floral scents were active in electroantennographic analyses, whereas a blend of three thereof, (*R*)-linalool (60), (*E*)-cinnamyl alcohol (154) and (*E*)-cinnamaldehyde (155), was as attractive to the beetles in field biotests as a blend of the nine compounds. Similarly, the ladybird beetle *Harmonia axyridis* visits *Sophora japonica* flowers to feed on pollen and nectar, with unknown effects on pollination.²³⁷ Physiological and behavioral experiments identified nonanal (126) as the floral scent responsible for the attraction of *H. axyridis*.²³⁷



Some plants are not only attractive to beetle pollinators but also to beetle florivores, and in palms (Arecaceae) it was shown that both pollinators and florivores are attracted by the same compounds. The macauba oil palm (*Acrocomia aculeata*) attracts weevil pollinators of the genus *Andranthobius* (Curculionidae) by *p*-methylanisole (156), 2-isopropyl-3-methoxypyrazine (143) and 2-*sec*-butyl-3-methoxypyrazine (157). Thereof, 156 was also attractive to florivorous *Cyclocephala forsteri*, and the two pyrazines to another florivorous cyclocephaline beetle, *Aspidolea bleuzeni*.²³⁸ The acuri palm (*Attalea phalerata*) releases almost exclusively methyl acetate (158) to attract its putative main beetle pollinators (*Mystrops* spp., Nitidulidae; *Andranthobius* spp., Curculionidae) and this communication channel is exploited by non-pollinating florivorous beetles, such as palm

borers (*Paratenthra martinsi*, Cerambycidae; *Parisoschoenus* sp., *Belopoeus* sp.; Curculionidae).²³⁹



Finally, some recent studies focused only on florivorous beetles and identified the floral attractants. The (3*S*)-isomer of (*E*)-nerolidol (113) mediated the attraction of florivorous *Cyclocephala paraguayensis* beetles to bottle gourd (*Lagenaria siceraria*, Cucurbitaceae) flowers,²⁴⁰ and 2-isopropyl-3-methoxypyrazine (143) and 2-isobutyl-3-methoxypyrazine (159), major compounds of *Acrocomia* and *Attalea* palms, attracted *Cyclocephala amazona* and *C. distincta* florivores.²⁴¹ *Aulacophora foveicollis* (Coleoptera: Chrysomelidae), a beetle pest that feeds on flowers and leaves of two Cucurbitaceae species (Gac, *Momordica cochinchinensis*; creeping cucumber, *Solena amplexicaulis*), was attracted by a blend of linalool oxide (furanoid) (145), 1-octanol (160), and nonanal (126).²⁴² These compounds occurred in flowers of both species and the blend was as attractive as the natural scent. Coccinelidae beetle florivores (*Epilachna dodecastigma*) of the bitter melon (*Momordica charantia*) were attracted to the flowers by sabinene (74), which was as attractive as the total flower scent.²⁴³ Studies with transgenic lines of *Petunia x hybrida* silenced in their ability to produce specific floral compounds revealed that methyl benzoate (100) is attractive to florivorous *Diabrotica undecimpunctata*.²⁴⁴



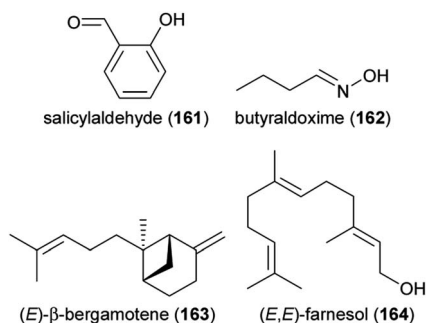
6.2 Moths, butterflies

Lepidoptera are the most species-rich group of pollinators,²¹⁷ and recent advances were made in understanding the molecular basis of olfaction in these insects, *i.e.* the detection of floral volatiles by olfactory receptors,^{245,246} and the importance of specific molecules in the attraction of Lepidoptera to flowers. Here, we will summarize the main findings of studies that identified floral volatiles attractive to hawkmoth, settling moth and butterfly pollinators.

6.2.1 Hawkmoths. Most work in hawkmoths (Sphingidae) was done on *Manduca sexta*, the tobacco hawkmoth. This moth species visits *Nicotiana* species and various other plants for nectaring, and Riffell *et al.*²⁴⁷ demonstrated that plants pollinated by this species converged in their scents. They are often dominated by the three oxygenated benzenoids methyl benzoate (100) (see also ref. 103), benzyl alcohol (99) and benzaldehyde (88). Though plants that have another composition are also visited by naive moths, they are less attractive.



Despite this behavior, the moth's olfactory system also responds to other compounds, such as salicylaldehyde (**161**), phenylacetaldehyde (**83**), methyl salicylate (**125**), linalool (**60** and/or **61**), limonene (**75**), caryophyllene (**64**), butyraldioxime (**162**), nonanal (**126**), and 3-methylbutylaldoxime (**114**). Through learning however, the attractiveness of originally less attractive blends increases, making it possible for the moths to effectively forage in habitats that have different plants available.²⁴⁷ The findings of Haverkamp *et al.*²⁴⁸ and Zhou *et al.*²⁴⁹ were highly innovative in this regard, showing that *Manduca sexta* has receptors on the tip of the proboscis that are sensitive to floral volatiles and influence feeding behaviors. Benzylacetone (**91**), a known attractant for moths looking for *Nicotiana attenuata* flowers (see also before), and (*E*)- α -bergamotene (**65**), a floral compound of *N. attenuata* not attractive to *M. sexta*, were both shown to be sensed by these receptors. This resulted in increased nectaring times, with positive consequences on the pollination efficiency of the moth and on nectar uptake.^{248,249} In contrast to volatiles sensed by the antennae, however, moths do not learn volatiles sensed by the proboscis and information learnt by the antennae cannot be retrieved by the proboscis.²⁵⁰



Some research was also done on *Hyles* and *Agrius* hawk-moths. Chemical ecological studies on two closely related *Ipomopsis* (Polemoniaceae) species, *I. aggregata* and *I. tenuituba*, showed that floral scents contribute to the reproductive isolation of the two species. *Hyles lineata* pollinators were only attracted to *I. tenuituba*, and key for the selective attraction was indole (**94**), which was only released from *I. tenuituba* flowers.²⁵¹ Quite different compounds seem to be involved in the attraction of *H. livornica* to flower scents of *Dianthus inoxianus*.²⁵² Several aliphatic ketones (e.g., 2-undecanone, 2-tridecanone, 2-pentadecanone) and terpenoids [e.g., (*E*)- β -bergamotene (**163**), (*E,E*)-farnesol (**164**)] elicited physiological responses in antennae of the moths. In elegant studies, von Arx *et al.*²⁵³ and Dahake *et al.*²⁵⁴ demonstrated that approaching and landing behaviors of foraging *H. lineata* and *M. sexta* hawkmoths are influenced by floral humidity. Floral humidity gradients are due to passive (nectar evaporation) and/or active (stomatal conductance) processes and allow the moths to sense nectar availability from a distance and to forage efficiently.^{253,254} Finally, *Agrius convolvuli* responded to various compounds of its *Gladiolus longicollis* host plant, among them aromatic compounds [e.g. benzaldehyde (**88**), benzyl acetate (**90**) phenylacetaldehyde (**83**), eugenol (**87**)] and nitrogen-bearing compounds [3-methylbutylaldoxime (**114**), 2-phenylacetoneitrile (**92**), 2-phenylnitroethane (**93**)].²⁵⁵

6.2.2 Settling moths. Several compounds were, in the last decade, identified as being biologically active in settling moths, among them compounds well known as moth attractants before [e.g., benzaldehyde (**88**), lilac aldehyde (**165**), phenylacetaldehyde (**83**)].^{256–259} Here, we focus on the chemical communication in brood site mutualisms, involving *Greya*, *Epicephala*, and yucca moth nursery pollinators. *Greya* moths (Prodoxidae) are highly specialized on the woodland stars, *Lithophragma*, and in olfactometer experiments, they were attracted by the plants' scents alone and preferred the scents of their local *Lithophragma* host over that of non-local hosts.²⁶⁰ Schiestl *et al.*²⁶¹ demonstrated that the antennae of the moths are responsive to a high number of compounds; among them widespread floral scents, such as 1,4-dimethoxybenzene (**122**), methyl salicylate (**125**), 2-phenylethyl acetate (**136**), 2-amino-benzaldehyde (**166**), but also the very rare (6*R*,10*R*)-6,10,14-trimethylpentadecan-2-one (**167**) (hexahydrofarnesyl acetone).



Epicephala moths are obligate nursery pollinators of several Phyllanthaceae and the first attractants were identified in 2010. Flowers of both sexes of monoecious *Breynia vitis-idaea* released similar scent compounds, though in sex-specific ratios, and several main compounds in at least one of the sexes [geranylacetone (**149**), 2-phenylethanol (**84**), 2-phenylacetoneitrile (**92**)] but also minor compounds [2-phenylethyl acetate (**136**), indole (**94**)] elicited signals in antennae of the *Epicephala* moths.²⁶² **84** and **92** were used in bioassays, which revealed that **84** but not **92** was attractive to the moths when tested alone. Still, only the mixture of both compounds was as attractive as the natural scent.²⁶² Further studies in Phyllanthaceae species pollinated by *Epicephala* moths revealed that the scents are highly dimorphic between female and male flowers^{263–266} and that moths are able to discriminate between them.^{264,265} This allows the moths to carry out different behaviors on different sex flowers: pollen collection on male flowers, active pollination and oviposition on female flowers. Regarding the interaction between *Yucca* and yucca moths, it has been known for a while that plants release strong floral scents that are attractive to the moth pollinators.²⁶⁷ Just recently, however, the first *Yucca* compounds were identified that are biologically active in yucca moths. Among them are several compounds new for science.²⁹ Antennae of *Tegeticula mexicana* pollinating moths and the parasitic, non-pollinating bogus yucca moth, *Prodoxus tam-aulipellus*, responded in electrophysiological experiments to (*Z*)-filamentolide (**24**) of *Yucca treculeana*, whereas antennae of pollinating *T. yucasella* and *T. cassandra*, and parasitic *P. decipiens* responded to **24**, (*Z*)-filamentol (**22**), (*Z*)-filamental (**23**),



(*E*)-4,8-dimethyl-1,3,7-nonatriene (**80**), and (*Z*)-9-nonadecene (**168**) of *Y. filamentosa*. **24** was used for field bioassays and was capable of attracting parasitic *Prodoxus* species, but the attraction of pollinating yucca moths to their host plants still needs to be resolved in the future.²⁹



6.2.3 Butterflies. Butterflies are believed to primarily use visual cues for finding appropriate host plants, but some species also use olfaction when looking for flowers.²⁶⁸ To the best of our knowledge, only a single study decoded the chemical communication in a butterfly pollination system in the last decade. In a very recent study Lehner *et al.*²⁶⁹ asked why the butterfly bush *Buddleja davidii* is extremely attractive to butterflies. It was shown that the peacock butterfly *Inachis io* is attracted by visual and olfactory cues, but only the olfactory cues elicit feeding responses. Bioassays with compounds that were previously shown to elicit antennal responses in butterflies²⁷⁰ demonstrated that 4-oxoisophorone (**169**) and oxoisophorone epoxide (**170**) were highly effective in eliciting feeding responses in peacock butterflies. Blends of the other terpenes [β -cyclocitral (**171**), farnesene (**172**), geranylacetone (**149**), (*E*)- β -ocimene (**62**)] and of aromatic compounds [benzaldehyde (**88**), benzyl alcohol (**99**), (*E*)-cinnamyl alcohol (**154**), (*E*)-cinnamaldehyde (**155**), phenylacetaldehyde (**83**), 2-phenylethanol (**84**)] were less effective. The butterfly bush releases **169** and **170** in higher amounts than other plants,²⁷¹ explaining its strong attractiveness to the peacock butterfly and potentially other Lepidoptera pollinators.^{269,272}



6.3 Flies

Many plants rely on flies as pollinators,^{217,273,274} and, in the last decade, various flower volatile attractants were identified in mutualistic as well as in deceptive pollination systems, and new olfactory based strategies of deceptive plants that exploit flies as pollinators were introduced.

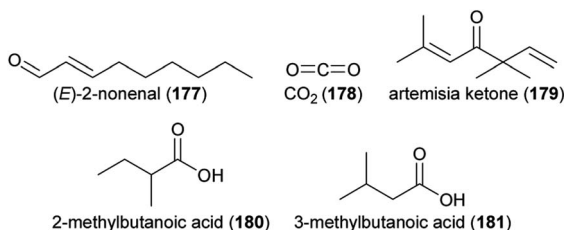
6.3.1 Mutualistic pollination systems. Hoverflies are frequent flower visitors of various plant species, such as the creeping thistle *Cirsium arvense*. *Episyrphus balteatus*, the marmalade hoverfly, uses olfactory cues to find flower heads of this species, and candidate compounds responsible for

attraction of the flies are phenylacetaldehyde (**83**), methyl salicylate (**125**), dimethyl salicylate (**173**), linalool oxide pyranoid (**174**), linalool (**60** and/or **61**) and 2-phenylethanol (**84**). These compounds elicited responses in electroantennographic analyses.²⁷⁵ The aromatic compounds thereof were also found to elicit antennal responses in another hoverfly species, *Eupeodes corollae*.¹⁷ Antennae of *E. corollae* responded to various further aromatic floral compounds, among them, *p*-cresol (**148**), and methyleugenol (**175**), to which an olfactory receptor is narrowly tuned.¹⁷ Hoverflies also visit stone fruit crops, such as apricot, cherry and peach, and floral scents of these crops attractive to hoverflies were 4-oxoisophorone (**169**) and 3,5-dimethoxytoluene (**176**).²⁷² Occasionally, hoverflies visit flowers of *Arabidopsis thaliana*, which releases various terpenes, among them linalool and derivatives thereof. Of these compounds, linalool (**60** and/or **61**) was shown to be attractive to hoverflies, whereas lilac aldehyde (**165**) was repellent.⁵⁹



Mosquitoes, among them species that are vectors of diseases (*e.g.* malaria, dengue), also frequently visit flowers for nectaring. Several recent reviews focused on (floral) volatiles attractive to mosquitoes and how such volatiles could be implemented in strategies for management of disease vectors.^{276–282} Here a few examples for studies that successfully identified compounds that elicit behavioral responses in mosquitoes are given. Otienoburu *et al.*²⁸³ analysed floral scents in the common milkweed *Asclepias syriaca*, which is visited by the northern house mosquito, *Culex pipiens*, and demonstrated that a blend of benzaldehyde (**88**), phenylacetaldehyde (**83**), and (*E*)-2-nonenal (**177**) is key in the attraction of the mosquitoes to the flowers. Similarly, acetophenone (**135**) from sweet alyssum (*Lobularia maritima*) flowers successfully attracted the yellow fever mosquito, *Aedes aegypti*.²⁸⁴ In a remarkable study, Peach *et al.*²⁸⁵ demonstrated that the northern house mosquito and the yellow fever mosquito are not only attracted by the floral scents (and visual plant cues) of the common tansy (*Tanacetum vulgare*), but also by plant-derived CO₂ (**178**). After sunset, plants became net producers of **178**, which increased the attractiveness of a synthetic floral blend that consisted of 20 compounds, among them benzaldehyde (**88**), acetophenone (**135**), artemisia ketone (**179**), and 2- and 3-methylbutanoic acid (**180** and **181**).





Various miterwort species (*Mitella* spp.) are mainly pollinated by nectaring fungus gnats, either by *Gnoriste mikado*, a long-tongued species, or by short-tongued species. The flowers of these plant species release mainly terpenoids, with the scent of species pollinated by *G. mikado* being dominated by lilac aldehyde (165).²⁸⁶ This compound was found to repel short-tongued gnats, but to frequently elicit nectaring behavior in *G. mikado*.

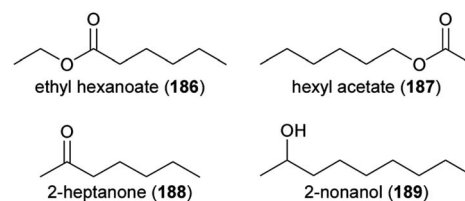
The globeflower *Trollius europaeus* is involved in a highly specific brood site mutualism with *Chiastocheta* (Anthomyiidae) flies. Its flowers release mainly terpenoids, but also a few fatty acid derivatives and miscellaneous cyclic and aromatic compounds, of which the terpenes (*E*)- β -caryophyllene (64), germacrene D (101), (*E,E*)- α -farnesene (63) and linalool (60 and/or 61), the aromatic compound methyl salicylate (125), and the fatty acid derivative (*Z*)-jasmone (127) elicited antennal responses in *Chiastocheta* fly pollinators.²⁸⁷ In another plant involved in a brood site mutualism with flies, *i.e.* *Rheum nobile*, the Sikkim rhubarb, the floral scents are dominated by methyl 2-methylbutyrate (139) and α -pinene (72). *Bradysia* fungus gnat pollinators were attracted by a blend of these two compounds and by 139 alone, but not by 72 alone.²⁸⁸



6.3.2 Deceptive pollination systems. Many deceptive plants chemically mimic brood sites of their pollinators, such as dung or carrion, and various of these brood-site deceptive plants exploit flies as pollinators.²⁸⁹ In various such sapromyophilous pollination systems, the oligosulfides dimethyl disulfide (182) and dimethyl trisulfide (183) were confirmed to be key mediators. As a blend, these compounds attracted pollinators (Anthomyiidae, Calliphoridae, Muscidae, and Sarcophagidae) of several brood-site deceptive plants (*e.g.* *Jaborosa rotacea*, *J. laciniata*, *Eucomis* spp., *Amorphophallus konjac*, *Rafflesia cantleyi*) that release these compounds.^{290–294} By adding blends of these compounds to non-saprophilous plant species pollinated by wasps and hawkmoths, Shuttleworth and Johnson²⁹³ and Moré *et al.*²⁹² could demonstrate in *Eucomis* (African pineapple lilies) and *Jaborosa*, respectively, that these compounds are sufficient to attract the fly pollinators of saprophilous congeners. In *Rafflesia cantleyi*, which is only pollinated by the calliphorid fly *Chrysomya chani*, the individual compounds as well as a blend of the two oligosulfides were attractive to the flies.²⁹⁴ In other calliphorid flies, however, only 183 elicited physiological and behavioral responses.²⁹⁵ Similarly, only 183 elicited

antennal responses in Muscidae (*Musca domestica*, *Atherigona* sp.) fly pollinators.^{295,296} Besides oligosulfides, several brood-site deceptive plants release nitrogen-bearing compounds, such as indole (94) and skatole (147),²⁸⁹ and a few studies confirmed their importance in attracting pollinators. In *Periploca laevigata* (Apocynaceae), for example, 183 and 94, and additionally the sesquiterpenes (*E*)- β -caryophyllene (64) and germacrene D (101) elicited antennal responses in the housefly (*Musca domestica*) pollinators, whereas 94 was as attractive as the blend of these four compounds in bioassays.²⁹⁷ In *Wurmbea elatior* (Colchicaceae), a species pollinated by various different saprophilous flies, 94 and 147 elicited antennal responses, but only 147 was needed to attract the same assemblage of flies as the flowers do.²⁹⁸ Finally, 94 and 147 were, together with 148, 7,8-dihydro- β -ionone (96), geranylacetone (149), and β -ionone (82), part of a mixture that attracted moth fly pollinators (Psychodidae) to the aroid *Typhonium eliosurum*.²³³ Nitrogen-bearing 1-pyrroline (184) was responsible for the human semen-like odor of *Stemona japonica* (Stemonaceae), which is mainly pollinated by *Atherigona* flies (Muscidae).²⁹⁹ The flowers additionally released 2-methyl-1-butanol (185), 3-methyl-1-butanol (129), 2-methylbutanal (120) and 3-methylbutanal (121). A synthetic mixture of all five compounds, but not 184 alone, attracted the *Atherigona* flies. Many flowers release not only fetid but additionally often various “sweet scents”, and Zito *et al.*³⁰⁰ demonstrated for *Caralluma europaea* (Apocynaceae) that they also are involved in the communication between brood-site deceptive plants and their pollinators. Main compounds in the floral scent of this plant species were terpinolene (152), α -terpinene (116), and linalool (60 and/or 61), and a mixture thereof attracted housefly pollinators in y-tube olfactometer assays.³⁰⁰

A very different set of chemicals is involved in plants that chemically mimic fermenting organic plant material such as fruits and vegetables, and target drosophilid flies as pollinators.³⁰¹ In an extraordinary work, Stökl *et al.*³⁷ decoded the mimicry strategy of the Solomon's lily, *Arum palaestinum* (Araceae). This brood-site deceptive trap plant attracted various drosophilid species, mostly *Drosophila simulans*, by using fermentation-associated volatiles. The floral scent was, to the human nose, reminiscent of a fruity wine and contained 14 compounds that elicited antennal responses in drosophilids. The most consistent responses were elicited by 2,3-butanediol acetate (55), acetoin acetate (56), ethyl hexanoate (186), hexyl acetate (187), 2-phenylethanol (84) and 2-phenylethyl acetate (136), compounds that also occurred in fermenting fruits.³⁷ Studies in the brain of *Drosophila* revealed the olfactory receptors involved.³⁷ As a synthetic mixture, these compounds were as attractive to the flies as the odor released from a fermenting banana in behavioral assays.

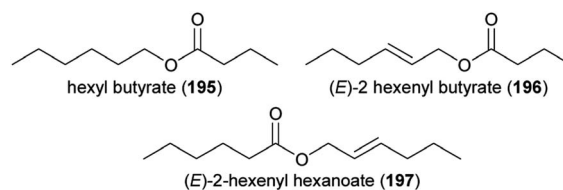


Studies on the chemical ecology of pollination in *Aristolochia* (Aristolochiaceae) and *Ceropegia* (Apocynaceae) species also revealed a new intriguing deceptive strategy in these plants, called kleptomyiophily. Species in these genera build trap flowers that mimic volatiles released from preyed-upon insects to attract and exploit kleptoparasitic flies as pollinators. Kleptoparasitic *Desmometopa* flies (Milichiidae) are main pollinators of the giant *Ceropegia*, *C. sandersonii*. These flies are well known to feed on honeybees that are eaten by spiders. Heiduk *et al.*²² found that the flies use compounds of the alarm pheromone of honeybees, that they release following an attack, to find appropriate feeding sites. These same compounds are released from *Ceropegia sandersonii* flowers to attract these flies as pollinators. Compounds that were released from both honeybees under simulated attack and *C. sandersonii* flowers, that elicited antennal responses in *Desmometopa* and that attracted the flies as a blend were geraniol (59), 2-heptanone (188), 2-nonanol (189) and (*E*)-2-octen-1-yl acetate (2).²²

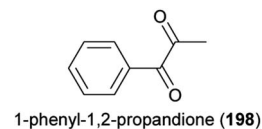
A congeneric species, *C. dolichophylla*, deceives kleptoparasitic *Desmometopa* flies by using different chemicals. Its flower scent is dominated by spiroacetals, mainly (2*S*,6*R*,8*S*)-8-methyl-2-propyl-1,7-dioxaspiro[5.5]undecane (190), (*E,E*)-2,8-dimethyl-1,7-dioxaspiro[5.5]undecane (191) and (*E,E*)-2,8-diethyl-1,7-dioxaspiro[5.5]undecane (192), as well as by *n*-tridecane (193) and *N*-(3-methylbutyl)acetamide (194). Thereof, the spiroacetals and 194 elicited antennal responses and the compounds most attractive to the flies in bioassays were 190 and mixtures that contained this compound, with 194 being less attractive.³⁰² Isomers of 190 and 194 are known from venom glands of paper wasps, suggesting that *C. dolichophylla* mimics pheromones of paper wasps, a potential food source of its pollinators.³⁰²



Aristolochia rotunda also has a kleptomyiophilous pollination strategy, but mimics volatiles of true bugs (Miridae) to deceive kleptoparasitic chloropid flies. Specifically, Oelschlägel *et al.*³⁰³ demonstrated that *A. rotunda* mimics pheromones/defensive compounds of the bugs. Compounds that overlapped between the flowers and the bugs and elicited physiological and behavioral responses (as a blend) in the chloropid pollinators (mainly *Trachysiphonella ruficeps*) were aliphatic esters, among them hexyl butyrate (195), (*E*)-2-hexenyl butyrate (196), and (*E*)-2-hexenyl hexanoate (197).³⁰³



Bioactive compounds, among them novel natural products, were also described from flowers of another deceptive *Ceropegia* species, *C. stenantha*. This species is pollinated by scatopsid flies and releases various aromatic compounds.³¹ Several thereof elicited antennal responses in fly pollinators, among them widespread floral scents [*e.g.* benzyl alcohol (99), 2-phenylethanol (84), 1-phenyl-1,2-propandione (198)], but also compounds so far only known from *C. stenantha* (37 and 38).



6.4 Bees

Bees are the most important pollinators when it comes to the number of plants pollinated,²¹⁷ and it has been known for a long time that they use floral scent for locating host plants, often together with visual floral cues.^{2,6,304} While some bees are generalist flower visitors that visit (mainly during the day, but sometimes at night) various species to collect pollen (and nectar) provisions for their offspring, others collect pollen only from plants of a single family or genus (oligoleges).³⁰⁵ Bees that collect fatty oils from oil-secreting flowers (oil-collecting bees) or those that collect floral volatiles (male euglossine bees) are highly specialized.³⁰⁵ For all these types of bees, major advances were recently made in our understanding of the specific chemicals involved in the communication between plants and their bee pollinators.

6.4.1 Generalist bees. Byers *et al.*⁵⁵ demonstrated that *Bombus vosnesenskii* bumblebees, main pollinators of the monkeyflower *Mimulus lewisii*, prefer the scent of this plant species over that of hummingbird-pollinated *M. cardinalis*. A blend of limonene (75), (*E*)- β -ocimene (62), and β -myrcene (76), which was as attractive to the bumblebees as the natural scent of *M. lewisii*, was responsible for this preference. These monoterpenes did not occur in the scent of *M. cardinalis* [(*E*)- β -ocimene (62), β -myrcene (76)] or were emitted at a much higher rate in *M. lewisii* (75). Another bumblebee species, *Bombus diversus diversus*, was attracted by a blend of the EAD-active compounds linalool (60 and/or 61), α -terpineol (71) and geraniol (59),³⁰⁶ released from the epiparasitic plant *Monotropastrum humile* (Ericaceae). Carpenter bees (*Xylocopa* sp.) were successfully attracted by β -ionone (82), a compound frequently released from plants pollinated by these bees.³⁰⁷

In *Brassica rapa*, Knauer and Schiestl³⁰⁸ identified an honest volatile signal for the availability of rewards. The flowers of this species released four compounds that elicited consistent antennal responses in *Bombus terrestris* pollinators. The aromatics phenylacetaldehyde (83), acetophenone (135) and *p*-

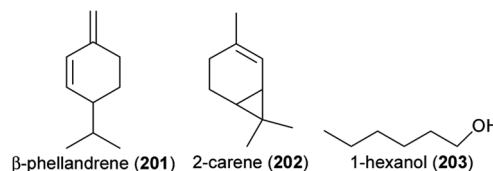


anisaldehyde (**111**), and the terpene (*E,E*)- α -farnesene (**63**). The amount of **83** positively correlated with pollen and nectar sugar availability, and while foraging on the flowers, bumblebees developed a strong preference for this compound over other EAD-active compounds. In *Salix caprea*, the temporal emission pattern of floral scent was found to correlate with the olfactory preferences of diurnal bee and nocturnal moth pollinators. Honeybee pollinators were strongly attracted to 1,4-dimethoxybenzene (**122**), a compound emitted in much higher amounts during the day than at night. They preferred this compound over lilac aldehyde (**165**) (strong attractant for moths, see above), which attracted only a few individuals in a choice setting and is released in higher amounts during night than day.²⁵⁷ Compounds bioactive to bees were, in the last decade, also described from some crop plants. In three highbush blueberry (*Vaccinium corymbosum*) cultivars, floral scents are dominated by (*E*)-cinnamyl alcohol (**154**), but various aliphatic compounds and terpenes also are released.³⁰⁹ A synthetic blend consisting of 16 compounds, among them **154**, α -pinene (**72**), β -pinene (**73**), linalool (**60** and/or **61**), and (*Z*)-3-hexenyl acetate (**110**), was attractive to honeybee pollinators, whereas single compounds were inactive. El-Sayed *et al.*²⁷² identified pollinator attractants of stone fruits. 4-Oxoisophorone (**169**) from apricot (*Prunus armeniaca*), cherry (*P. avium*), and plum (*P. domestica*), and 3,5-dimethoxytoluene (**176**) from peach (*P. persica*) and cherry successfully attracted bee pollinators in field trapping experiments.²⁷² Floral scents that elicit electroantennographic responses in honeybee pollinators were recently identified in the vegetable crops, apple and European pear. Compounds eliciting antennal responses were 2-aminobenzaldehyde (**166**), benzaldehyde (**88**), decanal (**115**) and nonanal (**126**) in Chinese cabbage (*Brassica rapa* breeds), nonanal (**126**), methyl salicylate (**125**), and phenylacetaldehyde (**83**) in carrot (*Daucus carota sativus*), and nonanal (**126**), **169**, **83** and 2-phenylethanol (**84**) in radish (*Raphanus raphanistrum sativus*).³¹⁰ In apple (*Malus domestica*), antennal responses were elicited by **88**, benzyl alcohol (**99**) (dominated the scent bouquet), (*E*)-cinnamyl alcohol (**154**), eugenol (**87**), methyleugenol (**175**), and two unknown compounds (Rachtersberger *et al.* 2019).³¹¹ About 20 compounds in the flower scent of pear (*Pyrus communis*) elicited antennal responses in honeybees, among them common [e.g., (*E*)- β -ocimene (**62**), 4-oxoisophorone (**169**)], but also rare [*N*-isobutylformamide (**199**), (*E*)-*N*-(3-methylbutyl)-1-phenylmethanimine (**200**)] floral scents and even compounds newly identified from nature [(*E*)-*N*-(2-methylbutyl)- and (*E*)-*N*-(3-methylbutyl)-1-(pyridin-3-yl)methanimine (**46** and **47**)].³¹²



Finally, one study tested for correlations between floral volatiles of greenhouse tomato (*Lycopersicon esculentum*, Solanaceae) and pollination by bumblebees.³¹³ Tomato flowers released β -phellandrene (**201**), 2-carene (**202**), α -pinene (**72**) and

p-cymene (**106**), and bumblebees preferably pollinated flowers that produced less **201** and **202** than flowers producing more of these compounds. The repellent properties of these compounds may protect flowers from damage (bees grasp the flowers with their mandibles for extracting pollen by buzzing) caused by over-pollination.³¹³

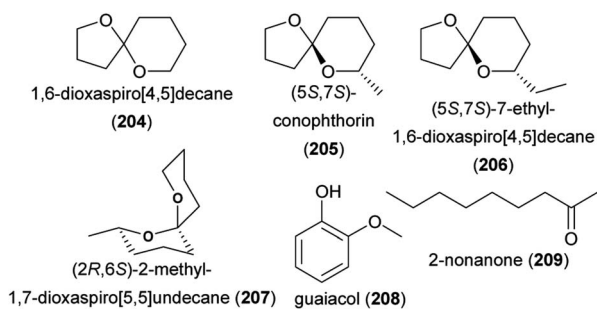


6.4.2 Nocturnal bees. While most bees are diurnal, there are about 250 species from four bee families (Andrenidae, Apidae, Colletidae, and Halictidae) that only fly during the dusk, dawn, or at night.³¹⁵ The first studies that proved the importance of floral scents in host finding of these bees identified the attractive cues of two of the plant species pollinated by these bees. Flowers of the Myrtaceae crop *Campomanesia phaea* (cambuci) open 1–1.5 h before sunrise and release a strong scent. Nocturnal bees (e.g. *Ptiloglossa latealcarata*, *Megalopta sodalis*, *Megommation insigne*) are the principal flower visitors before sunrise and the main pollinators of the plant.³¹⁴ Only later, other insects, such as (introduced) honeybees and other diurnal bee species, frequently visit the flowers but with no effect on pollination.³¹⁴ The flowers release compounds of various chemical classes, with the aliphatic compounds 1-hexanol (**203**) and 1-octanol (**160**) as well as the aromatic compounds 2-phenylethanol (**84**) and benzyl alcohol (**99**) dominating the scent. A blend of these main compounds and **160** alone, but not the other compounds alone, attracted nocturnal bee pollinators in a field bioassay.³¹⁴ Also the attractiveness of 2-phenylethanol alone to species pollinating cambuci was later confirmed.³¹⁵ *Paullinia cupana* (Sapindaceae), commonly known as guarana, is another plant species attractive to nocturnal bee pollinators (*Megalopta* spp., *Ptiloglossa* aff. *lucernarum*).^{316,317} A synthetic mixture composed of the main compounds released at night [(*E*)- β -ocimene (**62**), linalool (**60** and/or **61**), linalool oxide (furanoid) (**145**), (*E*)- β -caryophyllene (**64**)] effectively attracted *Megalopta* pollinators.³¹⁶ Further compounds attractive to nocturnal bees were found by chance in experiments aimed to attract male euglossine bees [e.g., methyl salicylate (**125**), 1,8-cineole (**70**), vanillin (**138**); reviewed in Martínez-Martínez *et al.*³¹⁵].

6.4.3 Pollen-specialist (oligolectic) bees. *Chelostoma rapunculi*, a bee specialized on *Campanula* species, responded in electroantennographic analyses to aromatic compounds, terpenoids, spiroacetals and other aliphatic compounds, all released from flowers of *Campanula trachelium*, one of its host plants.³¹⁸ Only blends of all compounds and of spiroacetals [1,6-dioxaspiro[4.5]decane (**204**), (5*S*,7*S*)-conophthorin (**205**), (5*S*,7*S*)-7-ethyl-1,6-dioxaspiro[4.5]decane (**206**), (2*R*,6*S*)-2-methyl-1,7-dioxaspiro[5.5]undecane (**207**)] were capable of attracting flower inexperienced bees. Flower-experienced bees additionally responded to the aromatic compounds [blend of phenylacetaldehyde (**83**), guaiacol (**208**), methyl salicylate (**125**),



2-phenylethyl acetate (**136**), 2-phenylethanol (**84**), the aliphatic 2-nonanone (**209**), and a blend of the terpenoids [(*E*)- β -ocimene (**62**), (*E*)-isomer of linalool oxide (furanoid) (**145**), terpinolene (**152**), linalool (**60** and/or **61**), α -copaene (**124**), geranylacetone (**149**)].



Recently, it was shown that many of these compounds also occur in *Malva moschata* and *Geranium sanguineum*, plants also occasionally visited by *Chelostoma rapunculi*.³¹⁹ *Campanula* species are also frequently visited by the second generation of *Andrena bicolor*, a bivoltine bee. While the first generation of this species visits various different plant species in spring, among them *Taraxacum officinale*, the second generation in summer prefers *Campanula* spp.³²⁰ Despite this difference, bees of the two generations did not differ in their floral preference when being offered *T. officinale* and *C. trachelium* in a choice setting, and also responded to the same compounds in electroantennographic measurements [*T. officinale*: benzaldehyde (**88**), acetophenone (**135**), benzoic acid (**210**), β -copaene (**68**); *C. trachelium*: 2-phenylethyl acetate (**136**), terpinolene (**152**), β -elemene (**211**), (*E*)- β -caryophyllene (**64**); both species: linalool (**60** and/or **61**), (*E*)- β -ocimene (**62**)], with *Campanula*-spiroacetals eliciting no responses in either generation of this bee species.³²⁰ In a comparative study on the olfactory systems of *Chelostoma rapunculi* and other bee species specialized on *Campanula* (*Chelostoma campanularum*, *Hoplitis mitis*) as well as of generalist bees (*Andrena bicolor*, *Bombus terrestris*, *Apis mellifera*), Brandt *et al.*³²¹ found that *Campanula* specialists are more sensitive to spiroacetals suggesting that they are well adapted in locating their specific host plants effectively. *Campanula* species release spiroacetals only in small amounts and the sensitivity of these bees might border on a pheromone-like attraction.³¹⁸



A similar conclusion was drawn on *Andrena vaga*, a bee specialized on willows (*Salix*). Measurements of the neural activity in the brain (antennal lobe) demonstrated that *A. vaga* is more sensitive to 1,4-dimethoxybenzene (**122**), a behavior-mediating odorant of *Salix* host flowers,³²² than honey bees.³²³ Schäffler *et al.*¹⁹ identified with diacetin [1,2- + 1,3-diacetin (**53** and **54**)] the first olfactory mediator of the interaction between

oil-secreting plants and oil-collecting bees. This compound was key in attracting *Macropis fulvipes* to *Lysimachia punctata* host flowers, with other compounds [heptanoic acid (**212**), (*E*)-2-dodecenal (**213**)] increasing the attractiveness of diacetin. Further, diacetin occurs not only in *L. punctata* but also in other *Lysimachia* species pollinated by *M. fulvipes* or other *Macropis* species,³²⁴ as well as in other oil-secreting plants pollinated by other oil-collecting bees.^{19,325} As this compound elicited antennal responses only in oil bees (*M. fulvipes*, *Redivia neliana*), but not in non-oil bees (*Apis mellifera*, *Melitta haemorrhoidalis*), it is a private communication channel between oil-secreting plants and oil-collecting bees.¹⁹



The aromatic 1,4-benzoquinone (**214**), a compound otherwise known to deter arthropods, was shown to be used by oligolectic *Hoplitis adunca* to recognize its *Echium* host plants.^{326,327} The flowers of *E. vulgare* released various compounds that were electroantennographically active, among them aromatics [e.g. **214**, 2-phenylethanol (**84**)] and terpenoids [(*Z*)- and (*E*)- β -ocimene (**62**), linalool (**60** and/or **61**)]. Synthetic mixtures of these EAD-active compounds only elicited attractive responses in the bees when **214** was present. Further, **214** alone was as attractive to flower-inexperienced bees as a complete synthetic mixture and a natural headspace sample demonstrating the key importance of this compound in host recognition of *Hoplitis adunca*.³²⁷



The narrowly oligolectic bee *Protodiscelis palpalis* is specialized on the aquatic plant *Hydrocleys martii* (Alismataceae), which itself depends on this bee species for sexual reproduction. The flower scent was strongly dominated by *p*-methylanisole (**156**), which successfully attracted the bees in bioassays. Compounds that occurred in smaller amounts [3,4-dimethoxytoluene (**215**), 3,4,5-trimethoxytoluene (**216**)] or not at all [2-methoxy-4-methylphenol (**217**)] in *H. martii*, but in higher amounts in the closely-related species *H. nymphoides*, a plant species not used by *P. palpalis* as host plant, were not attractive to *P. palpalis*.³²⁸

6.4.4 Male euglossine bees. Male euglossine bees (Apidae, Euglossini) are highly specialized pollinators as they collect floral scents, store them in their hind legs, and use them



during courtship behavior.^{329,330} They visit a high number of plants that are specialized on these bees as pollinators, and in turn a high number of compounds have been described to attract male euglossines.³³¹ Plants pollinated by these bees are typically strongly scented, but the bees also collect fragrances from plants that are only weakly or not at all scented to the human nose, such as from the bromeliad *Cryptanthus burlemarxii*. This species is mainly pollinated by *Eulaema* sp. and *Euglossa* sp. bees, and Milet-Pinheiro *et al.*²⁰ just recently identified from this bromeliad the least volatile and highest molecular weight compound known to attract euglossine males. Flowers released small amounts of the diterpene (+)-copalol (31), which were responsible for eliciting attraction, landing, and scent-gathering behavior of the bee pollinators.

6.5 Wasps

Around 10 000 species of wasps are pollinators,²¹⁷ and in the last decade bioactive compounds were identified for both social and solitary species. Brodmann *et al.*³³² studied the olfactory attractiveness of *Scrophularia umbrosa* (Scrophulariaceae) host plants to social *Vespula vulgaris* wasps and found that only a blend of antennal-active (Z)-3-hexenyl acetate (110), 1-hexanol (203), and (Z)-3-hexenol (118) was capable of attracting the wasps, whereas a blend of other antennal-active compounds [benzaldehyde (88), decanal (115), geranyl acetate (218), 6-methyl-5-hepten-2-one (109), nonanal (126), octanal (219), 1-octen-3-ol (220)] was inactive. In another study, social *Vespula* wasps responded to a high number of floral volatiles of their *Heracleum sphondylium* and *Hedera helix* host plants in electroantennographic measurements, among them aliphatic [e.g., 2-heptanone (188)], aromatic (e.g., 88) and nitrogen-bearing [e.g., 2-phenylacetone (92)] compounds as well as terpenoids [e.g., β -myrcene (76), 4-oxoisophorone (169)].³¹² For *H. helix* experiments with synthetic compounds were also performed and it was found that a synthetic mixture resembling the floral scent of this species was attractive to the wasps, but less than the natural olfactory cues. Besides social *Vespula* wasps, advances were made in the host finding of solitary fig wasps, which are involved in the highly specialized brood-site mutualisms with figs, and in generalist parasitoids. The Mediterranean fig, *Ficus carica*, attracts its highly specialized pollinator species, the fig wasp *Blastophaga psenes*, with volatiles, and, among the five major volatiles released by receptive figs, linalool (60 and/or 61) and oxides thereof (145, 174), the (–)-isomer of (E)- β -caryophyllene (64), and germacrene D (101) elicited antennal responses in *B. psenes*, whereas α -copaene (124) was inactive.³³³ In another fig species, *Ficus curtipes*, the scent is dominated by 6-methyl-5-hepten-2-ol (221) and 6-methyl-5-hepten-2-one (109). Both compounds elicited behavioral responses in the fig wasp pollinator *Eupristina* sp., but elicited different responses: 221 attracted the wasp from a distance, whereas 109 induced fig-entry behavior in the wasp.³³⁴ The egg parasitoid *Trissolcus basalis* uses buckwheat for nectaring and was successfully attracted by a blend of four floral [butanoic acid (222), 2-methylbutanoic acid (180), 3-methylbutanoic acid (181), pentanoic acid (223)] and two vegetative [(Z)-3-hexenyl acetate (110), (E,E)- α -farnesene (63)] volatiles.³³⁵



6.6 Other pollinators

Next to the common pollinators and flower visitors mentioned above, attractive compounds also were recently identified for uncommon pollinators/flower visitors. For example, ant pollinators (e.g., *Aphaenogaster senilis*) of *Cytinus hypocistis* (Cytinaceae) were shown to be attracted by (E)-cinnamaldehyde (155) and (E)-cinnamyl alcohol (154).³³⁶ Thrips (*Frankliniella occidentalis*) flower visitors of *Arabidopsis thaliana* were attracted by linalool (60 and/or 61) and repelled by lilac aldehyde (165) and lilac alcohol (224),^{59,337} and thrips were also shown to be attracted by (S)-verbenone (225), a compound released from pollen host plants (*Pinus* spp.).³³⁸ In a very recent intriguing study, Etl *et al.*²⁶ described a new plant bug pollination system, likely evolved from a cyclocephaline beetle pollination system, and decoded the chemical communication involved. They found that the aroid *Syngonium hastiferum* specializes on a single plant bug pollinator species, *Neella* sp. nov. (Miridae), that is attracted by the scent of the inflorescences. The scent was strongly dominated by the previously unknown natural product (Z)-3-isopropylpent-3-en-1-ol (12) (gambanol). In field bioassays, this compound specifically attracted the pollinator species.²⁶ Congeners of *Neella* sp. nov., such as *N. floridula*, visit cyclocephaline beetle pollinated aroids as non-pollinating florivores.³³⁹ One of those plants is *Dieffenbachia aurantiaca*. The floral scent of this species attracted as many plant bug florivores as natural inflorescences. It was dominated by (Z)-jasmone (127), and this compound was highly effective in attracting the plant bugs in field bioassays.³³⁹



The first floral attractant for ground-dwelling mammal pollinators was identified from *Cytinus visseri* (Cytinaceae). This species is pollinated by the striped fieldmouse, *R. pumilio*, the pygmy mouse, *Mus minutoides*, and the short-snouted elephant shrew, *E. brachyrhynchus*.³⁴⁰ Whole flowers of this species as well as separated nectar release a plastic-like scent, due to the emission of 1-hexen-3-one (226). Only whole flowers released another major compound, i.e. 3-hexanone (227). Behavioral experiments with the striped fieldmouse and the short-snouted elephant shrew showed that the natural floral scent is highly attractive to these mammal pollinators. Y-Maze experiments with the striped fieldmouse further demonstrated that this mouse species is attracted by 227,



whereas **226** has repellent properties, but only when tested alone and not when tested in combination with **227**.³⁴⁰ The African pineapple lily *Eucomis regia*, similar to *C. visseri*, is also pollinated by mice and elephant-shrews that are attracted by the scent of the plant species. Flowers of this lily released common [e.g. 2-heptanone (**188**), 2-heptanol (**228**)] and uncommon [e.g. *exo*-brevicomine (**229**), methional (**230**)] scents, of which **230** was responsible for the boiled-potato-like smell of the flowers and of separated nectar. This compound successfully attracted *Namaqua* rock mouse (*Micaelamys namaquensis*) pollinators in y-maze experiments.³⁴¹ Just recently, Johnson and Govender³⁴² tested the effects of various volatiles found in rodent pollinated plants and in insect pollinated congeners (aromatic compounds) on their attractiveness to mouse pollinators. Several compounds were attractive to the mice, among them the aliphatic ketones **227**, **188** and acetoin (**130**), the aliphatic ester ethyl butyrate (**231**) and the aliphatic acid butanoic acid (**222**). Dimethyl disulfide (**182**) and the aromatic compounds [benzaldehyde (**88**), methyl benzoate (**100**), 2-phenylethanol (**84**), phenylacetaldehyde (**83**)] were neutral or repellent to the mice. Adding unattractive compounds to attractive ones resulted in a non-attractive signal.³⁴²



Finally, bioactive compounds were identified for crab spiders that visit flowers to hunt for prey (pollinators, florivores). Knauer *et al.*³⁴³ asked whether the crab spider *Thomisus onustus*, which frequently visits flowers of the buckler mustard (*Biscutella laevigata*), is attracted by floral volatiles. Indeed, the spider was attracted by β -ocimene (**62**), whereas the other two main floral scent compounds [*p*-anisaldehyde (**111**), 2-aminobenzaldehyde (**166**)] did not elicit a behavioral response in the spiders. Florivores induced the emission of (**62**). Accordingly, spiders had an olfactory preference for plants infested with florivores.³⁴³

7 Functions of floral volatiles other than pollinator attraction

Flower volatiles did not only evolve to attract pollinators, but have various other functions,^{344–346} such as to repel florivores, to fight against pathogenic microorganisms, or to attract enemies of plant pests. Several of these functions have only recently been understood, and here we summarize the main advances and present the compounds involved.

7.1 Repelling florivores

Ants are frequent florivores of flowers, and in a study on *Phlox paniculata* and *Lasius niger* ants it was demonstrated that floral scent is repellent to the ants.³⁴⁷ The ant repellent properties were lost after inhibition of terpene biosynthesis pointing to the

importance of terpenes in repelling the ants. Indeed, linalool (**60** and/or **61**), which was strongly inhibited³⁴⁷ and known to be repellent to ants before,³⁴⁸ repelled the ants in olfactometer assays. However, linalool (**60** and/or **61**) was only repellent when offering the compound in concentrations higher than released from the flowers, suggesting that other compounds are also involved in repelling ants from the flowers.³⁴⁷ Other flower volatiles known to be repellent to ants are *p*-anisaldehyde (**111**) from *Petasites fragrans* flowers³⁴⁹ and 2-phenylethanol (**84**) from flowers of the alpine skypilot *Polemonium viscosum*.³⁵⁰ In the latter species, the flowers are polymorphic in the amount of **84** emitted. Ants avoided high emitters in the field and died when exposed to **84** in the laboratory.³⁵⁰ Other florivores than ants shown to be repelled by floral scents were beetles and crickets. By using genetically modified *Nicotiana attenuata*, Kessler *et al.*³⁵¹ demonstrated that benzylacetone (**91**) is repellent to the florivorous cucumber beetles *Diabrotica undecimpunctata*, and similar experiments with *Petunia x hybrida* revealed that isoeugenol (**232**) and benzyl benzoate (**233**) are repellent to *D. undecimpunctata* and the snowy tree cricket, *Oecanthus fultoni*.²⁴⁴ The bush cricket *Metrioptera bicolor*, which mainly feeds on vegetative plant parts, but occasionally also on flowers, was repelled by the common floral scents linalool (**60** and/or **61**) and β -caryophyllene (**64**).³⁵²

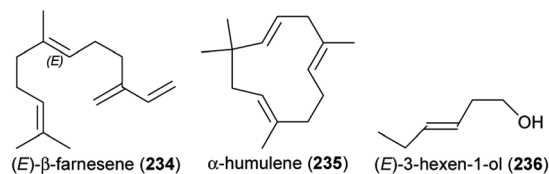


7.1.1 Defense against pathogenic microorganisms. A floral volatile shown to defend against pathogens is (*E*)- β -caryophyllene (**64**) in *A. thaliana*, where this compound is a major volatile of the stigma.³⁵³ Flowers of genetically modified plants that lacked emission of this compound showed greater bacterial growth on their stigmas than did flowers of wild-type plants, and their seeds were lighter and showed a high frequency of abnormal shape.³⁵³ Similar results were obtained in *Petunia hybrida*. Downregulation of a terpene synthase responsible for the production of floral germacrene D (**101**), bicyclgermacrene (**102**) and germacrene D-4-ol (**103**), resulted in an increase in a *Pseudomonas* bacterium on the pistils.¹⁶ Interestingly, these compounds are released from the petals into floral buds, accumulate in the stigma and in anthers, and do not only have defense functions, but also regulate pistil growth and seed yield.¹⁶ Another terpenoid, linalool (**60** and/or **61**), had a stronger negative effect on growth rate of bacteria that colonize flowers of the foxglove beard-tongue, *Penstemon digitalis*, than on bacteria colonizing leaves of this plant species.³⁵⁴ In addition to terpenoids, a nitrogen-bearing compound and an aromatic floral compound were also shown to be used as defense against bacteria. In *Saponaria officinalis*, the diversity of epiphytic bacteria was much lower on flowers than leaves, especially due to floral 2-phenylacetone nitrile (**92**) and 2-phenylethanol (**84**). These two compounds had strong growth-inhibitory effects on various bacteria, whereas floral methyl benzoate (**100**) and vegetative (*Z*)-3-



hexenol (**118**) and (*Z*)-3-hexenyl acetate (**110**) had only weak inhibitory effects.³⁵⁵ Finally, in strawberry, which varies in floral scents among genotypes, statistical models suggest that terpenes [e.g. α - and β -pinene (**72** and **73**), isomers of ocimene, e.g. **62**] and aromatic compounds [e.g. *p*-anisaldehyde (**111**), benzaldehyde (**88**)] especially shape the floral bacterial and fungal communities.¹⁷⁹

7.1.2 Attraction of enemies of plant pests. Two studies recently identified compounds attractive to beetle predators that feed on aphid plant pests. The first study showed that pyrethrum (*Tanacetum cinerariifolium*) flowers release (*E*)- β -farnesene (**234**), a compound well known as an alarm pheromone of aphids, to attract ladybird beetles.³⁴⁶ In addition, **234** accumulates in the vascular system of the flower receptacle and the peduncle, preferred feeding sites of aphids. If aphids feed on these tissues, they take it up, and excrete it together with honeydew. From there, **234** is emitted and induces an alarm response in the aphids. Thus, in pyrethrum, **234** prevents attack by aphids and recruits aphid predators.³⁴⁶ The second study tested 58 floral volatiles in olfactometer assays on the ladybirds (Coleoptera: Coccinellidae) *Harmonia axyridis* and *Coccinella septempunctata*, and found that 3-methylbutyl acetate (**132**), α -humulene (**235**), (*E*)-3-hexen-1-ol (**236**), methyl salicylate (**125**) and β -pinene (**73**) are attractive to both of them.⁴⁵ Another study demonstrated that the wasp *Apanteles taragamae*, a parasitoid of the legume pod borer *Maruca vitrata*, is attracted by floral volatiles (compounds were not identified) of cowpea (*Vigna unguiculata*) that were induced by florivory of *M. vitrata*.³⁵⁶

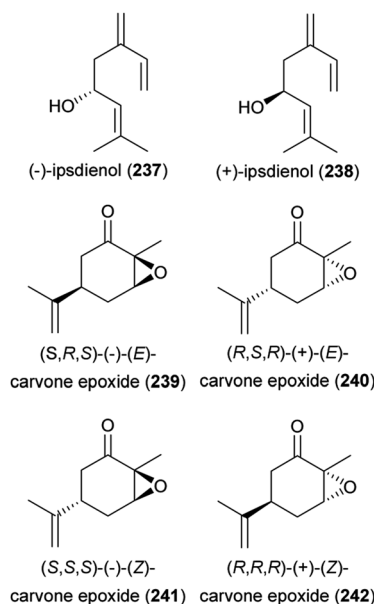


8 Chirality in flower scents and its importance for interspecific interactions

Many floral scent components are optically active and it is known that chirality is of relevance in flower-animal interactions.^{5,357–359} However, despite this knowledge there are very few recent studies that determined the absolute configuration of compounds from floral scents and tested single enantiomers or reconstructed enantiomeric ratios on flower visitors in physiological and/or behavioral experiments (see also Perkins *et al.*¹⁸).

In the pollinator-plant and herbivore-plant association between *Manduca sexta* and *Datura wrightii* it was shown that floral (*S*)-linalool (**61**) elicited oviposition behavior, while floral (*R*)-linalool (**60**) had negative effects on oviposition behaviour.³⁶⁰ Neither the single enantiomers nor mixtures of the two enantiomers had effects on the feeding behavior of the moths, which is triggered by other floral volatiles (mixture of benzaldehyde (**88**), benzyl alcohol (**99**), geraniol (**59**) and nerol). Field bean (*Vicia faba*) flowers only released **60**, and this enantiomer

attracted the bean seed beetle *Bruchus rufimanus*, a pest of the field bean.²³⁶ In the pollination mutualism between Phyllanthaceae and *Epicephala* moths, the different enantiomers of linalool (**60** and **61**) might elicit different behaviors in the moths. In both *Glochidion obovatum* and *Glochidion rubrum* female flowers released (*S*)-linalool (**61**) and male flowers (*R*)-linalool (**60**). Okamoto *et al.*²⁶⁵ speculated that this sexual difference in the emission of linalool enantiomers allows moth pollinators to discriminate between the flower sexes and collect pollen on male flowers and actively pollinate and oviposit on female flowers (see above). In another nursery pollination mutualism, between *Lithophragma bolanderi* and *Greya politella* moths, it was just recently demonstrated that the flowers produce only the (6*R*,10*R*)-enantiomer of 6,10,14-trimethylpentadecan-2-one (**167**), and that this enantiomer is elicits antennal responses in the moth pollinators.²⁶¹ This same compound was also detected in tibial fragrances of male euglossine bees (*Euglossa* spp.), which likely obtained this compound from aroid or orchid flowers, and in behavioral assays, it attracted more individuals and species of euglossines than a racemate of **167**.³⁶¹



Another optically active attractant for male euglossines is ipsdienol. This compound is released from various male euglossine bee pollinated plants,^{331,362} and it was demonstrated for *Euglossa cyanura* males that only (-)-ipsdienol (**237**) is attractive, whereas the (+)-isomer (**238**) is ignored by the bees.³⁶³ Further, antennae of these bees showed larger physiological responses to the (-)-isomer than to the (+)-isomer. Another bee species, *E. mixta*, which is not attracted by ipsdienol, did not discriminate between the enantiomers in electroantennographic measurements.³⁶³ New data were also obtained for carvone epoxide. This compound is released from plants of several families (e.g., Annonaceae, Araceae, Gesneriaceae, Orchidaceae) and is a main attractant for male euglossine bees of the genus *Eulaema*.³³¹ The plants mainly release (*E*)-carvone epoxides, though some also release smaller amounts of the (*Z*)-



isomers (summarized by Brandt *et al.*³⁶⁴). For each diastereomer two enantiomers are possible, summing up to a total of four stereoisomers of carvone epoxide (239–242). All five orchid species recently studied for the presence of the stereoisomers of carvone epoxide released only the (–)-(E)-isomer (239). In *Eulaema* bees, this enantiomer elicited the highest antennal responses and was the most attractive.³⁶⁴

9 Conclusions

Our review provides clear evidence that the number of described floral volatiles is still increasing. While this increase is due partly to advances in analytical procedures, it results mainly from the increasing number of plant species being studied motivated by both basic research on pollination biology and the search for new volatiles by the flavor and fragrance industry. Some of the newly discovered volatiles are derived from common floral volatiles, such as (1*S*,2*S*)-isojasmol (3), (1*S*,2*S*)-isojasmyl acetate (4), and (S)-dehydrojasmone (5) detected in various Araceae species, which are derived from (Z)-jasmone (Fig. 25).¹¹

Another example is a group of new volatiles of *Yucca* species, including (Z)- and (E)-filamentol (22 and 25), (Z)- and (E)-filamentol (23 and 26), (Z)- and (E)-filamentolide (24 and 27) and filamentone (28), and of (E)-4,8-dimethyl-1,3,7-nonatrien-5-yl acetate (29) and the (Z)-isomer thereof from *Philodendron squamiferum*, all derived from the widespread 4,8-dimethyl-1,3,7-nonatriene (see Fig. 26).^{11,29}

The continuous increase in the number of known floral volatiles also shows that our understanding of the floral scent

chemistry is far from being complete, calling for studies that analyse floral volatiles (including their absolute configuration) from plant lineages and species where no data are available so far.

Major advances during the period of this review were also made in understanding the biosynthesis of floral volatiles and the regulation of volatile emissions. Several new genes and biosynthetic routes were discovered, among them alternative pathways, which sometimes work in parallel in specific flowers, *e.g.* for production of 2-phenylethanol (84), of one of the most widespread floral volatiles. Similarly, several new transcription factors and other factors that regulate floral scent production and emission, such as epigenetic influences and long non-coding RNAs, were discovered. A breakthrough was the identification of a transporter protein that exports floral volatiles from cells.

Major advances were also achieved in our understanding of how floral scents are shaped by abiotic factors, such as air pollution and increased temperatures, but how, *e.g.* temperature-induced changes in scent translate into pollinator behavior is less well understood. Several studies demonstrated in the last years that not only herbivores but also microorganisms (*e.g.*, mycorrhizal fungi, plant endophytes, flower-dwelling bacteria and yeasts) have effects on flower scent emissions and sometimes also on the communication with flower visitors. Most of these studies were performed in the laboratory or using experimental settings, and it remains to be tested how microorganisms affect floral scents and flower–animal interactions in natural settings.

Since 2010, a large number of studies identified compounds attractive to flower visitors. The behaviorally active compounds were main compounds of the released scent in some pollination systems, *e.g.*,³⁶⁵ while they were only minor or even trace compounds in others. We now have a much better understanding how plants communicate with beetle, lepidopteran, bee and fly pollinators, and the first attractants were also identified for less common pollinators, such as rodents, shrews and true bugs. Despite these advances, our knowledge about the communication between flowers and their visitors is still very limited, especially when considering the high number of plants pollinated by animals and the high number of animals involved. Also, most advances were made in specialized pollination systems, and more studies are required to elucidate olfactory communication in generalist systems. Considering the chemical properties of the compounds involved in pollinator attraction, such as volatility and the functional groups present (*e.g.*, alcohol, olefin, ester, ketone), there seem to be no trends for different groups of pollinators (bees, beetles, flies, Lepidoptera, *etc.*). However, there are some trends regarding overall



Fig. 25 The widespread floral scent (Z)-jasmone and uncommon putative derivatives thereof.

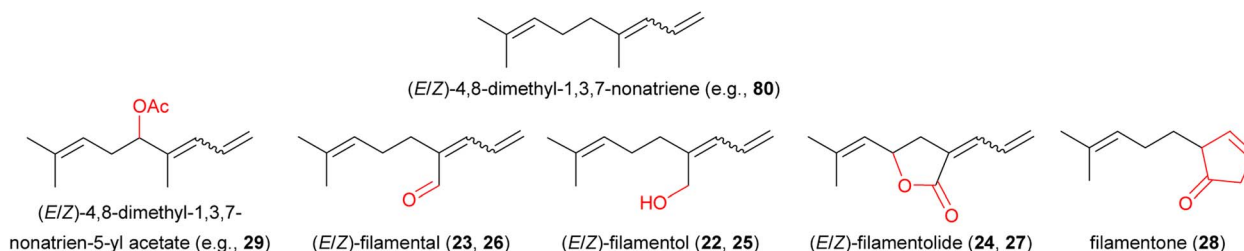


Fig. 26 The widespread floral scent (E/*Z*)-4,8-dimethyl-1,3,7-nonatriene (*e.g.*, 80) and uncommon putative derivatives thereof.



biosynthetic class. For instance, among reported beetle attractants fatty acid derivatives are well represented with a smaller number of compounds from other classes.

Lepidoptera seem to respond mainly to phenylpropanoids/benzenoids, terpenes and nitrogen-bearing compounds, and flies additionally to fatty acid derivatives and sulphur-bearing compounds. For bees, reported attractants were mainly phenylpropanoids/benzenoids, terpenes, and fatty acid derivatives, but only a few nitrogen- and no sulphur-bearing compounds have been described. For wasps, there are a lot of reports of fatty acid derivatives, but also several terpenes and some compounds from other classes (see also ref. 18), and for ground-dwelling mammals mainly fatty acid derivatives are listed as attractants with the conspicuous absence of terpenes and phenylpropanoids/benzenoids. In fact, phenylpropanoids/benzenoids, which are otherwise common among floral scents and attractive to other pollinators [e.g. phenylacetaldehyde (**83**), 2-phenylethanol (**84**)] were even shown to be repellent to rodents. Finally, a few floral volatiles were identified during the period of this review that serve plants in functions other than pollinator attraction, such as by repelling florivores, inhibiting pathogenic microorganisms, and attracting enemies of herbivores. Considering the abundance and chemical diversity of floral volatiles, there are relatively few studies in this “young” field of research and many new insights can be expected in the near future.

10 Author contributions

Both authors worked together on the manuscript.

11 Conflicts of interest

There are no conflicts to declare.

12 Acknowledgements

SD thanks Roman Fuchs for help with the chemical structures and Karin Gross, Danae Laina, Monica Barman and other members of the Plant Ecology group as well as Robert R. Junker for comments on an earlier version of the manuscript. JG thanks Kirsten Heinrichs for help in literature searching, and Eran Pichersky for comments on an earlier draft. SD and JG thank two anonymous reviewers for helpful comments on an earlier version of the manuscript.

13 References

- H. Bouwmeester, R. C. Schuurink, P. M. Bleeker and F. Schiestl, *Plant J.*, 2019, **100**, 892–907.
- H. E. M. Dobson, in *Biology of Floral Scent*, ed. N. Dudareva and E. Pichersky, CRC Press, Boca Raton, 2006, pp. 147–198.
- R. R. Junker and A. L. Parachnowitsch, *J. Indian Inst. Sci.*, 2015, **95**, 43–67.
- J. T. Knudsen, R. Eriksson, J. Gershenzon and B. Ståhl, *Bot. Rev.*, 2006, **72**, 1–120.
- R. A. Raguso, *Annu. Rev. Ecol. Evol. Syst.*, 2008, **39**, 549–569.
- R. A. Raguso, in *Biology of Plant Volatiles*, eds. E. Pichersky and N. Dudareva, CRC Press, Boca Raton, 2020, pp. 297–325.
- J. Ollerton, R. Winfree and S. Tarrant, *Oikos*, 2011, **120**, 321–326.
- A. M. Klein, B. E. Vaissière, J. H. Cane, I. Steffan-Dewenter, S. A. Cunningham, C. Kremen and T. Tscharrntke, *Proc. R. Soc. B*, 2007, **274**, 303–313.
- B. Bohman, M. M. Y. Tan, R. D. Phillips, A. Scaffidi, A. N. Sobolev, S. A. Moggach, G. R. Flematti and R. Peakall, *Angew. Chem., Int. Ed.*, 2020, **59**, 1124–1128.
- K. Lukas, T. Harig, S. Schulz, J. Hadersdorfer and S. Dötterl, *Chemoecology*, 2019, **29**, 211–223.
- A. C. D. Maia, C. Grimm, M. Schubert, F. Etl, E. G. Gonçalves, D. M. Do Amaral Ferraz Navarro, S. Schulz and S. Dötterl, *J. Chem. Ecol.*, 2019, **45**, 204–213.
- H. Teichert, S. Dötterl and G. Gottsberger, *Plant Syst. Evol.*, 2018, **304**, 831–839.
- L. A. Burkle and J. B. Runyon, *Funct. Ecol.*, 2019, **33**, 2116–2129.
- R. R. Junker, N. Höcherl and N. Blüthgen, *J. Anim. Ecol.*, 2010, **79**, 818–823.
- A. Kantsa, R. A. Raguso, A. G. Dyer, J. M. Olesen, T. Tscheulin and T. Petanidou, *Nat. Commun.*, 2018, **9**, 1041.
- B. Boachon, Y. Burdloff, J.-X. Ruan, R. Rojo, R. R. Junker, B. Vincent, F. Nicolè, F. Bringel, A. Lesot, L. Henry, J.-E. Bassard, S. Mathieu, L. Allouche, I. Kaplan, N. Dudareva, S. Vuilleumier, L. Miesch, F. André, N. Navrot, X.-Y. Chen and D. Werck-Reichhart, *Plant Cell*, 2019, **31**, 2947–2972.
- H. M. Li, W. B. Liu, L. L. Yang, H. Q. Cao, P. Pelosi, G. R. Wang and B. Wang, *J. Agric. Food Chem.*, 2020, **68**, 12212–12220.
- J. Perkins, T. Hayashi, R. Peakall, G. R. Flematti and B. Bohman, *Nat. Prod. Rep.*, 2023, **40**, 819–839.
- I. Schäffler, K. E. Steiner, M. Haid, S. S. van Berkel, G. Gerlach, S. D. Johnson, L. Wessjohann and S. Dötterl, *Sci. Rep.*, 2015, **5**, 12779.
- P. Milet-Pinheiro, A. Domingos-Melo, J. B. Olivera, N. S. L. Albuquerque, A. C. G. Costa, S. Albuquerque-Lima, M. F. R. Silva, D. M. A. F. Navarro, A. C. D. Maia, L.-L. Gundersen, M. Schubert, S. Dötterl and I. C. Machado, *Curr. Biol.*, 2021, **31**, 860–868.
- T. Hayashi, B. Bohman, A. Scaffidi, R. Peakall and G. R. Flematti, *Curr. Biol.*, 2021, **31**, 1954–1961.
- A. Heiduk, I. Brake, M. von Tschirnhaus, M. Göhl, A. Jürgens, S. D. Johnson, U. Meve and S. Dötterl, *Curr. Biol.*, 2016, **26**, 2787–2793.
- P. Stamm, F. Etl, A. C. D. Maia, S. Dötterl and S. Schulz, *J. Org. Chem.*, 2021, **86**, 5245–5254.
- R. Kaiser, *Scent of the vanishing flora*, Wiley VCH, 2011.
- C. Cohen, W. R. Liltved, J. F. Colville, A. Shuttleworth, J. Weissflog, A. Svatoš, B. Bytebier and S. D. Johnson, *Curr. Biol.*, 2021, **31**, 1962–1969.
- F. Etl, C. Kaiser, O. Reiser, M. Schubert, S. Dötterl and J. Schönenberger, *Curr. Biol.*, 2022, **32**, 4688–4698.



- 27 B. Bohman, A. M. Weinstein, R. D. Phillips, R. Peakall and G. R. Flematti, *J. Nat. Prod.*, 2019, **82**, 1107–1113.
- 28 R. Peakall, D. Ebert, J. Poldy, R. A. Barrow, W. Francke, C. C. Bower and F. P. Schiestl, *New Phytol.*, 2010, **188**, 437–450.
- 29 A. Tröger, G. P. Svensson, H.-M. Galbrecht, R. Twele, J. M. Patt, S. Bartram, P. H. G. Zarbin, K. A. Segraves, D. M. Althoff, S. von Reuss, R. A. Raguso and W. Francke, *J. Chem. Ecol.*, 2021, **47**, 1025–1041.
- 30 C. E. P. Nunes, M. F. G. V. Peñafior, J. M. S. Bento, M. J. Salvador and M. Sazima, *Oecologia*, 2016, **182**, 933–946.
- 31 A. Heiduk, J.-P. Haenni, U. Meve, S. Schulz and S. Dötterl, *J. Comp. Physiol.*, A, 2019, **205**, 301–310.
- 32 B. Bohman, L. Jeffares, G. Flematti, L. T. Byrne, B. W. Skelton, R. D. Phillips, K. W. Dixon, R. Peakall and R. A. Barrow, *J. Nat. Prod.*, 2012, **75**, 1589–1594.
- 33 B. Bohman, L. Jeffares, G. Flematti, R. D. Phillips, K. W. Dixon, R. Peakall and R. A. Barrow, *Org. Lett.*, 2012, **14**, 2576–2578.
- 34 A. C. D. Maia, S. Dötterl, R. Kaiser, I. Silberbauer-Gottsberger, H. Teichert, M. Gibernau, D. M. do Amaral Ferraz Navarro, C. Schlindwein and G. Gottsberger, *J. Chem. Ecol.*, 2012, **38**, 1072–1080.
- 35 B. Bohman, R. D. Phillips, G. R. Flematti, R. A. Barrow and R. Peakall, *Angew. Chem., Int. Ed.*, 2017, **56**, 8455–8458.
- 36 S. Wakamura, N. Arakaki, D. Moriyama, S. Kanayama, M. Oike, A. Kimura, S. Wajima, H. Ono and H. Yasui, *Chemoecology*, 2020, **30**, 49–57.
- 37 J. Stökl, A. Strutz, A. Dafni, A. Svatos, J. Doubsky, M. Knaden, S. Sachse, B. S. Hansson and M. C. Stensmyr, *Curr. Biol.*, 2010, **20**, 1846–1852.
- 38 F. Chen, J. C. D'Auria, D. Tholl, J. R. Ross, J. Gershenzon, J. P. Noel and E. Pichersky, *Plant J.*, 2003, **36**, 577–588.
- 39 Y. Jiang, G. Liu, W. Zhang, C. Zhang, X. Chen, Y. Chen, C. Yu, D. Yu, J. Fu and F. Chen, *Phytochemistry*, 2021, **191**, 112899.
- 40 Q. Xu, S. Wang, H. Hong and Y. Zhou, *BMC Genomics*, 2019, **20**, 125.
- 41 F. Chen, D. Tholl, J. Bohlmann and E. Pichersky, *Plant J.*, 2011, **66**, 212–229.
- 42 J. Degenhardt, T. G. Köllner and J. Gershenzon, *Phytochemistry*, 2009, **70**, 1621–1637.
- 43 X. Zeng, C. Liu, R. Zheng, X. Cai, J. Luo, J. Zou and C. Wang, *Front. Plant Sci.*, 2016, **6**, 1232.
- 44 M. Lv, X. Sun, D. Li, G. Wei, L. Liu, F. Chen, Y. Cai and H. Fan, *Ind. Crops Prod.*, 2022, **182**, 114875.
- 45 J. Zhao, Z. Wang, Z. Li, J. Shi, L. Meng, G. Wang, J. Cheng and Y. Du, *J. Asia-Pac. Entomol.*, 2020, **23**, 1023–1029.
- 46 H. Huang, Y.-W. Kuo, Y.-C. Chuang, Y.-P. Yang, L.-M. Huang, M.-F. Jeng, W.-H. Chen and H.-H. Chen, *Front. Plant Sci.*, 2021, **12**, 700958.
- 47 C. Zhao, Z. Yu, J. A. T. d. Silva, C. He, H. Wang, C. Si, M. Zhang, D. Zeng and J. Duan, *Int. J. Mol. Sci.*, 2020, **21**, 7005.
- 48 Y. Yue, R. Yu and Y. Fan, *Planta*, 2014, **240**, 745–762.
- 49 G. Liu, M. Yang, X. Yang, X. Ma and J. Fu, *J. Plant Physiol.*, 2021, **258–259**, 153358.
- 50 T. Bao, K. Shadrack, S. Yang, X. Xue, S. Li, N. Wang, Q. Wang, L. Wang, X. Gao and Q. Cronk, *Plant Cell Physiol.*, 2020, **61**, 1733–1749.
- 51 J. Jin, M. J. Kim, S. Dhandapani, J. G. Tjhang, J.-L. Yin, L. Wong, R. Sarojam, N.-H. Chua and I.-C. Jang, *J. Exp. Bot.*, 2015, **66**, 3959–3975.
- 52 E. Pichersky and E. Lewinsohn, *Annu. Rev. Plant Biol.*, 2011, **62**, 549–566.
- 53 J.-i. Hattori, K. Shindo, T. Sasaki, F. Ohno, H. Tokuda, K. Ishikawa and N. Misawa, *Sci. Rep.*, 2018, **8**, 12474.
- 54 A. Fährnrich, A. Brosemann, L. Teske, M. Neumann and B. Piechulla, *Plant Mol. Biol.*, 2012, **79**, 537–553.
- 55 K. J. R. P. Byers, J. P. Vela, F. Peng, J. A. Riffell and H. D. Bradshaw Jr, *Plant J.*, 2014, **80**, 1031–1042.
- 56 J.-L. Magnard, A. Rocca, J.-C. Caissard, P. Vergne, P. Sun, R. Hecquet, A. Dubois, L. Hibrand-Saint Oyant, F. Jullien, F. Nicolè, O. Raymond, S. Huguet, R. Baltenweck, S. Meyer, P. Claudel, J. Jeauffre, M. Rohmer, F. Foucher, P. Hugueney, M. Bendahmane and S. Baudino, *Science*, 2015, **349**, 81–83.
- 57 L. Sheng, S. Zang, J. Wang, T. Wei, Y. Xu and L. Feng, *PeerJ*, 2021, **9**, e11098.
- 58 J.-F. Ginglinger, B. Boachon, R. Höfer, C. Paetz, T. G. Köllner, L. Miesch, R. Lugan, R. Baltenweck, J. Mutterer, P. Ullmann, F. Beran, P. Claudel, F. Verstappen, M. J. C. Fischer, F. Karst, H. Bouwmeester, M. Miesch, B. Schneider, J. Gershenzon, J. Ehling and D. Werck-Reichhart, *Plant Cell*, 2013, **25**, 4640–4657.
- 59 B. Boachon, R. R. Junker, L. Miesch, J.-E. Bassard, R. Höfer, R. Caillieaudeaux, D. E. Seidel, A. Lesot, C. Heinrich, J.-F. Ginglinger, L. Allouche, B. Vincent, D. S. C. Wahyuni, C. Paetz, F. Beran, M. Miesch, B. Schneider, K. Leiss and D. Werck-Reichhart, *Plant Cell*, 2015, **27**, 2972–2990.
- 60 S. Lee, S. Badieyan, D. R. Bevan, M. Herde, C. Gatz and D. Tholl, *Proc. Natl. Acad. Sci. U. S. A.*, 2010, **107**, 21205–21210.
- 61 A. Richter, C. Schaff, Z. Zhang, A. E. Lipka, F. Tian, T. G. Köllner, C. Schnee, S. Preiß, S. Irmisch, G. Jander, W. Boland, J. Gershenzon, E. S. Buckler and J. Degenhardt, *Plant Cell*, 2016, **28**, 2651–2665.
- 62 R. Sohrabi, J.-H. Huh, S. Badieyan, L. H. Rakotondraibe, D. J. Kliebenstein, P. Sobrado and D. Tholl, *Plant Cell*, 2015, **27**, 874–890.
- 63 S. Baldermann, M. Kato, M. Kurosawa, Y. Kurobayashi, A. Fujita, P. Fleischmann and N. Watanabe, *J. Exp. Bot.*, 2010, **61**, 2967–2977.
- 64 A. Rubio-Moraga, J. L. Rambla, A. Fernández-de-Carmen, A. Trapero-Mozos, O. Ahrazem, D. Orzáez, A. Granell and L. Gómez-Gómez, *Plant Mol. Biol.*, 2014, **86**, 555–569.
- 65 Y. Kaminaga, J. Schnepf, G. Peel, C. M. Kish, G. Ben-Nissan, D. Weiss, I. Orlova, O. Lavie, D. Rhodes, K. Wood, D. M. Porterfield, A. J. L. Cooper, J. V. Schloss, E. Pichersky, A. Vainstein and N. Dudareva, *J. Biol. Chem.*, 2006, **281**, 23357–23366.



- 66 M. Farhi, O. Lavie, T. Masci, K. Hendel-Rahmanim, D. Weiss, H. Abeliovich and A. Vainstein, *Plant Mol. Biol.*, 2010, **72**, 235–245.
- 67 M. Gutensohn, A. Klempien, Y. Kaminaga, D. A. Nagegowda, F. Negre-Zakharov, J.-H. Huh, H. Luo, R. Weizbauer, T. Mengiste, D. Tholl and N. Dudareva, *Plant J.*, 2011, **66**, 591–602.
- 68 M. Sakai, H. Hirata, H. Sayama, K. Sekiguchi, H. Itano, T. Asai, H. Dohra, M. Hara and N. Watanabe, *Biosci., Biotechnol., Biochem.*, 2007, **71**, 2408–2419.
- 69 X.-M. Chen, H. Kobayashi, M. Sakai, H. Hirata, T. Asai, T. Ohnishi, S. Baldermann and N. Watanabe, *J. Plant Physiol.*, 2011, **168**, 88–95.
- 70 A. Rocca, L. Hibrand-Saint Oyant, E. Cavel, J.-C. Caissard, J. Machenaud, T. Thouroude, J. Jeauffre, A. Bony, A. Dubois, P. Vergne, J. Szécsi, F. Foucher, M. Bendahmane and S. Baudino, *Plant Physiol.*, 2019, **179**, 1064–1079.
- 71 T. Koeduka, Y. Ueyama, S. Kitajima, T. Ohnishi and K. Matsui, *J. Plant Physiol.*, 2020, **252**, 153245.
- 72 H. Hirata, T. Ohnishi, K. Tomida, H. Ishida, M. Kanda, M. Sakai, J. Yoshimura, H. Suzuki, T. Ishikawa, H. Dohra and N. Watanabe, *Sci. Rep.*, 2016, **6**, 20234.
- 73 L. Sheng, Y. Zeng, T. Wei, M. Zhu, X. Fang, X. Yuan, Y. Luo and L. Feng, *Genes*, 2018, **9**, 576.
- 74 S. Dhandapani, J. Jin, V. Sridhar, N.-H. Chua and I.-C. Jang, *Plant Physiol.*, 2019, **180**, 171–184.
- 75 J. Günther, N. D. Lackus, A. Schmidt, M. Huber, H.-J. Stödtler, M. Reichelt, J. Gershenzon and T. G. Köllner, *Plant Physiol.*, 2019, **180**, 767–782.
- 76 F. Dong, Z. Yang, S. Baldermann, Y. Kajitani, S. Ota, H. Kasuga, Y. Imazeki, T. Ohnishi and N. Watanabe, *J. Plant Physiol.*, 2012, **169**, 217–225.
- 77 F. Dong, Y. Zhou, L. Zeng, Q. Peng, Y. Chen, L. Zhang, X. Su, N. Watanabe and Z. Yang, *Molecules*, 2016, **21**, 1106.
- 78 Y. Zhou, Q. Peng, L. Zeng, J. Tang, J. Li, F. Dong and Z. Yang, *Food Chem.*, 2018, **258**, 352–358.
- 79 Y. Zhou, Q. Peng, L. Zhang, S. Cheng, L. Zeng, F. Dong and Z. Yang, *Food Chem.*, 2019, **280**, 27–33.
- 80 T. Koeduka, E. Fridman, D. R. Gang, D. G. Vassão, B. L. Jackson, C. M. Kish, I. Orlova, S. M. Spassova, N. G. Lewis, J. P. Noel, T. J. Baiga, N. Dudareva and E. Pichersky, *Proc. Natl. Acad. Sci. U. S. A.*, 2006, **103**, 10128–10133.
- 81 N. F. M. Shaipulah, J. K. Muhlemann, B. D. Woodworth, A. Van Moerkercke, J. C. Verdonk, A. A. Ramirez, M. A. Haring, N. Dudareva and R. C. Schuurink, *Plant Physiol.*, 2015, **170**, 717–731.
- 82 T. Zhang, T. Huo, A. Ding, R. Hao, J. Wang, T. Cheng, F. Bao and Q. Zhang, *PLoS One*, 2019, **14**, e0223974.
- 83 H. Yan, S. Baudino, J.-C. Caissard, N. Florence, H. Zhang, K. Tang, S. Li and S. Lu, *Plant Physiol. Biochem.*, 2018, **129**, 21–26.
- 84 J. R. Widhalm and N. Dudareva, *Mol. Plant*, 2015, **8**, 83–97.
- 85 A. V. Qualley, J. R. Widhalm, F. Adebisin, C. M. Kish and N. Dudareva, *Proc. Natl. Acad. Sci. U. S. A.*, 2012, **109**, 16383–16388.
- 86 T. A. Akhtar and E. Pichersky, *Plant Physiol.*, 2013, **162**, 52–62.
- 87 K. Van Gelder, T. Forrester and T. A. Akhtar, *Planta*, 2020, **252**, 3.
- 88 A. K. Gupta, T. A. Akhtar, A. Widmer, E. Pichersky and F. P. Schiestl, *BMC Plant Biol.*, 2012, **12**, 158.
- 89 T. Koeduka, M. Kajiyama, T. Furuta, H. Suzuki, T. Tsuge and K. Matsui, *J. Biosci. Bioeng.*, 2016, **122**, 679–684.
- 90 T. Koeduka, M. Kajiyama, H. Suzuki, T. Furuta, T. Tsuge and K. Matsui, *Planta*, 2016, **244**, 725–736.
- 91 Y. Wang, H. Zhang, C. Wan, X. He, J. Huang, M. Lyu, Y. Yuan and B. Wu, *Plants*, 2022, **11**, 13.
- 92 M. Kang, Y. Choi, H. Kim and S.-G. Kim, *New Phytol.*, 2022, **234**, 527–544.
- 93 H. Guo, N. D. Lackus, T. G. Köllner, R. Li, J. Bing, Y. Wang, I. T. Baldwin and S. Xu, *Mol. Biol. Evol.*, 2019, **37**, 1090–1099.
- 94 I. Abe, Y. Takahashi, H. Morita and H. Noguchi, *Eur. J. Biochem.*, 2001, **268**, 3354–3359.
- 95 R. A. Raguso, A. Kelber, M. Pfaff, R. A. Levin and L. A. McDade, *Ann. Mo. Bot. Gard.*, 2007, **94**, 236–257.
- 96 L. J. Nielsen and B. L. Møller, *Phytochemistry*, 2015, **120**, 3–18.
- 97 S. Irmisch, A. Clavijo McCormick, G. A. Boeckler, A. Schmidt, M. Reichelt, B. Schneider, K. Block, J.-P. Schnitzler, J. Gershenzon, S. B. Unsicker and T. G. Köllner, *Plant Cell*, 2013, **25**, 4737–4754.
- 98 M. Sørensen, E. H. J. Neilson and B. L. Møller, *Mol. Plant*, 2018, **11**, 95–117.
- 99 S. Irmisch, A. Clavijo McCormick, J. Günther, A. Schmidt, G. A. Boeckler, J. Gershenzon, S. B. Unsicker and T. G. Köllner, *Plant J.*, 2014, **80**, 1095–1107.
- 100 J. Martínez-Harms, A.-C. Warskulat, B. Dudek, G. Kunert, S. Lorenz, B. S. Hansson and B. Schneider, *ChemBioChem*, 2018, **19**, 1553–1562.
- 101 E. C. Tatsis, H. Böhm and B. Schneider, *Phytochemistry*, 2013, **92**, 105–112.
- 102 J. C. Verdonk, M. A. Haring, A. J. van Tunen and R. C. Schuurink, *Plant Cell*, 2005, **17**, 1612–1624.
- 103 U. Klahre, A. Gurba, K. Hermann, M. Saxenhofer, E. Bossolini, P. M. Guerin and C. Kuhlemeier, *Curr. Biol.*, 2011, **21**, 730–739.
- 104 B. Spitzer-Rimon, E. Marhevka, O. Barkai, I. Marton, O. Edelbaum, T. Masci, N.-K. Prathapani, E. Shklarman, M. Ovadis and A. Vainstein, *Plant Cell*, 2010, **22**, 1961–1976.
- 105 A. Van Moerkercke, M. A. Haring and R. C. Schuurink, *Plant J.*, 2011, **67**, 917–928.
- 106 B. Spitzer-Rimon, M. Farhi, B. Albo, A. Cna'ani, M. M. Ben Zvi, T. Masci, O. Edelbaum, Y. Yu, E. Shklarman, M. Ovadis and A. Vainstein, *Plant Cell*, 2012, **24**, 5089–5105.
- 107 M. R. Boersma, R. M. Patrick, S. L. Jillings, N. F. M. Shaipulah, P. Sun, M. A. Haring, N. Dudareva, Y. Li and R. C. Schuurink, *Plant J.*, 2022, **109**, 1134–1151.
- 108 A. Cna'ani, B. Spitzer-Rimon, J. Ravid, M. Farhi, T. Masci, J. Aravena-Calvo, M. Ovadis and A. Vainstein, *New Phytol.*, 2015, **208**, 708–714.



- 109 F. Liu, Z. Xiao, L. Yang, Q. Chen, L. Shao, J. Liu and Y. Yu, *New Phytol.*, 2017, **215**, 1490–1502.
- 110 J. Ravid, B. Spitzer-Rimon, Y. Takebayashi, M. Seo, A. Cna'ani, J. Aravena-Calvo, T. Masci, M. Farhi and A. Vainstein, *New Phytol.*, 2017, **215**, 411–422.
- 111 J. H. Lynch, Y. C. Qian, L. Y. Guo, I. Maoz, X. Q. Huang, A. S. Garcia, G. Louie, M. E. Bowman, J. P. Noel, J. A. Morgan and N. Dudareva, *Nat. Chem. Biol.*, 2020, **16**, 850–856.
- 112 M. M. Ben Zvi, E. Shklarman, T. Masci, H. Kalev, T. Debener, S. Shafir, M. Ovadis and A. Vainstein, *New Phytol.*, 2012, **195**, 335–345.
- 113 Z. Yang, Y. Li, F. Gao, W. Jin, S. Li, S. Kimani, S. Yang, T. Bao, X. Gao and L. Wang, *J. Exp. Bot.*, 2020, **71**, 4140–4158.
- 114 G.-J. Hong, X.-Y. Xue, Y.-B. Mao, L.-J. Wang and X.-Y. Chen, *Plant Cell*, 2012, **24**, 2635–2648.
- 115 Z. Yu, G. Zhang, J. A. Teixeira da Silva, C. Zhao and J. Duan, *Plant Sci.*, 2021, **309**, 110952.
- 116 F. Abbas, Y. Ke, Y. Zhou, Y. Yu, M. Waseem, U. Ashraf, C. Wang, X. Wang, X. Li, Y. Yue, R. Yu and Y. Fan, *Front. Plant Sci.*, 2021, **12**, 623742.
- 117 Y. Ke, F. Abbas, Y. Zhou, R. Yu and Y. Fan, *Front. Plant Sci.*, 2021, **12**, 710826.
- 118 F. Abbas, Y. Ke, Y. Zhou, Y. Yu, M. Waseem, U. Ashraf, X. Li, R. Yu and Y. Fan, *Plant Cell Rep.*, 2021, **40**, 1269–1284.
- 119 W. Ding, Q. Ouyang, Y. Li, T. Shi, L. Li, X. Yang, K. Ji, L. Wang and Y. Yue, *Tree Physiol.*, 2019, **40**, 557–572.
- 120 H.-Y. Li, Y.-Z. Yue, W.-J. Ding, G.-W. Chen, L. Li, Y.-L. Li, T.-T. Shi, X.-L. Yang and L.-G. Wang, *Genes*, 2020, **11**, 353.
- 121 Y.-Y. Yang, B. Ma, Y.-Y. Li, M.-Z. Han, J. Wu, X.-F. Zhou, J. Tian, W.-H. Wang, P.-S. Leng and Z.-H. Hu, *Front. Plant Sci.*, 2022, **13**, 1021576.
- 122 K. Yoshida, N. Oyama-Okubo and M. Yamagishi, *Mol. Breed.*, 2018, **38**, 144.
- 123 J. Cai, P. Zu and F. P. Schiestl, *Sci. Rep.*, 2016, **6**, 36966.
- 124 M. Oliva, R. Ovadia, A. Perl, E. Bar, E. Lewinsohn, G. Galili and M. Oren-Shamir, *Plant Biotechnol. J.*, 2015, **13**, 125–136.
- 125 V. Kumar, Y. Elazari, R. Ovadia, E. Bar, A. Nissim-Levi, N. Carmi, E. Lewinsohn and M. Oren-Shamir, *Postharvest Biol. Technol.*, 2021, **181**, 111657.
- 126 F. Abbas, X. Nian, Y. Zhou, Y. Ke, L. Liu, R. Yu and Y. Fan, *Plant Physiol. Biochem.*, 2021, **167**, 619–629.
- 127 R. M. Patrick, X.-Q. Huang, N. Dudareva and Y. Li, *J. Exp. Bot.*, 2021, **72**, 3704–3722.
- 128 S. Shi, S. Zhang, J. Wu, X. Liu and Z. Zhang, *Front. Plant Sci.*, 2022, **13**, 996474.
- 129 J. R. Widhalm, R. Jaini, J. A. Morgan and N. Dudareva, *Trends Plant Sci.*, 2015, **20**, 545–550.
- 130 F. Adebesein, J. R. Widhalm, B. Boachon, F. Lefèvre, B. Pierman, J. H. Lynch, I. Alam, B. Junqueira, R. Benke, S. Ray, J. A. Porter, M. Yanagisawa, H. Y. Wetzstein, J. A. Morgan, M. Boutry, R. C. Schuurink and N. Dudareva, *Science*, 2017, **356**, 1386–1388.
- 131 P. Liao, S. Ray, B. Boachon, J. H. Lynch, A. Deshpande, S. McAdam, J. A. Morgan and N. Dudareva, *Nat. Chem. Biol.*, 2021, **17**, 138–145.
- 132 A. Cna'ani, R. Shavit, J. Ravid, J. Aravena-Calvo, O. Skaliter, T. Masci and A. Vainstein, *Front. Plant Sci.*, 2017, **8**, 1898.
- 133 B. Boachon, J. H. Lynch, S. Ray, J. Yuan, K. M. P. Caldo, R. R. Junker, S. A. Kessler, J. A. Morgan and N. Dudareva, *Nat. Chem. Biol.*, 2019, **15**, 583–588.
- 134 R. Delle-Vedove, B. Schatz and M. Dufay, *Ann. Bot.*, 2017, **120**, 1–20.
- 135 P. J. Zu, W. U. Blanckenhorn and F. P. Schiestl, *New Phytol.*, 2016, **209**, 1208–1219.
- 136 F. P. Schiestl, *New Phytol.*, 2015, **206**, 571–577.
- 137 I. Paul, M. P. Sarkar and P. B. S. Bhadoria, *Chemoecology*, 2022, **32**, 49–68.
- 138 J. M. W. Ryalls, B. Langford, N. J. Mullinger, L. M. Bromfield, E. Nemitz, C. Pfrang and R. D. Girling, *Environ. Pollut.*, 2022, **297**, 118847.
- 139 E. Agathokleous, Z. Feng and J. Peñuelas, *Trends Ecol. Evol.*, 2022, **37**, 939–941.
- 140 X. Y. Yuan, Z. Z. Feng, C. F. Hu, K. Zhang, L. Y. Qu and E. Paoletti, *Environ. Pollut.*, 2021, **291**, 118141.
- 141 A. Saunier and J. D. Blande, *Environ. Pollut.*, 2019, **255**, 113257.
- 142 W. R. Glenney, J. B. Runyon and L. A. Burkle, *New Phytol.*, 2018, **220**, 785–798.
- 143 R. Atkinson, J. Arey, S. M. Aschmann, S. B. Corchnoy and Y. H. Shu, *Int. J. Chem. Kinet.*, 1995, **27**, 941–955.
- 144 J. D. Blande, *Curr. Opin. Environ. Sci. Health*, 2021, **19**, 100228.
- 145 J. D. Blande, J. K. Holopainen and U. Niinemets, *Plant, Cell Environ.*, 2014, **37**, 1892–1904.
- 146 J. D. Fuentes, M. Chamecki, T. Roulston, B. C. Chen and K. R. Pratt, *Atmos. Environ.*, 2016, **141**, 361–374.
- 147 A. Jürgens and M. Bischoff, *Funct. Ecol.*, 2017, **31**, 56–64.
- 148 Q. S. McFrederick, J. D. Fuentes, T. Roulston, J. C. Kathilankal and M. Lerdau, *Oecologia*, 2009, **160**, 411–420.
- 149 G. Farré-Armengol, J. Peñuelas, T. Li, P. Yli-Pirilä, I. Filella, J. Llusà and J. D. Blande, *New Phytol.*, 2016, **209**, 152–160.
- 150 A. Saunier, P. Grof-Tisza and J. D. Blande, *Environ. Pollut.*, 2023, **316**, 120573.
- 151 B. Cook, A. Haverkamp, B. S. Hansson, T. Roulston, M. Lerdau and M. Knaden, *J. Chem. Ecol.*, 2020, **46**, 987–996.
- 152 J. D. Fuentes, T. H. Roulston and J. Zenker, *Environ. Res. Lett.*, 2013, **8**, 014048.
- 153 R. D. Girling, I. Lusebrink, E. Farthing, T. A. Newman and G. M. Poppy, *Sci. Rep.*, 2013, **3**, 2779.
- 154 I. Lusebrink, R. D. Girling, E. Farthing, T. A. Newman, C. W. Jackson and G. M. Poppy, *J. Chem. Ecol.*, 2015, **41**, 904–912.
- 155 F. Démare, L. Gibert, P. Creusot, B. Lapeyre and M. Proffit, *Sci. Total Environ.*, 2022, **827**, 154342.
- 156 S. Dötterl, M. Vater, T. Rupp and A. Held, *J. Chem. Ecol.*, 2016, **42**, 486–489.
- 157 M. Vanderplanck, B. Lapeyre, M. Brondani, M. Opsommer, M. Dufay, M. Hossaert-McKey and M. Proffit, *Antioxidants*, 2021, **10**, 636.



- 158 IPCC, *Climate Change 2021: The Physical Science Basis. Contribution of Working Group I to the Sixth Assessment Report of the Intergovernmental Panel on Climate Change*, ed. V. Masson-Delmotte, P. Zhai, A. Pirani, S. L. Connors, C. Péan, S. Berger, N. Caud, Y. Chen, L. Goldfarb, M. I. Gomis, M. Huang, K. Leitzell, E. Lonnoy, J. B. R. Matthews, T. K. Maycock, T. Waterfield, O. Yelekçi, R. Yu and B. Zhou, Cambridge University Press, Cambridge, United Kingdom, New York, NY, USA, 2021.
- 159 M. Borghi, L. P. de Souza, T. Yoshida and A. R. Fernie, *New Phytol.*, 2019, **224**, 1425–1441.
- 160 M. Gérard, M. Vanderplanck, T. Wood and D. Michez, *Emerging Top. Life Sci.*, 2020, **4**, 77–86.
- 161 M. Barman and A. Mitra, *Environ. Exp. Bot.*, 2021, **181**, 104296.
- 162 A. Cna'ani, J. K. Mühlemann, J. Ravid, T. Masci, A. Klempien, T. T. H. Nguyen, N. Dudareva, E. Pichersky and A. Vainstein, *Plant, Cell Environ.*, 2015, **38**, 1333–1346.
- 163 G. Farré-Armengol, I. Filella, J. Llusia, U. Niinemets and J. Peñuelas, *Global Change Biol.*, 2014, **20**, 3660–3669.
- 164 G. Farré-Armengol, I. Filella, J. Llusia, U. Niinemets and J. Peñuelas, *Funct. Plant Biol.*, 2015, **42**, 851–857.
- 165 Z. H. Hu, H. X. Zhang, P. S. Leng, J. Zhao, W. H. Wang and S. D. Wang, *Acta Physiol. Plant.*, 2013, **35**, 1691–1700.
- 166 S. H. Cheng, X. M. Fu, X. Mei, Y. Zhou, B. Du, N. Watanabe and Z. Y. Yang, *Plant Physiol. Biochem.*, 2016, **107**, 1–8.
- 167 J. X. Fu, D. Hou, C. Zhang, Z. Y. Bao, H. B. Zhao and S. Q. Hu, *Molecules*, 2017, **22**, 430.
- 168 D. R. Campbell, P. Sosenski and R. A. Raguso, *Ann. Bot.*, 2019, **123**, 601–610.
- 169 L. A. Burkle and J. B. Runyon, *Global Change Biol.*, 2016, **22**, 1644–1654.
- 170 C. C. Rering, J. G. Franco, K. M. Yeater and R. E. Mallinger, *Ecosphere*, 2020, **11**, e03254.
- 171 I. N. A. Salman, A. Cna'ani, V. Tzin and M. Seifan, *Entomol. Exp. Appl.*, 2022, **170**, 666–680.
- 172 C. C. Jaworski, B. Geslin, M. Zakardjian, C. Lecareux, P. Caillault, G. Nève, J. Y. Meunier, S. Dupouyet, A. C. T. Sweeney, O. T. Lewis, L. V. Dicks and C. Fernandez, *J. Ecol.*, 2022, **110**, 2628–2648.
- 173 R. J. Höfer, M. Ayasse and J. Kuppler, *Front. Plant Sci.*, 2021, **11**, 564802.
- 174 V. J. Luizzi, M. Friberg and H. Petré, *Funct. Ecol.*, 2021, **35**, 1655–1665.
- 175 M. Friberg, M. T. Waters and J. N. Thompson, *Ann. Bot.*, 2017, **120**, 471–478.
- 176 C. J. Majetic, A. M. Fettes, O. M. Beck, E. F. Stachnik and K. M. Beam, *Flora*, 2017, **232**, 183–193.
- 177 T. Dorey and F. P. Schiestl, *Evolution*, 2022, **76**, 2930–2944.
- 178 K. M. Becklin, G. Gamez, B. Uelk, R. A. Raguso and C. Galen, *Am. J. Bot.*, 2011, **98**, 1299–1308.
- 179 N. Wei, R. L. Whyte, T.-L. Ashman and M. A. Jamieson, *Hortic. Res.*, 2022, **9**, uh0005.
- 180 L. Shapiro, C. M. De Moraes, A. G. Stephenson and M. C. Mescher, *Ecol. Lett.*, 2012, **15**, 1430–1438.
- 181 A. Cellini, V. Giacomuzzi, I. Donati, B. Farneti, M. T. Rodriguez-Estrada, S. Savioli, S. Angeli and F. Spinelli, *ISME J.*, 2019, **13**, 847–859.
- 182 R. R. Junker and D. Tholl, *J. Chem. Ecol.*, 2013, **39**, 810–825.
- 183 P. Gaube, R. R. Junker and A. Keller, *Basic Appl. Ecol.*, 2021, **50**, 1–15.
- 184 R. R. Junker and A. Keller, *FEMS Microbiol. Ecol.*, 2015, **91**, fiv097.
- 185 A. Crowley-Gall, C. C. Rering, A. B. Rudolph, R. L. Vannette and J. J. Beck, *Curr. Opin. Insect. Sci.*, 2021, **44**, 23–34.
- 186 E. D. Fenner, T. Scapini, M. da Costa Diniz, A. Giehl, H. Treichel, S. Álvarez-Pérez and S. L. Alves Jr., *J. Fungi*, 2022, **8**, 984.
- 187 J. S. Francis, A. R. Tatarko, S. K. Richman, A. D. Vaudo and A. S. Leonard, *Curr. Opin. Insect. Sci.*, 2021, **44**, 16–22.
- 188 J. Klaps, B. Lievens and S. Álvarez-Pérez, *Fungal Biol. Biotechnol.*, 2020, **7**, 1.
- 189 R. L. Vannette, *Annu. Rev. Ecol. Evol. Syst.*, 2020, **51**, 363–386.
- 190 A. M. Golonka, B. Obi Johnson, J. Freeman and D. W. Hinson, *East. Biol.*, 2014, **3**, 1–26.
- 191 C. C. Rering, J. J. Beck, G. W. Hall, M. M. McCartney and R. L. Vannette, *New Phytol.*, 2018, **220**, 750–759.
- 192 C. C. Rering, A. B. Rudolph and J. J. Beck, *Environ. Microbiol.*, 2021, **23**, 4141–4150.
- 193 R. N. Schaeffer, C. C. Rering, I. Maalouf, J. J. Beck and R. L. Vannette, *Biol. Lett.*, 2019, **15**, 20190132.
- 194 I. S. Sobhy, D. Baets, T. Goelen, B. Herrera-Malaver, L. Bosmans, W. Van den Ende, K. J. Verstrepen, F. Wäckers, H. Jacquemyn and B. Lievens, *Front. Plant Sci.*, 2018, **9**, 1009.
- 195 A. Colda, S. Bossaert, C. Verreth, B. Vanhoutte, O. Honnay, W. Keulemans and B. Lievens, *PLoS One*, 2021, **16**, e0250203.
- 196 R. L. Vannette, M. P. L. Gauthier and T. Fukami, *Proc. R. Soc. B: Biol. Sci.*, 2013, **280**, 20122601.
- 197 A. Cusumano, P. Bella, E. Peri, M. Rostás, S. Guarino, B. Lievens and S. Colazza, *Microb. Ecol.*, 2023, **86**, 364–376.
- 198 C. Helletsgruber, S. Dötterl, U. Ruprecht and R. R. Junker, *J. Chem. Ecol.*, 2017, **43**, 1073–1077.
- 199 J. Peñuelas, G. Farré-Armengol, J. Llusia, A. Gargallo-Garriga, L. Rico, J. Sardans, J. Terradas and I. Filella, *Sci. Rep.*, 2014, **4**, 6727.
- 200 L. Huang, Y. Liu, L. Dou, S. Pan, Z. Li, J. Zhang and J. Li, *PeerJ*, 2022, **10**, e14107.
- 201 A. Kessler and A. Chautá, *Emerging Top. Life Sci.*, 2020, **4**, 33–43.
- 202 D. Lucas-Barbosa, *Trends Plant Sci.*, 2016, **21**, 125–133.
- 203 D. Lucas-Barbosa, J. J. A. van Loon and M. Dicke, *Phytochemistry*, 2011, **72**, 1647–1654.
- 204 Q. Rusman, D. Lucas-Barbosa, E. H. Poelman and M. Dicke, *Trends Plant Sci.*, 2019, **24**, 725–740.
- 205 P. Tunes, S. Dötterl and E. Guimarães, *Front. Plant Sci.*, 2022, **13**, 813418.
- 206 C. E. Reisenman, J. A. Riffell, K. Duffy, A. Pesque, D. Mikles and B. Goodwin, *J. Chem. Ecol.*, 2013, **39**, 76–89.



- 207 M. Pareja, E. Qvarfordt, B. Webster, P. Mayon, J. Pickett, M. Birkett and R. Glinwood, *PLoS One*, 2012, **7**, e31971.
- 208 Q. Rusman, E. H. Poelman, F. Nowrin, G. Polder and D. Lucas-Barbosa, *Plant, Cell Environ.*, 2019, **42**, 1882–1896.
- 209 F. P. Schiestl, H. Kirk, L. Bigler, S. Cozzolino and G. A. Desurmont, *New Phytol.*, 2014, **203**, 257–266.
- 210 M. Bruinsma, D. Lucas-Barbosa, C. J. M. ten Broeke, N. M. van Dam, T. A. van Beek, M. Dicke and J. J. A. van Loon, *J. Chem. Ecol.*, 2014, **40**, 39–49.
- 211 M. Hoffmeister, N. Wittkopper and R. R. Junker, *Oikos*, 2016, **125**, 1241–1249.
- 212 A. Kessler, R. Halitschke and K. Poveda, *Ecology*, 2011, **92**, 1769–1780.
- 213 D. Kessler, C. Diezel and I. T. Baldwin, *Curr. Biol.*, 2010, **20**, 237–242.
- 214 S. Cozzolino, S. Fineschi, M. Litto, G. Scopece, J. Trunschke and F. P. Schiestl, *J. Chem. Ecol.*, 2015, **41**, 622–630.
- 215 S. E. Ramos and F. P. Schiestl, *Science*, 2019, **364**, 193–196.
- 216 S. E. Ramos and F. P. Schiestl, *Front. Ecol. Evol.*, 2020, **8**, 30.
- 217 J. Ollerton, *Annu. Rev. Ecol. Evol. Syst.*, 2017, **48**, 353–376.
- 218 D. Schneider, M. Wink, F. Sporer and P. Lounibos, *Naturwissenschaften*, 2002, **89**, 281–294.
- 219 G. Gottsberger, *Plant Divers. Evol.*, 2016, **131**, 263–362.
- 220 M. R. Moore, R. D. Cave and M. A. Branham, *Zookeys*, 2018, **745**, 1–99.
- 221 M. R. Moore and M. L. Jameson, *J. Insect Sci.*, 2013, **13**, 100.
- 222 G. Gottsberger, I. Silberbauer-Gottsberger, R. S. Seymour and S. Dötterl, *Flora*, 2012, **207**, 107–118.
- 223 S. Dötterl, A. David, W. Boland, I. Silberbauer-Gottsberger and G. Gottsberger, *J. Chem. Ecol.*, 2012, **38**, 1539–1543.
- 224 A. C. D. Maia, M. Gibernau, S. Dötterl, D. M. D. F. Navarro, K. Seifert, T. Müller and C. Schlindwein, *Phytochemistry*, 2013, **93**, 71–78.
- 225 J. Pereira, C. Schlindwein, Y. Antonini, A. C. D. Maia, S. Dötterl, C. Martins, D. M. D. F. Navarro and R. Oliveira, *Biol. J. Linn. Soc.*, 2014, **111**, 679–691.
- 226 A. C. D. Maia, C. Grimm, M. Schubert, F. Etl, E. G. Gonçalves, D. M. D. F. Navarro, S. Schulz and S. Dötterl, *J. Chem. Ecol.*, 2019, **45**, 214–215.
- 227 P. Stamm, F. Etl, A. C. D. Maia, S. Dötterl and S. Schulz, *J. Org. Chem.*, 2021, **86**, 5245–5254.
- 228 T. N. Suinyuy, J. S. Donaldson and S. D. Johnson, *Proc. R. Soc. B: Biol. Sci.*, 2015, **282**, 20152053.
- 229 T. N. Suinyuy and S. D. Johnson, *Bot. J. Linn. Soc.*, 2021, **195**, 233–248.
- 230 S. Salzman, A. Dahake, W. Kandalaft, W. A. Valencia-Montoya, M. Calonje, C. D. Specht and R. A. Raguso, *Curr. Biol.*, 2023, **33**, 1654–1664.
- 231 S. L. Steenhuisen, A. Jürgens and S. D. Johnson, *J. Chem. Ecol.*, 2013, **39**, 438–446.
- 232 L. Blažytė-Čereškienė, V. Apšegaitė and V. Būda, *Arthropod-Plant Interact.*, 2019, **13**, 735–743.
- 233 T. D. J. Sayers, M. J. Steinbauer, K. Farnier and R. E. Miller, *Bot. J. Linn. Soc.*, 2020, **193**, 375–401.
- 234 B. S. Brodie, A. Renyard, R. Gries, H. M. Zhai, S. Ogilvie, J. Avery and G. Gries, *Arthropod-Plant Interact.*, 2018, **12**, 591–599.
- 235 J. Vuts, C. M. Woodcock, J. C. Caulfield, S. J. Powers, J. A. Pickett and M. A. Birkett, *Pest Manage. Sci.*, 2018, **74**, 2069–2075.
- 236 T. J. A. Bruce, J. L. Martin, L. E. Smart and J. A. Pickett, *Pest Manage. Sci.*, 2011, **67**, 1303–1308.
- 237 C. L. Xiu, B. Xu, H. S. Pan, W. Zhang, Y. Z. Yang and Y. H. Lu, *J. Integr. Agric.*, 2019, **18**, 873–883.
- 238 A. C. D. Maia, L. K. Reis, D. M. D. F. Navarro, F. Aristone, C. A. Colombo, J. Carreño-Barrera, L. A. Núñez-Avellaneda and G. K. N. Santos, *J. Appl. Entomol.*, 2020, **144**, 33–40.
- 239 A. C. D. Maia, D. M. D. F. Navarro, L. A. Núñez-Avellaneda, J. Carreño-Barrera, L. Iannuzzi, J. Cardona-Duque and W. A. G. Nantes, *Sci. Nat.*, 2021, **108**, 21.
- 240 A. P. Favaris, A. C. Tuler, W. D. Silva, S. R. Rodrigues, W. S. Leal and J. M. S. Bento, *PLoS One*, 2020, **15**, e0235028.
- 241 A. C. D. Maia, G. K. N. Santos, E. G. Gonçalves, D. M. D. F. Navarro and L. A. Nuñez-Avellaneda, *Pest Manage. Sci.*, 2018, **74**, 2053–2058.
- 242 A. Karmakar, A. Mukherjee and A. Barik, *Entomol. Exp. Appl.*, 2016, **158**, 133–141.
- 243 N. Sarkar, S. Mitra and A. Barik, *Int. J. Pest Manage.*, 2017, **63**, 138–145.
- 244 D. Kessler, C. Diezel, D. G. Clark, T. A. Colquhoun and I. T. Baldwin, *Ecol. Lett.*, 2013, **16**, 299–306.
- 245 A. de Fouchier, W. B. Walker, N. Montagné, C. Steiner, M. Binyameen, F. Schlyter, T. Chertemps, A. Maria, M. C. François, C. Monsempes, P. Anderson, B. S. Hansson, M. C. Larsson and E. Jacquín-Joly, *Nat. Commun.*, 2017, **8**, 5709.
- 246 M. B. Guo, L. X. Du, Q. Y. Chen, Y. L. Feng, J. Zhang, X. X. Zhang, K. Tian, S. Cao, T. Y. Huang, E. Jacquín-Joly, G. R. Wang and Y. Liu, *Mol. Biol. Evol.*, 2021, **38**, 1413–1427.
- 247 J. A. Riffell, H. Lei, L. Abrell and J. G. Hildebrand, *Science*, 2013, **339**, 200–204.
- 248 A. Haverkamp, F. Yon, I. W. Keeseey, C. Missbach, C. Koenig, B. S. Hansson, I. T. Baldwin, M. Knaden and D. Kessler, *eLife*, 2016, **5**, e15039.
- 249 W. W. Zhou, A. Kuegler, E. McGale, A. Haverkamp, M. Knaden, H. Guo, F. Beran, F. Yon, R. Li, N. Lackus, T. G. Köllner, J. Bing, M. C. Schuman, B. S. Hansson, D. Kessler, I. T. Baldwin and S. Q. Xu, *Curr. Biol.*, 2017, **27**, 1336–1341.
- 250 E. Adam, B. S. Hansson and M. Knaden, *J. Exp. Biol.*, 2021, **224**, jeb242780.
- 251 M. Bischoff, R. A. Raguso, A. Jürgens and D. R. Campbell, *Evolution*, 2015, **69**, 1–13.
- 252 F. Balao, J. Herrera, S. Talavera and S. Dötterl, *Phytochemistry*, 2011, **72**, 601–609.
- 253 M. von Arx, J. Goyret, G. Davidowitz and R. A. Raguso, *Proc. Natl. Acad. Sci. U. S. A.*, 2012, **109**, 9471–9476.
- 254 A. Dahake, P. Jain, C. C. Vogt, W. Kandalaft, A. D. Stroock and R. A. Raguso, *Nat. Commun.*, 2022, **13**, 7773.
- 255 A. Shuttleworth and S. D. Johnson, *Entomol. Exp. Appl.*, 2020, **168**, 191–197.
- 256 C. Eby, M. G. T. Gardiner, R. Gries, G. J. R. Judd, G. Khaskin and G. Gries, *Entomol. Exp. Appl.*, 2013, **147**, 82–92.



- 257 A. Jürgens, U. Glück, G. Aas and S. Dötterl, *Bot. J. Linn. Soc.*, 2014, **175**, 624–640.
- 258 B. P. Molnár, Z. Kárpáti, A. Nagy, I. Szarukán, J. Csabai, S. Koczor and M. Tóth, *J. Chem. Ecol.*, 2019, **45**, 657–666.
- 259 G. Thöming and G. K. Knudsen, *Phytochemistry*, 2014, **100**, 66–75.
- 260 M. Friberg, C. Schwind, L. C. Roark, R. A. Raguso and J. N. Thompson, *J. Chem. Ecol.*, 2014, **40**, 955–965.
- 261 F. P. Schiestl, E. A. Wallin, J. J. Beck, M. Friberg and J. N. Thompson, *Arthropod-Plant Interact.*, 2021, **15**, 209–221.
- 262 G. P. Svensson, T. Okamoto, A. Kawakita, R. Goto and M. Kato, *New Phytol.*, 2010, **186**, 995–1004.
- 263 D. H. Huang, J. Mamut, X. F. Yang, F. C. Shi and H. H. Li, *Diversity*, 2022, **14**, 266.
- 264 T. Okamoto, in *Obligate Pollination Mutualism. Ecological Research Monographs*, ed. M. Kato and A. Kawakita, Springer, Tokyo, 2017.
- 265 T. Okamoto, A. Kawakita, R. Goto, G. P. Svensson and M. Kato, *Proc. R. Soc. B*, 2013, **280**, 20132280.
- 266 T. Okamoto, G. P. Svensson, R. Goto, A. Kawakita and M. Kato, *Plant Species Biol.*, 2022, **37**, 197–208.
- 267 G. P. Svensson, O. Pellmyr and R. A. Raguso, *Oikos*, 2011, **120**, 1577–1583.
- 268 M. Kinoshita, F. J. Stewart and H. Omura, *BioEssays*, 2017, **39**, 1600086.
- 269 S. Lehner, S. Schulz and S. Dötterl, *Front. Plant Sci.*, 2022, **13**, 994851.
- 270 S. Andersson, *Chemoecology*, 2003, **13**, 13–20.
- 271 S. Andersson, L. A. Nilsson, I. Groth and G. Bergström, *Bot. J. Linn. Soc.*, 2002, **140**, 129–153.
- 272 A. M. El-Sayed, A. Sporle, K. Colhoun, J. Furlong, R. White and D. M. Suckling, *Chemoecology*, 2018, **28**, 39–49.
- 273 R. A. Raguso, *Appl. Entomol. Zool.*, 2020, **55**, 1–7.
- 274 T. S. Woodcock, B. M. H. Larson, P. G. Kevan, D. W. Inouye and K. Lunau, *J. Pollinat. Ecol.*, 2014, **12**, 63–94.
- 275 C. Primante and S. Dötterl, *J. Chem. Ecol.*, 2010, **36**, 1207–1210.
- 276 L. Dormont, M. Mulatier, D. Carrasco and A. Cohuet, *J. Chem. Ecol.*, 2021, **47**, 351–393.
- 277 R. Ignell and S. R. Hill, *Curr. Opin. Insect. Sci.*, 2020, **40**, 6–10.
- 278 E. K. Lutz, C. Lahondère, C. Vinauger and J. A. Riffell, *Curr. Opin. Insect. Sci.*, 2017, **20**, 75–83.
- 279 A. Mafra-Neto and T. Dekker, *Curr. Opin. Insect. Sci.*, 2019, **34**, 105–111.
- 280 J. M. Muema, J. L. Bargul, S. N. Njeru, J. O. Onyango and S. S. Imbahale, *Parasites Vectors*, 2017, **10**, 184.
- 281 V. O. Nyasembe and B. Torto, *Phytochem. Lett.*, 2014, **8**, 196–201.
- 282 M. Wooding, Y. Naudé, E. Rohwer and M. Bouwer, *Parasites Vectors*, 2020, **13**, 80.
- 283 P. E. Otienoburu, B. Ebrahimi, P. L. Phelan and W. A. Foster, *J. Chem. Ecol.*, 2012, **38**, 873–881.
- 284 S. von Oppen, H. Masuh, S. Licastro, E. Zerba and P. Gonzalez-Audino, *Entomol. Exp. Appl.*, 2015, **155**, 184–192.
- 285 D. A. H. Peach, R. Gries, H. M. Zhai, N. Young and G. Gries, *Sci. Rep.*, 2019, **9**, 3908.
- 286 T. Okamoto, Y. Okuyama, R. Goto, M. Tokoro and M. Kato, *J. Evol. Biol.*, 2015, **28**, 590–600.
- 287 S. Ibanez, S. Dötterl, M. C. Anstett, S. Baudino, J. C. Caissard, C. Gallet and L. Despres, *New Phytol.*, 2010, **188**, 451–463.
- 288 B. Song, G. Chen, J. Stöcklin, D. L. Peng, Y. Niu, Z. M. Li and H. Sun, *New Phytol.*, 2014, **203**, 1109–1118.
- 289 A. Jürgens, S. L. Wee, A. Shuttleworth and S. D. Johnson, *Ecol. Lett.*, 2013, **16**, 1157–1167.
- 290 G. Chen, X. K. Ma, A. Jürgens, J. Lu, E. X. Liu, W. B. Sun and X. H. Cai, *J. Chem. Ecol.*, 2015, **41**, 808–815.
- 291 M. Moré, A. A. Cocucci and R. A. Raguso, *Int. J. Plant Sci.*, 2013, **174**, 863–876.
- 292 M. Moré, P. Mulieri, M. Battán-Horenstein, A. A. Cocucci and R. A. Raguso, *Arthropod-Plant Interact.*, 2019, **13**, 375–386.
- 293 A. Shuttleworth and S. D. Johnson, *Proc. R. Soc. B: Biol. Sci.*, 2010, **277**, 2811–2819.
- 294 S. L. Wee, S. B. Tan and A. Jürgens, *Phytochemistry*, 2018, **153**, 120–128.
- 295 P. Zito, M. Sajeva, A. Raspi and S. Dötterl, *Chemoecology*, 2014, **24**, 261–267.
- 296 A. Shuttleworth, S. D. Johnson and A. Jürgens, *Flora*, 2017, **232**, 92–103.
- 297 P. Zito, S. Dötterl and M. Sajeva, *J. Chem. Ecol.*, 2015, **41**, 340–349.
- 298 S. D. Johnson, J. Sivechurran, S. Doarsamy and A. Shuttleworth, *New Phytol.*, 2020, **228**, 1662–1673.
- 299 G. Chen, A. Jürgens, L. D. Shao, Y. Liu, W. B. Sun and C. F. Xia, *J. Chem. Ecol.*, 2015, **41**, 244–252.
- 300 P. Zito, S. Guarino, E. Peri, M. Sajeva and S. Colazza, *Arthropod-Plant Interact.*, 2013, **7**, 485–489.
- 301 K. R. Goodrich and A. Jürgens, *New Phytol.*, 2018, **217**, 74–81.
- 302 A. Heiduk, H. Kong, I. Brake, M. von Tschirnhaus, T. Tolasch, A. G. Tröger, E. Wittenberg, W. Francke, U. Meve and S. Dötterl, *Front. Ecol. Evol.*, 2015, **3**, 66.
- 303 B. Oelschlägel, M. Nuss, M. von Tschirnhaus, C. Pätzold, C. Neinhuis, S. Dötterl and S. Wanke, *New Phytol.*, 2015, **206**, 342–351.
- 304 S. Dötterl and N. Vereecken, *Can. J. Zool.*, 2010, **88**, 668–697.
- 305 C. D. Michener, *The bees of the world*, The John Hopkins University Press, Baltimore, 2nd edn, 2007.
- 306 R. Kubo and M. Ono, *Entomol. Sci.*, 2014, **17**, 432–434.
- 307 G. Rabeschi, P. J. Bergamo and C. E. P. Nunes, *J. Chem. Ecol.*, 2021, **47**, 444–454.
- 308 A. C. Knauer and F. P. Schiestl, *Ecol. Lett.*, 2015, **18**, 135–143.
- 309 C. Rodríguez-Saona, L. Parra, A. Quiroz and R. Isaacs, *Ann. Bot.*, 2011, **107**, 1377–1390.
- 310 F. Mas, A. Harper, R. Horner, T. Welsh, P. Jaksons and D. M. Suckling, *J. Sci. Food Agric.*, 2018, **98**, 4445–4453.
- 311 M. Rachtersberger, G. D. Cordeiro, I. Schäffler and S. Dötterl, *J. Agric. Food Chem.*, 2019, **67**, 13221–13227.
- 312 K. Lukas, S. Dötterl, M. Ayasse and H. Burger, *Front. Ecol. Evol.*, 2020, **8**, 1–14.



- 313 A. Morse, P. Kevan, L. Shipp, S. Khosla and B. McGarvey, *Environ. Entomol.*, 2012, **41**, 855–864.
- 314 G. D. Cordeiro, M. Pinheiro, S. Dötterl and I. Alves-dos-Santos, *Plant Biol.*, 2017, **19**, 132–139.
- 315 C. A. Martínez-Martínez, G. D. Cordeiro, H. O. J. Martins, R. O. A. C. Kobal, P. Milet-Pinheiro, M. A. Stanton, E. L. Franco, C. Krug, S. Mateus, C. Schlindwein, S. Dötterl and I. Alves-dos-Santos, *Front. Ecol. Evol.*, 2021, **9**, 676743.
- 316 C. Krug, G. D. Cordeiro, I. Schäffler, C. I. Silva, R. Oliveira, C. Schlindwein, S. Dötterl and I. Alves-dos-Santos, *Front. Plant Sci.*, 2018, **9**, 1072.
- 317 C. Krug, M. V. B. Garcia and F. B. Gomes, *Apidologie*, 2015, **46**, 164–166.
- 318 P. Milet-Pinheiro, M. Ayasse, H. E. M. Dobson, C. Schlindwein, W. Francke and S. Dötterl, *J. Chem. Ecol.*, 2013, **39**, 1347–1360.
- 319 H. Burger, N. Joos and M. Ayasse, *Front. Ecol. Evol.*, 2021, **9**, 682960.
- 320 P. Milet-Pinheiro, K. Herz, S. Dötterl and M. Ayasse, *BMC Ecol.*, 2016, **16**, 20.
- 321 K. Brandt, S. Dötterl, W. Francke, M. Ayasse and P. Milet-Pinheiro, *J. Chem. Ecol.*, 2017, **43**, 4–12.
- 322 S. Dötterl, U. Füssel, A. Jürgens and G. Aas, *J. Chem. Ecol.*, 2005, **31**, 2993–2998.
- 323 H. Burger, M. Ayasse, S. Dötterl, S. Kreissl and C. G. Galizia, *J. Comp. Physiol., A*, 2013, **199**, 751–761.
- 324 I. Schäffler, F. Balao and S. Dötterl, *Ann. Bot.*, 2012, **110**, 125–138.
- 325 M. Castañeda-Zárate, S. D. Johnson and T. van der Niet, *Curr. Biol.*, 2021, **31**, 238–246.
- 326 H. Burger, M. Ayasse, C. M. Häberlein, S. Schulz and S. Dötterl, *S. Afr. J. Bot.*, 2010, **76**, 788–795.
- 327 H. Burger, S. Dötterl, C. Häberlein, S. Schulz and M. Ayasse, *Oecologia*, 2012, **168**, 727–736.
- 328 A. T. Carvalho, S. Dötterl and C. Schlindwein, *J. Chem. Ecol.*, 2014, **40**, 1126–1134.
- 329 T. Eltz, D. W. Roubik and M. W. Whitten, *Physiol. Entomol.*, 2003, **28**, 251–260.
- 330 S. Vogel, *Österr. Bot. Z.*, 1966, **113**, 302–361.
- 331 S. Ramírez, R. L. Dressler and M. Ospina, *Biota Colomb.*, 2002, **3**, 7–118.
- 332 J. Brodmann, D. Emer and M. Ayasse, *Plant Biol.*, 2012, **14**, 500–505.
- 333 C. C. L. Soler, M. Proffit, J.-M. Bessière, M. Hossaert-McKey and B. Schatz, *Ecol. Lett.*, 2012, **15**, 978–985.
- 334 D. Gu, D.-R. Yang, P. Yang, Y.-Q. Peng and Z.-J. Wang, *Chem. Ecol.*, 2016, **32**, 733–741.
- 335 M. C. Foti, M. Rostás, E. Peri, K. C. Park, T. Slimani, S. D. Wratten and S. Colazza, *J. Pest Sci.*, 2017, **90**, 299–310.
- 336 C. de Vega, C. M. Herrera and S. Dötterl, *Perspect. Plant Ecol. Evol. Syst.*, 2014, **16**, 32–42.
- 337 T. Yang, G. Stoop, M. Thoen, G. Wieggers and M. A. Jongsma, *Plant Biotechnol. J.*, 2013, **11**, 875–882.
- 338 Z. S. Abdullah, K. J. Ficken, B. P. J. Greenfield and T. M. Butt, *J. Chem. Ecol.*, 2014, **40**, 534–540.
- 339 F. Etl, A. Berger, A. Weber, J. Schönenberger and S. Dötterl, *J. Chem. Ecol.*, 2016, **42**, 300–304.
- 340 S. D. Johnson, P. M. Burgoyne, L. D. Harder and S. Dötterl, *Proc. R. Soc. B*, 2011, **278**, 2303–2310.
- 341 P. Wester, S. D. Johnson and A. Pauw, *New Phytol.*, 2019, **222**, 1624–1637.
- 342 S. D. Johnson and K. Govender, *Philos. Trans. R. Soc., B*, 2022, **377**, 20210167.
- 343 A. C. Knauer, M. Bakhtiari and F. P. Schiestl, *Nat. Commun.*, 2018, **9**, 1367.
- 344 A. Hammerbacher, T. A. Coutinho and J. Gershenzon, *Plant, Cell Environ.*, 2019, **42**, 2827–2843.
- 345 R. R. Junker and N. Blüthgen, *Ann. Bot.*, 2010, **105**, 777–782.
- 346 J. Li, H. Hu, J. Mao, L. Yu, G. Stoop, M. Wang, R. Mumm, N. C. A. de Ruijter, M. Dicke, M. A. Jongsma and C. Wang, *New Phytol.*, 2019, **223**, 1607–1620.
- 347 R. R. Junker, J. Gershenzon and S. B. Unsicker, *J. Chem. Ecol.*, 2011, **37**, 1323–1331.
- 348 R. R. Junker and N. Blüthgen, *Evol. Ecol. Res.*, 2008, **10**, 295–308.
- 349 J. G. Patrick, T. Shepherd, W. Hoppitt, N. S. Plowman and W. Pat, *Arthropod-Plant Interact.*, 2017, **11**, 623–627.
- 350 C. Galen, R. Kaczorowski, S. L. Todd, J. Geib and R. A. Raguso, *Am. Nat.*, 2011, **177**, 258–272.
- 351 D. Kessler, J. Bing, A. Haverkamp and I. T. Baldwin, *Funct. Ecol.*, 2019, **33**, 1223–1232.
- 352 R. R. Junker, I. M. M. Heidinger and N. Blüthgen, *J. Orthoptera Res.*, 2010, **19**, 69–74.
- 353 M. Huang, A. M. Sanchez-Moreiras, C. Abel, R. Sohrabi, S. Lee, J. Gershenzon and D. Tholl, *New Phytol.*, 2012, **193**, 997–1008.
- 354 R. C. F. Burdon, R. R. Junker, D. G. Scofield and A. L. Parachnowitsch, *Chemoecology*, 2018, **28**, 11–19.
- 355 R. R. Junker, C. Loewel, R. Gross, S. Dötterl, A. Keller and N. Blüthgen, *Plant Biol.*, 2011, **13**, 918–924.
- 356 E. A. Dannon, M. Tamò, A. Van Huis and M. Dicke, *J. Chem. Ecol.*, 2010, **36**, 1083–1091.
- 357 M. Ayasse, F. P. Schiestl, H. F. Paulus, F. Ibarra and W. Francke, *Proc. R. Soc. B*, 2003, **270**, 517–522.
- 358 S. Dötterl, D. Burkhardt, B. Weißbecker, A. Jürgens, S. Schütz and A. Mosandl, *J. Chromatogr. A*, 2006, **1113**, 231–238.
- 359 R. A. Raguso, *Curr. Opin. Plant Biol.*, 2016, **32**, 31–36.
- 360 C. E. Reisenman, J. A. Riffell, E. A. Bernays and J. G. Hildebrand, *Proc. R. Soc. B*, 2010, **277**, 2371–2379.
- 361 T. Eltz, E. Hedenström, J. Bång, E. A. Wallin and J. Andersson, *J. Chem. Ecol.*, 2010, **36**, 1322–1326.
- 362 M. W. Whitten, H. G. Hills and N. H. Williams, *Phytochemistry*, 1988, **27**, 2759–2760.
- 363 D. L. P. Schorkopf, L. Mitko and T. Eltz, *J. Chem. Ecol.*, 2011, **37**, 953–960.
- 364 K. Brandt, S. Dötterl, R. Fuchs, D. M. A. F. Navarro, I. C. S. Machado, D. Dobler, O. Reiser, M. Ayasse and P. Milet-Pinheiro, *J. Chem. Ecol.*, 2019, **45**, 464–473.
- 365 F. Etl, W. Francke, J. Schönenberger and S. Dötterl, *J. Chem. Ecol.*, 2022, **48**, 263–269.

