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Bacterial volatile compound-based tools for crop management and quality

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Trends in Plant Science

Bacterial volatile compound-based tools for crop management and quality --Manuscript Draft--

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Abstract:	Bacteria produce a huge diversity of metabolites, many of which mediate ecological relations. Among these, volatile compounds allow broad-range effects at low doses and may therefore be exploited for applications in plant defence and agricultural production. Such applications are still in their early development. Here we review the latest technologies involving the use of bacterial volatile compounds for phytosanitary inspection, biological control, plant growth promotion, and crop quality. We highlight a variety of effects with a potential applicative interest, based on either live biocontrol and/or biostimulant agents, or the isolated metabolites responsible for the interaction with hosts or competitors. Future agricultural technologies may benefit from the clarification of bacterial interactions with the environment, and the development of new analytical tools.		

Highlights

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• Plant-associated bacteria interact with their environment through exchange of chemicals, including volatile compounds. Innovative agricultural technologies may exploit the inherent advantages of bacterial airborne signals, including diffusibility, independence from water availability and physical connection, and absence of pesticide residuals.

• Volatile compounds resulting from plant-pathogens interactions allow non-destructive disease diagnosis on bulk samples of asymptomatic plant material.

• Volatile compounds, expressing a direct biocidal activity, interfering with signalling, or stimulating plant host defences, contribute to biological control of pests and pathogens.

• Bacterial volatile compounds modulate plant hormones enhancing plant growth, stress tolerance, crop quality, aroma and nutraceutical characteristics, and reduce post-harvest losses.

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1 Bacterial volatile compound-based tools for crop management and quality

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9 Keywords

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12

13 Abstract

Bacteria produce a huge diversity of metabolites, many of which mediate ecological relations. 14 Among these, volatile compounds cause broad-range effects at low doses and may therefore be 15 16 exploited for plant defence strategies and agricultural production, but such applications are still in 17 their early development. Here we review the latest technologies involving the use of bacterial 18 volatile compounds for phytosanitary inspection, biological control, plant growth promotion, and crop quality. We highlight a variety of effects with a potential applicative interest, based on either 19 live biocontrol and/or biostimulant agents, or the isolated metabolites responsible for the 20 interaction with hosts or competitors. Future agricultural technologies may benefit from the 21 22 development of new analytical tools to understand bacterial interactions with the environment.

23

24 Bacterial volatile compounds in plant ecological interactions

25 Bacterial metabolic products characterised by low vapour pressure, high lipophylicity and a 26 molecular weight below 300 Da are likely to be released as Volatile Organic Compounds (VOCs, see glossary). Considering the enormous metabolic diversity of bacteria, such compounds may derive 27 from a large variety of chemical pathways, and are generally emitted as complex mixtures [1]. The 28 composition of the bacterial **volatilome** (see glossary) is highly influenced by the growth conditions 29 [2-5], including soil chemistry and structure, pH, availability of water and oxygen, presence of plant 30 exudates or other organic compounds, and light irradiation. In horticultural crop management, 31 32 these parameters may be modified by agricultural (tillage, cover cropping, fertilisation, watering,

and plant protection) or post-harvest (refrigeration, atmosphere control, and ethylene modulation)
 practices, thus affecting BVC emissions as well [6].

35 Along with the characterisation of a growing number of bacterial volatile compounds (BVCs, see glossary) [3,4,7], their roles in intra- or inter-specific signalling or competition are being discovered 36 37 (Figure 1). It should also be noted that the most commonly adopted analytical techniques in studies concerning bacterial volatile compounds are unable to detect molecules with a low molecular 38 weight, such as CO₂, ethylene, nitrogen oxides, ethanol and H₂S (Box 1). Thus, part of the biological 39 effects mediated by bacterial airborne signals may still be eluding the researchers' efforts. In 40 41 comparison to water-soluble compounds, inherent advantages of VOCs in ecological interactions 42 reside in their high diffusibility, enabling both above- and belowground action, the ability to diffuse 43 through lipophilic barriers (such as cell membranes and plant cuticle), and the independence from 44 water and physical connection among the VOC-producing organism and the signal recipient.

Technological applications based on the release or the exchange of VOCs are most likely to succeed, when they take advantage of these characteristics. Volatility is possibly one of the key points, as it allows a relative uniformity of the gas phase even in cases of poor accessibility of the target (for instance, in the soil, in stored bulk samples, or in internal plant tissues). In addition, biogenic VOCs do not pose problems with residues and environmental accumulation. By contrast, one should consider possible drawbacks deriving from the generally low concentration, impermanence, low target specificity, and difficult handling of VOCs used for plant treatment.

The above reasons may be responsible for the so-far limited enactment of VOC-based technologies. 52 However, some work-arounds may be envisaged to reduce the weight of drawbacks. The use of 53 microbes that form stable populations on plant hosts, exploiting naturally occurring resources and 54 constantly delivering their bio-active function, or that survive harsh conditions (e.g. Bacillus spp. 55 spores), may grant a durable effect of the treatment. In this light, the screening of bacterial species 56 57 forming endophytic and/or specialised symbiosis offers a source of biological functions expressed in an efficient and highly focused way. Recent technological advancement in genomics, 58 metagenomics and gnotobiotics (see glossary) has enabled breeding programs centered on the 59 plant **holobiont** (see glossary), in which, in addition to plant genetic resources, microbial diversity 60 61 (overlooked in traditional breeding, and even possibly lost during domestication or selection) is also explored [8-10]. Alternatively, when live bacteria cannot be used, their active principles could still 62 63 be considered for field application with encapsulation methods allowing a controlled release [11].

Caution should be taken for such treatments in dosing the active principle's release rates (as plant
stress may derive from its excess) and avoiding wastes due to volatilisation.

66

67 VOC profiling for plant disease diagnosis

68 Along with visual inspection, immunochemical and molecular methods represent the standard techniques for disease diagnosis, due to their reliability, sensitivity, specificity, and reasonable 69 70 practicality in terms of costs and work effort [12,13]. However, these methods still pose a number of issues. Since all of them are destructive, the assessment of plant or fruit health status causes an 71 72 economic loss and cannot be applied to unique samples. The development and production of 73 specific antibodies, and the design and validation of PCR primers require a laborious set-up and are 74 conditioned by the availability of sequence or protein data. Moreover, both PCR and immunochemical methods are targeted to single organisms, and the screening for multiple 75 76 pathogens results consequently in a multiplication of work. Finally, representativity at the sampling 77 stage is a major constraint, particularly when pathogen populations are relatively small, and the 78 chance of false negatives must be minimised by increasing the sample size.

79 As a consequence of the bacterial metabolism and the concurrent activation of plant defences, the 80 VOC emission by pathogen-infected plants is, in principle, discernible from that of healthy ones [14]. 81 Thus, the possibility of a VOC-assisted plant diagnosis has been put forward, and the recognition of bacterial diseases by volatile fingerprinting has been attempted in several species and/or crops 82 (Table 1). VOC screening is non-destructive and can be applied to crops or live plants without 83 compromising their economic value or viability. Unlike molecular and immunochemical methods, 84 VOC-assisted plant diagnosis, in principle, allows the screening for multiple pathogens in the same 85 86 run [15,17,25,30,31]. Finally, bulk samples can be analysed as a whole with minimal risks of sampling 87 errors.

88

89 Analytical technologies and methods

90 GC-MS, PTR-MS, SIFT-MS, E-NOSE and FAIMS (see glossary) are among the technologies available 91 for VOC-based diagnosis (Table 2). A major distinction can be made between techniques allowing 92 the analytical determination of the chemical components of the VOC blend (e.g. GC-MS), and 93 techniques that only allow overall VOC profiling (e.g. most E-NOSE models), with in-between cases 94 of techniques with analytical power restricted to certain conditions. Techniques of the first class 95 may be used to identify distinctive marker compounds for determinate pathogens [15,17,18,20-

22,29]. Alternatively, the recognition of infected samples may be done through multivariate
statistical analysis or artificial neural networks. These recognition procedures may be applied to any
technique, but they are an obligate choice for non-analytical methods.

99 Based on the technology and the recognition method, several sampling systems are available. 100 Portable instruments may allow the direct application to ambient air. In these cases, the 101 instrumental sensitivity can be adjusted by regulating the input flow rate (i.e. the air volume 102 screened). Ambient air analysis or headspace sampling in odourless gas bags or canisters are the 103 simplest options, with the smallest chance of artefactual results. However, when such options are 104 not practical (e.g. gas samples are too small, or marker compounds are in trace amounts), the use 105 of VOC-sorbent materials may be envisaged to concentrate the VOC sample [14].

106

107 Sample recognition and applicative perspectives

Besides pathogen infection, other irrelevant factors (such as plant genotype, secondary microbial colonisation, influence of growing substrate on BVC emission, tissue age and environmental conditions during sampling) are predicted to influence VOC emissions with additional levels of complexity. Furthermore, diagnostic power is influenced by disease severity [15,32,33], thus latent infection stages or sporadic pathogen presence are harder to detect. Physico-chemical factors influencing volatilisation and relative composition of air samples (temperature, relative humidity, sorbent saturation or chemical affinity) also require to be accounted for.

Thus, the main challenge for VOC-based diagnosis seems to be the development of feature 115 116 extraction methods, to isolate disease-related information from background noise. For this reason, E-NOSE methods have not progressed beyond the proof-of-principle status so far. In fact, 117 118 discrimination power is related to the independence among the components of data variability [35], while E-NOSE models tested for plant diagnosis include no more than 32 sensors with partially 119 120 overlapping chemical sensitivity [14]. Recent developments in E-NOSE construction, such as coupling with chromatographic separation [36] and colorimetry [37] may overcome current 121 122 discrimination power limits.

With regard to techniques with high analytical power, methods based on unsupervised machine learning and bacteria-VOCs association database were studied for human diagnostic purposes [38]. The implementation of such methods in plant health monitoring could be integrated with existing microbial VOC databases [3]. An effort is required to expand such databases, currently limited in the number of bacterial species covered.

The current technology readiness of VOC-based diagnostic systems may support standard phytosanitary inspection by pre-screening plant material, to focus more in-depth, time- and resource-consuming analyses on dubious cases. Significant advances may come in the future with the development of more versatile instruments [36,37]. Whatever technology may become dominant, coordination among researchers, field operators and industry is a requisite for the setting of standards, databases and accepted practices.

134

135 Bacterial volatile compounds in biological control

Biological control has raised an interest over time, as a tool to achieve a stable level of disease 136 137 control by environmentally sustainable means. Biocontrol agents (BCAs, see glossary) are organisms 138 that reduce a pathogen's population size, or its chance to cause disease, by directly killing the pathogen (with antibiotics, lytic enzymes and other toxic compounds), by interfering with its 139 signalling or regulatory metabolism, or by direct competition (i.e. better exploitation of resources, 140 141 determining the pathogen's starvation). These interactions contribute to microbial antagonism 142 (Figure 1). In this scenario, several BVCs have drawn attention as possible mediators of long-range effects. While it may be expected that their gas-phase concentrations never reach biologically active 143 144 levels as they diffuse in the atmosphere, the competition among microbes in the phyllosphere takes 145 place in matrices (such as biofilms, mucilages, plant waxes) or sites (sub-stomatal chambers, soil pores) where local BVC may attain substantial concentrations [39]. In this light, the identification of 146 147 BVC-releasing symbiotic endophytes may be desirable, as their beneficial effects would be delivered 148 close to their target and in a concentrated form.

Alternatively, BVCs can induce systemic plant defences (Box 2). Notably, such induction occurs at low BVC rates, acts systemically and persists after the removal of the emitter bacterium, whereas BVC-mediated microbial antagonism would require a local and continuous emission at higher rates. Low BVC rates prime, rather than activate plant defences, i.e. responses are more prompt and intense upon pathogen attack, but no phenotypic changes (including in yield and crop quality) are expressed otherwise. In addition, the same plant may release pre-alert signals to neighbouring ones [40]. Thus, even signals in low concentrations may lead to significant large-scale consequences.

BVCs may also influence plant-pest interactions, insect behaviour and survival rate, and thus could be possibly used for pest control strategies (Box 3).

In spite of the potential applications, the use of BCAs presents some inherent difficulties, such as an
 inconstant effectiveness, depending on environmental, agricultural and ecological factors that may

vary in different areas, plant species or growing seasons. In addition, the efficacy of BCAs depends on their population size, which usually decreases steeply after the release, and co-formulants are often required to extend the BCA's field life. For these reasons, BVCs have not yet found specific applications in biological control. As a promising perspective, the development of synthetic bacterial communities may overcome some of these drawbacks, by achieving a better stability or resilience of the microbial biocoenosis, along with the integration of multiple mechanisms of action [41,42].

166

167 Direct toxicity against pathogens

In exerting direct toxicity against plant pathogens, BVCs are influenced by several factors. Along with the chemical nature of the compound, its release rate, the occurrence of the conditions for its production, and the gas phase dynamics that regulate its volatility and stability all contribute to its ecological role and the technological usefulness [39]. The best-studied bacteria, in relation to characterisation of BVC toxicity, include several actinomycetes, *Pseudomonas, Bacillus, Serratia, Burkholderia, Xanthomonas* and *Erwinia* species.

174 Among the volatile compounds hindering the growth of competitors, ammonia, cyanide and sulfurcontaining metabolites are believed to play a major role [43]. However, bacterial strains not 175 176 releasing such compounds can still display antimicrobial properties, indicating that other BVCs 177 substantially contribute to inhibition of microbial growth, and that synergistic effects exist between different compounds. Antimicrobial effects were described for alkanes, alkenes, alcohols, 178 aldehydes, ketones, esters, terpenoids, pyrazines, phenolics, amines, guinolones, chlorine and 179 180 sulfur compounds (Table 3) [44-57]. The most common molecular targets of toxic BVCs include metal cofactors, sulfhydryl groups, and protein folding. 181

Fungi and oomycetes often show a considerable sensitivity to BVCs, both for the elongation of mycelia and for spore germination [43,46-49]. In contrast, fewer cases are reported regarding BVCmediated control of bacterial pathogens, namely, *Agrobacterium* species [48,58], *Clavibacter michiganensis* [59,60], *Xanthomonas oryzae* pv. *oryzae* [61] and *Ralstonia solanacearum* [62]. Reasons for such difference in susceptibility may reside in differences between bacteria and eukaryotic organisms, for instance in plasma membrane composition or gene expression.

The importance of BVC toxicity for interspecific competition in real conditions is debated [63], because of its dependence on BVC production rates and chances of accumulation. Thus, BVCmediated suppression of pathogens was not considered as a trait for selection of new biocontrol agents until recently [49,64]. Nonetheless, this mechanism has been documented for several

commercial biocontrol agents, and some of the BVCs involved, such as benzothiazole and dimethyl 192 193 sulfides [49,64] have been adopted as active principles in exogenous biocide treatments. 194 Biomimicry, i.e. the simulation of biological processes and interactions for applicative purposes, may be advisable in field conditions for a number of reasons, including the caution in introducing 195 196 organisms into a new environment with potentially irreversible effects, and the higher control of chemical nature, dosage and timing of the treatments [65]. Conditions for exogenous VOC 197 198 treatments, however, include the technological feasibility of gas application to the target (soil, 199 canopy, stored crops) and the low toxicity at the treatment dosages for the operator and for non-200 target organisms.

201

202 Disruption of quorum sensing

The complex of regulatory functions connecting the perception of intra- or inter-specific bacterial 203 population density to the expression of 'social' phenotypes is termed Quorum Sensing (QS, see 204 glossary). The typical QS signalling circuit consists of the production of a signal compound, along 205 206 with the expression of specific receptors for the same signal(s). Among the traits governed by QS, 207 bacterial motility, formation of biofilms, biosynthesis of secondary metabolites and virulence factors 208 have been observed, implying their role in improving bacterial fitness in a crowded, diverse and competitive environment [66]. As a consequence, several species may form stable symbiotic 209 consortia, based on the reciprocal exchange of nutrients and signals [67,68]. In the case of 210 pathogenic bacteria, the full expression of virulence can require a stimulation by other microbial 211 212 neighbours [69].

N-acyl-homoserine lactones (AHLs) are the best-studied example of QS signals, since they are employed by a wide array of Gram-negative bacteria, including plant pathogens such as *Pseudomonas, Erwinia, Pectobacterium* and *Agrobacterium* species. Other compounds mediating QS include peptide, aminoacid or fatty acid derivatives. While QS activity was only studied in aqueous solutions for most of these compounds, their semi-volatile nature also suggests an airborne signalling action.

The disruption of QS systems (Quorum Quenching, QQ) may be pursued to reduce the population of pathogens and/or to control the incidence and severity of plant diseases [70]. Interference in QS has been demonstrated for some BVCs. Among these, DMDS reduces the production of AHLs in *P. chlororaphis* [71]. Linear ketones (2-heptanone, 2-nonanone, 2-undecanone) and 2-aminoacetophenone showed an activity on engineered AHL biosensors [72,73].

Indole and its derivatives may act as QS signals, being produced by some bacterial species in a population-dependent manner and eliciting specific responses. Non-producing pathogenic bacteria may also perceive it, possibly by means of AHL receptors [74]. Although auxin (indole-3-acetic acid) and indole are structurally related, in *Agrobacterium tumefaciens* (specialised in auxin biosynthesis, but not releasing indole) only indole can induce bacterial motility, biofilm formation, antibiotic resistance and expression of virulence genes, while reducing bacterial growth in the 0.2-1 mM concentration range [75].

231

232 Biostimulation

233 After the initial observation of plant growth promotion by 2,3-butanediol-emitting bacteria [76], it 234 has become evident that BVC-mediated biostimulation is a widespread phenomenon involving numerous bacterial species and compounds [2,77], with potential applications still to be tested. The 235 adaptive rationale of some biological effects promoted by plant-associated bacteria is, in many 236 237 cases, evident. Symbiotic organisms, for instance, take benefit from increasing root growth and 238 plant nutritional status. Conversely, pathogens can release BVCs to modify plant metabolism to their own advantage [78,79]. However, some methodological caveats should be pointed out in the study 239 240 of influences of VOCs on complex plant traits. Several molecules of great importance for plant 241 metabolism, but not easily detected in the most common experimental settings, for instance, may 242 be neglected (Box 1).

Plant growth is the result of several factors, such as hormonal signalling, nutrition, stress tolerance (Figure 1). Thus, the observation of a plant growth-promoting effect by a bacterial strain in laboratory conditions, in absence of nutritional or cultural constraints, is possibly not indicative of the applicability of the same strain in field. Secondly, plant growth may not correlate (or even inversely correlate) with crop yield, for which not only carbon fixation, but also reallocation of photosynthates is relevant. Thirdly, the effects of BVCs are generally pleiotropic, i.e. they interact with multiple signalling pathways, and are not specific to a definite target organism [43].

250

251 Plant growth promotion and nutrition

Growth promotion by BVCs was shown on several cultivated species, including alfalfa, barley, basil, broccoli, lettuce, poplar, soybean, tobacco, tomato [43,77]. These effects have been related to modulation of plant hormones, such as cytokinins [76], auxins, brassinosteroids [80], gibberellins, ethylene [81] and strigolactones [82].

256 While hormonal effects may shape the allocation of resources within the plant and its phenological 257 progression [79], plant growth promotion and biomass increase should come with a corresponding 258 nutritional enhancement. The stimulation of auxin metabolism and/or signalling, for instance, leads to changes in plant root architecture, which contributes to the uptake of water and nutrients [83]. 259 260 Ethylene participates in the activation of mineral uptake systems [84]. BVCs were implied in counteracting carbohydrate- and ABA-mediated inhibition of photosynthesis, thus enabling higher 261 CO₂ fixation rates [85], and stimulating iron uptake [86,87]. It should be noted that most research 262 was conducted in laboratory or controlled conditions, where nutrients and water are generally not 263 264 limiting.

265

266 Abiotic stress tolerance

Plant growth promotion by microbes can result from increased tolerance to environmental stresses.
Although mechanisms are often far from being elucidated [88], such effects are generally induced
by the release of volatile hormones by the microbes (notably ethylene, methyl-jasmonate and
methyl-salicylate), or involve the signalling cascade of plant hormones [89].

Current knowledge on BVC-induced tolerance refers mainly to osmotic, salt and/or water stress (Box 4). In addition, this research area may provide significant advancement to phytoremediation and to the adaptation of crops to stresses related to climate change [90]. The recovery of marginal soils, for instance, may be enhanced by the root branching stimulation, exerted by some symbiotic bacteria to increase the release of organic carbon into the rhizosphere. Among the BVCs implicated in this plant-microbe interaction, 1-butanol and the QS signal butyrolactone may play a role [91].

277

278 Crop quality

279 A list of examples of bacterial interactions with crops, influencing crop quality, is shown in Table 4. 280 In all the cases in which crop quality depends on secondary metabolites, such as essential oils and aromas, a close link between the elicitation of plant defences and an increased crop value is easily 281 282 explained. In fact, essential oils form one of the first lines of plant defence and inter-plant communication, and their contents are raised by BVCs from several defence-inducing bacteria. Thus, 283 284 an increased essential oil content was obtained by exposing aromatic plants, including peppermint and basil [92,93], or medicinal plants such as Atractylodes lancea [94], to BVCs emitted by several 285 286 Pseudomonas, Bacillus and Azospirillum spp.

In the determination of the aromatic profiles of fruit, a remarkable role has been observed for the associated microflora. *Methylobacterium* spp. include endophytic species expressing alcohol dehydrogenase (ADH) activity, which converts plant-derived alcohols to the corresponding aldehydes or ketones [95,96]. The substrate-specificity of bacterial ADH is low, but distinct from that of plant ADH, thus explaining the diversity, along with the higher intensity, of VOC emissions from microbe-colonised plants.

293 During post-harvest storage, crops may incur in spoilage by pathogens, with consequent loss of produce and/or contamination by mycotoxins. Because of the use of controlled atmosphere on a 294 295 large variety of crops, VOC-based technologies may fit well in post-harvest disease control. In fact, 296 on one hand, relatively high concentrations of bioactive volatiles can be obtained in a closed storage 297 cell; on the other hand, in comparison to synthetic fungicides, the application of biogenic VOCs and BVC-emitting bacteria to products for human consumption poses lower concerns. Several Bacillus 298 299 spp. strains releasing antifungal BVCs were identified and tested on citrus, mango, cherry, litchi and 300 peach [97-102], while Streptomyces spp. were tested on strawberry, citrus, tomato and chili [103-107]. Some compounds mediating antifungal effects, such as cedrol, 2-pentylfuran [102] and 301 acetophenone [107], are commonly found among fruit aromas, and their efficacy was proved on a 302 303 large spectrum of pathogens. Therefore, their technical use may encounter few restrictions by 304 policy-makers, and possibly even higher appreciation by consumers.

The microbial population living on grapevine berries (including *Paenibacillus* spp.) produces volatile compounds possibly improving the quality of wine [108]. Thus, while fruit technology has been so far oriented to the limitation of microbial populations on the crop, future work should address and exploit the contribution of the microflora to aromatic properties of fruit or derivate products.

309

310 Concluding remarks

Despite the great diversity of bacterial metabolites and of biological relations mediated by them, which form a huge reservoir of resources with a potential applicative interest, BVC applications have been explored so far only marginally, and their practical use is at its dawn especially for improving crop tolerance and quality [95]. The present overview was limited to volatile compounds emitted by bacteria, but other organisms (including fungi, moulds and, to some extent, plants) also release bioactive compounds. In addition, biogenic inorganic volatile compounds were only marginally considered. Thus, future agricultural and environmental engineering applications may benefit from

the study of a wider range of biological relations, or by the development of new analytical tools andprotocols.

In modern agricultural systems, there is a growing interest in finding environmentally sustainable, effective and inexpensive solutions for the problems encountered at each step of the production chain. The diffusion of biological control methods, and the programs for phytoremediation or recovery of marginal soils are examples of such dynamics. However, the novelty of these solutions also poses some legislation and registration issues [113,114]. In addition, live biostimulant or biocontrol agents may not adapt to all cultural conditions [90], and promising biological functions may come along with potential risks for human health or environmental equilibrium.

327 Therefore, the mechanisms of interaction among different bacterial species and with their 328 eukaryotic (plant, insect) hosts deserve in-depth investigation, to develop more efficient and flexible solutions for emerging problems. Volatile compounds may show inherent advantages related to 329 their diffusibility, low dose of action and absence of toxic residues [89]. Conversely, environmental 330 331 influences on BVCs (e.g. growing substrate determining volatile blend composition and release rate, interaction among different compounds) may negatively affect their reliability in agricultural 332 practice. Extensive field testing is required as a key step to the commercial and industrial application 333 334 of technologies based on BVCs (see also the outstanding questions).

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336 References

- Schulz, S. and Dickschat, J.S. (2007) Bacterial volatiles: The smell of small organisms. Nat. Prod.
 Rep. 24, 814–42
- Blom, D. et al. (2011) Production of plant growth modulating volatiles is widespread among
 rhizosphere bacteria and strongly depends on culture conditions. Environ. Microbiol. 13, 3047–
 58
- 342 3. Lemfack, M.C. et al. (2018) mVOC 2.0: a database of microbial volatiles. Nucleic Acids Res. 46,
 343 D1261-D1265.
- Tyc, O. et al. (2017) The ecological role of volatile and soluble secondary metabolites produced
 by soil bacteria. Trends Microbiol. 25, 280–92
- Rath, M., Mitchell, T.R. and Gold, S.E. (2018) Volatiles produced by *Bacillus mojavensis* RRC101
 act as plant growth modulators and are strongly culture-dependent. Microbiol. Res. 208, 76–
 84

- van Agtmaal, M. et al. (2015) Legacy effects of anaerobic soil disinfestation on soil bacterial
 community composition and production of pathogen-suppressing volatiles. Front. Microbiol. 6,
 701
- 352 7. Effmert, U. et al. (2012) Volatile mediated interactions between bacteria and fungi in the soil.
 353 J. Chem. Ecol. 38, 665–703
- 354 8. Gopal, M. and Gupta, A. (2016) Microbiome selection could spur next-generation plant
 355 breeding strategies. Front. Microbiol. 7, 1971
- 9. Wei, Z. and Jousset, A. (2017) Plant breeding goes microbial. Trends Plant Sci. 22, 555–8
- 10. Kroll, S., Agler, M.T. and Kemen, E. (2017) Genomic dissection of host-microbe and microbe microbe interactions for advanced plant breeding. Curr. Opin. Plant. Biol. 36, 71–8
- Sharifi, R. and Ryu, C.M. (2020) Formulation and Agricultural Application of Bacterial Volatile
 Compounds. In: Bacterial Volatile Compounds as Mediators of Airborne Interactions (Ryu C.M.,
 Weisskopf, L. and Piechulla, B., eds.), pp. 317-336, Springer
- 362 12. Sankaran, S. et al. (2010) A review of advanced techniques for detecting plant diseases.
 363 Comput. Electron. Agric. 72, 1–13
- 364 13. Martinelli, F. et al. (2015) Advanced methods of plant disease detection. A review. Agron.
 365 Sustain. Dev. 35, 1–25
- 14. Cellini, A. et al. (2017) Potential applications and limitations of electronic nose devices for plant
 disease diagnosis. Sensors 17, 2596
- Cellini, A. et al. (2016) Early detection of bacterial diseases in apple plants by analysis of volatile
 organic compounds profiles and use of electronic nose. Ann. Appl. Biol. 168, 409–20
- 16. Pothakos, V. et al. (2014) Spoilage potential of psychrotrophic lactic acid bacteria (LAB) species:
- 371 *Leuconostoc gelidum* subsp. *gasicomitatum* and *Lactococcus piscium*, on sweet bell pepper
- 372 (SBP) simulation medium under different gas compositions. Int. J. Food Microbiol. 178, 120–9
- 373 17. Vikram, A. et al. (2006) Metabolic fingerprinting to discriminate diseases of stored carrots. Ann.
 374 Appl. Biol. 148, 17–26
- 18. Aksenov, A.A. et al. (2014) Detection of huanglongbing disease using differential mobility
 spectrometry. Anal Chem. 86, 2481–8
- 19. Werbrouck, S., Verholle, P. and Van Langenhove, H. (2017) Volatile indicators of contamination
- in tissue cultures. Acta Hortic. 1155, 239–244

- Zhang, A. and Hartung, J.S. (2005) Phenylacetaldehyde o-Methyloxime: A Volatile Compound
 Produced by Grapefruit Leaves Infected with the Citrus Canker Pathogen, *Xanthomonas axonopodis* pv. *citri*. J. Agric. Food Chem. 53, 5134–5137
- Blasioli, S. et al. (2010) Electronic nose as an innovative tool for the diagnosis of grapevine
 crown gall. Anal. Chim. Acta 672, 20–24
- 22. Cellini, A. et al. (2016) Characterization of volatile organic compounds emitted by kiwifruit
 plants infected with *Pseudomonas syringae* pv. *actinidiae* and their effects on host defences.
 Trees 30, 795-806
- 23. Ioannidis, A-G. et al. (2018) Characterization of spoilage markers in modified atmosphere
 packaged iceberg lettuce. Int. J. Food Microbiol. 279, 1–13
- 24. Li, C., Schmidt, N.E. and Gitaitis, R. (2011) Detection of onion postharvest diseases by analyses
 of headspace volatiles using a gas sensor array and GC-MS. LWT Food Sci. Technol. 44, 1019–
 25
- Prithiviraj, B. et al. (2004) Volatile metabolite profiling for the discrimination of onion bulbs
 infected by *Erwinia carotovora* ssp. *carotovora*, *Fusarium oxysporum* and *Botrytis allii*. Eur. J.
 Plant Pathol. 110, 371–7
- Sinha, R. et al. (2018) FAIMS based volatile fingerprinting for real-time postharvest storage
 infections detection in stored potatoes and onions. Postharvest Biol. Technol. 135, 83–92
- Sinha, R., Khot, L.R. and Schroeder, B.K. (2017) FAIMS based sensing of *Burkholderia cepacia* caused sour skin in onions under bulk storage condition. J. Food Meas. Charact. 11, 1578–85
- Wilson, A.D., Lester, D.G. and Oberle, C.S. (2004) Development of conductive polymer analysis
 for the rapid detection and identification of phytopathogenic microbes. Phytopathology 94,
 419–431
- 402 29. Blasioli, S. et al. (2014) Identification of volatile markers in potato brown rot and ring rot by
 403 combined GC-MS and PTR-MS techniques: Study on in vitro and in vivo samples. J. Agric. Food
 404 Chem. 62, 337–47
- 30. de Lacy Costello, D.L. et al. (1999) Identification of volatiles generated by potato tubers
 (Solanum tuberosum CV: Maris Piper) infected by *Erwinia carotovora, Bacillus polymyxa* and *Arthrobacter* sp. Plant Pathol. 48, 345–51
- 408 31. Kushalappa, A.C. et al. (2002) Volatile fingerprinting (SPME-GC FID) to detect and discriminate
 409 diseases of potato tubers. Plant Dis. 86, 131–137

- 32. Biondi, E. et al. (2014) Detection of potato brown rot and ring rot by electronic nose: From
 laboratory to real scale. Talanta 129, 422-430
- 33. Rutolo, M.F. et al. (2016) Early identification of potato storage disease using an array of metaloxide based gas sensors. Postharvest Biol. Technol. 116, 50–58
- 34. Rutolo, M. et al. (2014) Detection of potato storage disease via gas analysis: a pilot study using
 field asymmetric ion mobility spectrometry. Sensors 14, 15939–52
- 35. Sun, R., Du, H. and Zheng, Y. (2020) Discriminative power of independent component analysis
 applied to an electronic nose. Meas. Sci. Technol. 31, 035108
- 418 36. Wilson, A.D. (2018) Applications of Electronic-Nose Technologies for Noninvasive Early
 419 Detection of Plant, Animal and Human Diseases. Chemosensors 6, 45
- 420 37. Li, Z. et al. (2019) Non-invasive plant disease diagnostics enabled by smartphone-based finger
 421 printing of leaf volatiles. Nat. Plants 5, 856–866
- 38. Palma, S.I.C.J. et al. (2018) Machine learning for the metaanalyses of microbial pathogens'
 volatile signatures. Sci. Rep. 8, 3360
- 39. Bailly, A. and Weisskopf, L. (2017) Mining the volatilomes of plant-associated microbiota for
 new biocontrol solutions. Front. Microbiol. 8, 1638
- 426 40. Heil, M. and Karban, R. (2010) Explaining evolution of plant communication by airborne signals.
 427 Trends Ecol. Evol. 25, 137–144
- 41. Mazzola, M. and Freilich, S. (2017) Prospects for biological soilborne disease control:
 application of indigenous versus synthetic microbiomes. Phytopathology 107, 256–63
- 42. Busby, P.E. et al. 2017. Research priorities for harnessing plant microbiomes in sustainable
 agriculture. PLOS Biol. 15, e2001793
- 43. Piechulla, B., Lemfack, M.C. and Kai, M. (2017) Effects of discrete bioactive microbial volatiles
 433 on plants and fungi. Plant Cell Environ. 40, 2042–67
- 434 44. Kilani-Feki, O. et al. (2012) Correlation between synthesis variation of 2-alkylquinolones and
 435 the antifungal activity of a *Burkholderia cepacia* strain collection. World J. Microbiol.
 436 Biotechnol. 28, 275–81
- 437 45. Yuan, J. et al. (2012) Antifungal activity of *Bacillus amyloliquefaciens* NJN-6 volatile compounds
 438 against *Fusarium oxysporum* f. sp. *cubense*. Appl. Environ. Microbiol. 78, 5942–5944
- 439 46. Velázquez-Becerra, C. et al. (2013) The rhizobacterium *Arthrobacter agilis* produces
 440 dimethylhexadecylamine, a compound that inhibits growth of phytopathogenic fungi in vitro.
 441 Protoplasma 250, 1251-1262

- 442 47. Wang, Z. et al. (2013) Fumigant activity of volatiles from *Streptomyces alboflavus* TD-1 against
 443 *Fusarium moniliforme* Sheldon. J. Microbiol. 51, 477–83
- 444 48. Popova, A.A. et al. (2014) Inhibitory and toxic effects of volatiles emitted by strains of
 445 *Pseudomonas* and *Serratia* on growth and survival of selected microorganisms, *Caenorhabditis* 446 *elegans*, and *Drosophila melanogaster*. Biomed. Res. Int.125704
- 447 49. De Vrieze, M. et al. (2015) Volatile organic compounds from native potato-associated
 448 *Pseudomonas* as potential anti-oomycete agents. Front. Microbiol. 6, 1–15
- 50. Hunziker, L. et al. (2015) Pseudomonas strains naturally associated with potato plants produce
 volatiles with high potential for inhibition of *Phytophthora infestans*. Appl. Environ. Microbiol.
 81, 821–830
- 452 51. Rybakova, D. et al. (2017) Aerial warfare: a volatile dialogue between the plant pathogen
 453 *Verticillium longisporum* and its antagonist *Paenibacillus polymyxa*. Plant Sci. 8, 1294
- 454 52. Xing, M. et al. (2018) Antifungal activity of natural volatile organic compounds against litchi
 455 downy blight pathogen *Peronophythora litchii*. Molecules 23, 358
- 456 53. Guevara-Avendaño, E. et al. (2019) Avocado rhizobacteria emit volatile organic compounds
 457 with antifungal activity against *Fusarium solani*, *Fusarium* sp. associated with Kuroshio shot
 458 hole borer, and *Colletotrichum gloeosporioides*. Microbiol. Res. 219, 74-83
- 459 54. Rajaofera, M.J.N. et al. (2019) Volatile organic compounds of *Bacillus atrophaeus* HAB-5 inhibit
 460 the growth of *Colletotrichum gloeosporioides*. Pestic. Biochem. Physiol. 156, 170–176
- 55. Zhang, Y. et al. (2019) Volatile organic compounds produced by *Pseudomonas chlororaphis*subsp. *aureofaciens* SPS-41 as biological fumigants to control *Ceratocystis fimbriata* in
 postharvest sweet potatoes. J. Agr. Food Chem. 67, 3702–3710
- 464 56. Lyu, A. et al. (2020) High efficacy of the volatile organic compounds of *Streptomyces*465 *yanglinensis* 3-10 in suppression of *Aspergillus* contamination on peanut kernels. Front.
 466 Microbiol. 11, 1–17
- 467 57. Ye, X. et al. (2020) Biocidal effects of volatile organic compounds produced by the
 468 myxobacterium *Corrallococcus* sp. EGB against fungal phytopathogens. Food Microbiol. 91,
 469 103502
- 470 58. Dandurishvili, N. et al. (2011) Broad-range antagonistic rhizobacteria *Pseudomonas fluorescens*
- 471 and *Serratia plymuthica* suppress *Agrobacterium* crown gall tumours on tomato plants. J. Appl.
- 472 Microbiol. 110, 341–52

- 473 59. Lanteigne, C. et al. (2012) Production of DAPG and HCN by *Pseudomonas* sp. LBUM300
 474 contributes to the biological control of bacterial canker of tomato. Phytopathology 102, 967–
 475 73
- 476 60. Rajer, F.U. et al (2017) Volatile organic compounds produced by a soil-isolate, *Bacillus subtilis*477 FA26 induce adverse ultra-structural changes to the cells of *Clavibacter michiganensis* ssp.
- 478 *sepedonicus*, the causal agent of bacterial ring rot of potato. Microbiology 163, 523-530
- 479 61. Xie, S. et al. (2018) Antibacterial effects of volatiles produced by *Bacillus* strain D13 against
 480 *Xanthomonas oryzae* pv. *oryzae*. Mol. Plant Pathol. 19, 49–58
- 481 62. Raza, W. et al. (2016) Volatile organic compounds produced by *Pseudomonas fluorescens* WR482 1 restrict the growth and virulence traits of *Ralstonia solanacearum*. Microbiol. Res. 192, 103–
 483 13
- 63. Sharifi, R. and Ryu, C.M. (2016) Are bacterial volatile compounds poisonous odors to a fungal
 pathogen *Botrytis cinerea*, alarm signals to Arabidopsis seedlings for eliciting induced
 resistance, or both? Front. Microbiol. 7, 1–10
- 487 64. Fernando, W.G.D. et al. (2005) Identification and use of potential bacterial organic antifungal
 488 volatiles in biocontrol. Soil Biol. Biochem. 37, 955–64
- 489 65. Gabriel K.T., Joseph Sexton, D. and Cornelison, C.T. (2018) Biomimicry of volatile-based 490 microbial control for managing emerging fungal pathogens. J. Appl. Microbiol. 124, 1024–31
- 66. Hassani, M.A, Durán, P. and Hacquard, S. (2018) Microbial interactions within the plant
 holobiont. Microbiome 6, 58
- 493 67. Morris, B.E.L. et al. (2013). Microbial syntrophy: interaction for the common good. FEMS
 494 Microbiol. Rev. 37, :384–406
- 68. Mee, M.T. et al. (2014) Syntrophic exchange in synthetic microbial communities. PNAS 111,
 E2149-56
- 497 69. Venturi, V. et al. (2010) Locality versus globality in bacterial signalling: can local communication
 498 stabilize bacterial communities? Biol. Direct. 5, 30
- 499 70. Helman, Y. and Chernin, L. (2015) Silencing the mob: disrupting quorum sensing as a means to
 500 fight plant disease. Mol. Plant. Pathol. 16, 316–29
- 501 71. Chernin, L. et al. (2011) Quorum-sensing quenching by rhizobacterial volatiles. Environ.
 502 Microbiol. Rep. 3, 698–704
- 72. Plyuta, V.A. et al. (2014) The ability of natural ketones to interact with bacterial quorum sensing
 systems. Mol. Genet. Microbiol. Virol. 29, 167–71

- 505 73. Kviatkovski, I. et al. (2015) *Pseudomonas aeruginosa* activates the quorum sensing LuxR
 506 response regulator through secretion of 2-aminoacetophenone. Chem. Commun. 51, 3258–61
- 507 74. Lee, J.-H. and Lee, J. (2010) Indole as an intercellular signal in microbial communities. FEMS
 508 Microbiol. Rev. 34, 426–44
- 509 75. Lee, J.-H. et al. (2015) The multifaceted roles of the interspecies signalling molecule indole in
 510 Agrobacterium tumefaciens. Environ. Microbiol. 17, 1234–44
- 511 76. Ryu, C.M. et al. (2003) Bacterial volatiles promote growth in Arabidopsis. PNAS 100:4927–32
- 512 77. Fincheira, P. and Quiroz, A. (2018) Microbial volatiles as plant growth inducers. Microbiol. Res.
 513 208, 63–75
- 514 78. Weingart, H. et al. (2001) The role of ethylene production in virulence of *Pseudomonas syringae*515 pvs. *glycinea* and *phaseolicola*. Phytopathology 91, 511–8
- 516 79. Sánchez-López, Á.M. et al. (2016) Volatile compounds emitted by diverse phytopathogenic
 517 microorganisms promote plant growth and flowering through cytokinin action. Plant Cell
 518 Environ. 39, 2592-2608
- 80. Bhattacharyya, D., Garladinne, M. and Lee, Y.H. (2015) Volatile indole produced by
 rhizobacterium *Proteus vulgaris* jbls202 stimulates growth of *Arabidopsis thaliana* through
 auxin, cytokinin, and brassinosteroid pathways. J. Plant Growth Regul. 34, 158–68
- 522 81. Tahir, H.A.S. et al. (2017) Plant growth promotion by volatile organic compounds produced by
 523 Bacillus subtilis SYST2. Front. Microbiol. 8, 171
- 524 82. Jiang, C.H. et al. (2019) Volatile organic compounds emitted by *Bacillus* sp. JC03 promote plant
 525 growth through the action of auxin and strigolactone. J Plant Growth Regul 87, 317-328
- 83. Bailly, A. et al. (2014) The inter-kingdom volatile signal indole promotes root development by
 interfering with auxin signalling. Plant J. 80, 758–71
- 84. Iqbal, N. et al. (2013) Current understanding on ethylene signaling in plants: The influence of
 nutrient availability. Plant Physiol. Biochem. 73, 128–38
- 530 85. Zhang, H. et al. (2008) Soil bacteria augment Arabidopsis photosynthesis by decreasing glucose
 531 sensing and abscisic acid levels in planta. Plant J. 56, 264–73
- 86. Wang, J.F. et al. (2017) Enhanced Iron and Selenium Uptake in Plants by Volatile Emissions of *Bacillus amyloliquefaciens* (BF06). Appl. Sci. 7, 85
- 87. Hernández-Calderón, E. et al. (2018) Volatile compounds from beneficial or pathogenic bacteria
 differentially regulate root exudation, transcription of iron transporters, and defense signaling
 pathways in *Sorghum bicolor*. Plant Mol. Biol. 96, 291–304

- 537 88. Garbeva, P. and Weisskopf, L. (2020) Airborne medicine: bacterial volatiles and their influence
 538 on plant health. New Phytol. 226, 32-43
- Kanchiswamy, C.N., Malnoy, M. and Maffei, M.E. (2015) Bioprospecting bacterial and fungal
 volatiles for sustainable agriculture. Trends Plant Sci. 20, 206–11
- 90. Sangiorgio, D. et al. (2020) Facing Climate Change: Application of Microbial Biostimulants to
 Mitigate Stress in Horticultural Crops. Agronomy 10,794
- 543 91. Gutiérrez-Luna, F.M. et al. (2010) Plant growth-promoting rhizobacteria modulate root-system
 544 architecture in *Arabidopsis thaliana* through volatile organic compound emission. Symbiosis
 545 51, 75–83
- 546 92. Banchio, E. et al. (2009) Soil bacteria elevate essential oil accumulation and emissions in sweet
 547 basil. J. Agric. Food Chem. 57, 653–7
- 548 93. Santoro, M.V. et al. (2011) Volatile organic compounds from rhizobacteria increase
 549 biosynthesis of essential oils and growth parameters in peppermint (*Mentha piperita*). Plant
 550 Physiol. Biochem. 49, 1177–82
- 551 94. Zhou, J.Y. et al. (2016) Volatiles released by endophytic *Pseudomonas fluorescens* promoting
 552 the growth and volatile oil accumulation in *Atractylodes lancea*. Plant Physiol. Biochem. 101,
 553 132–40
- 95. Verginer, M. et al. (2010) Monitoring the plant epiphyte *Methylobacterium extorquens* DSM
 21961 by real-time PCR and its influence on the strawberry flavor. FEMS Microbiol. Ecol. 74,
 136–45
- 96. Nasopoulou, C. et al. (2014) Localization of strawberry (*Fragaria* x *ananassa*) and
 Methylobacterium extorquens genes of strawberry flavor biosynthesis in strawberry tissue by
 in situ hybridization. J. Plant Physiol. 171, 1099–105
- 560 97. Arrebola, E., Sivakumar, D. and Korsten, L. (2010) Effect of volatile compounds produced by
 561 *Bacillus* strains on postharvest decay in citrus. Biol. Control 53, 122–8
- 562 98. Chen, K. et al. (2018) Antagonistic activity and the mechanism of *Bacillus amyloliquefaciens* DH563 4 against citrus green mold. Phytopathology 108, 1253–62
- 564 99. Zheng, M. et al. (2013) Antimicrobial effects of volatiles produced by two antagonistic *Bacillus*565 strains on the anthracnose pathogen in postharvest mangos. Biol. Control 65, 200–6
- 566 100. Gotor-Vila, A. et al. (2017) Antifungal effect of volatile organic compounds produced by *Bacillus*
- 567 *amyloliquefaciens* CPA-8 against fruit pathogen decays of cherry. Food Microbiol. 64, 219–25

- 101.Gao, H. et al. (2018) Research on volatile organic compounds from *Bacillus subtilis* CF-3:
 biocontrol effects on fruit fungal pathogens and dynamic changes during fermentation. Front.
 Microbiol. 9, 456
- 571 102. Liu, C. et al. (2018) Antagonistic activities of volatiles produced by two *Bacillus* strains against
 572 *Monilinia fructicola* in peach fruit. J. Sci. Food Agric. 98, 5756–63
- 573 103. Wan, M. et al. (2008) Effect of volatile substances of *Streptomyces platensis* F-1 on control of
 574 plant fungal diseases. Biol. Control 46, 552–9
- 575 104. Li, Q. et al. (2010) Fumigant activity of volatiles of *Streptomyces globisporus* JK-1 against
 576 *Penicillium italicum* on *Citrus microcarpa*. Postharvest Biol. Technol. 58, 157–65
- 105. Wang, C. et al. (2013) Antifungal activity of volatile organic compounds from *Streptomyces alboflavus* TD-1. FEMS Microbiol. Lett. 341, 45–51
- 106. Li, Q. et al. (2012) Effects of volatile substances of *Streptomyces globisporus* JK-1 on control of
 Botrytis cinerea on tomato fruit. Biol. Control 61, 113–20
- 107. Boukaew, S. et al. (2018) Fumigant activity of volatile compounds of *Streptomyces philanthi* RM-1-138 and pure chemicals (acetophenone and phenylethyl alcohol) against anthracnose
 pathogen in postharvest chili fruit. Crop Prot. 103, 1–8
- 108. Verginer, M., Leitner, E. and Berg, G. (2010) Production of volatile metabolites by grape associated microorganisms. J. Agric. Food Chem. 58, 8344–50
- 109. Perpetuini, G. et al. (2019) Genetic and functional characterization of the bacterial community
 on fruit of three raspberry (*Rubus idaeus*) cultivars. J. Berry Res. 9, 227-247
- 110. Feron, G. et al. (2007) Microbial production of 4-hydroxybenzylidene acetone, the direct
 precursor of raspberry ketone. Lett. Appl. Microbiol. 45, 29–35
- 111. Abanda-Nkpwatt, D. and Schwab, W. (2004) Microbial transformation of aliphatic aldehydes by
 Bacillus megaterium to 2,3-dialkylacroleins. J Agric. Food Chem. 52, 5939-5942
- 592 112. Deshmukh, Y., Khare, P. and Patra, D. (2016) Rhizobacteria elevate principal basmati aroma
 593 compound accumulation in rice variety. Rhizosphere 1, 53–7
- 594 113. Calvo, P., Nelson, L. and Kloepper, J.W. (2014) Agricultural uses of plant biostimulants. Plant
 595 Soil 383, 3–41
- 596 114. Backer, R. et al. (2018) Plant growth-promoting rhizobacteria: context, mechanisms of action,
- and roadmap to commercialization of biostimulants for sustainable agriculture. Front. Plant Sci.
- 598 9, 1473

- 115. Jakobson, L. et al. (2016) Natural Variation in Arabidopsis Cvi-0 Accession Reveals an Important
 Role of MPK12 in Guard Cell CO₂ Signaling. PLoS Biol. 14, 1–25
- 116. Piechulla, B. (2017) Considering Microbial CO₂ during Microbe-Plant Cocultivation. Plant
 Physiol. 173, 1529
- 117. Rho, H., Doty, S.L. and Kim, S.H. (2018) Estimating microbial respiratory CO₂ from endophytic
 bacteria in rice. Plant Signal Behav. 13, e1500067
- 118. Nascimento, F.X., Rossi Márcio, J. and Glick, B.R. (2018) Ethylene and 1-Aminocyclopropane-1 carboxylate (ACC) in Plant–Bacterial Interactions. Frontiers in Plant Science 9, 114
- 607 119. Chen, Y. et al. (2020) Root ethylene mediates rhizosphere microbial community reconstruction
 608 when chemically detecting cyanide produced by neighbouring plants. Microbiome 8, 4
- 120. Ravanbakhsh, M. et al. (2018) Microbial modulation of plant ethylene signaling: ecological and
 evolutionary consequences. Microbiome 6, 52
- 611 121. Arasimowicz-Jelonek, M. and Floryszak-Wieczorek, J. (2014) Nitric oxide: An effective weapon
 612 of the plant or the pathogen? Mol. Plant Pathol. 15, 406-416
- 122. Hermenau, R. et al. (2019) Genomics-Driven Discovery of NO-Donating Diazeniumdiolate
 Siderophores in Diverse Plant-Associated Bacteria. Angew. Chem. Int. Ed. 58, 13024-13029
- final formation in tomato. Planta 221, 297-303
 final formation in tomato. Planta 221, 297-303
- 617 124. Mukherjee, S. and Corpas, F.J. (2020) Crosstalk among hydrogen sulfide (H2S), nitric oxide (NO)
- and carbon monoxide (CO) in root-system development and its rhizosphere interactions: A
 gaseous interactome. Plant Physiol. and Biochem. 155, 800-814
- 125. Yamasaki, H. and Cohen, M.F. (2016) Biological consilience of hydrogen sulfide and nitric oxide
 in plants: Gases of primordial earth linking plant, microbial and animal physiologies. Nitric Oxide
 55–56, 91-100
- 126. Hann, C.T. et al. (2014) Methanol and ethanol modulate responses to danger-and microbe associated molecular patterns. Front. Plant Sci. 5, 550
- 127. Ryu, C.M. et al. (2004) Bacterial volatiles induce systemic resistance in Arabidopsis. Plant
 Physiol. 134, 1017–26
- 128. Kong, H.G. et al. (2018) Stereoisomers of the bacterial volatile compound 2,3-butanediol
 differently elicit systemic defense responses of pepper against multiple viruses in the field.
 Front. Plant Sci. 9, 90

- 129. Han, S.H. et al. (2006) GacS-dependent production of 2R,3R-butanediol by *Pseudomonas chlororaphis* o6 is a major determinant for eliciting systemic resistance against *Erwinia carotovora* but not against *Pseudomonas syringae* pv. *tabaci* in tobacco. Mol. Plant-Microbe
 Interact. 19, 924–30
- 130. Marquez-Villavicencio, M.P. et al. (2011) The 3-hydroxy-2-butanone pathway is required for
 Pectobacterium carotovorum pathogenesis. PLoS One 6, e22974
- 131. Spinelli, F. et al. (2012) Emission of volatile compounds by *Erwinia amylovora*: biological activity
 in vitro and possible exploitation for bacterial identification. Trees 26, 141–52
- 132. Rudrappa, T. et al. (2010) The rhizobacterial elicitor acetoin induces systemic resistance in
 Arabidopsis thaliana. Commun. Integr. Biol. 3, 130–8
- 133. Cofer, T.M., Seidl-Adams, I. and Tumlinson, J.H. (2018) From acetoin to (Z)-3-hexen-1-ol: the
- diversity of volatile organic compounds that induce plant responses. J. Agric. Food Chem. 66,
 11197–208
- 134. Huang, C.J. et al. (2012) Dimethyl disulfide is an induced systemic resistance elicitor produced
 by *Bacillus cereus* C1L. Pest Manag. Sci. 68, 1306–1310
- 135. Farag, M., Zhang, H. and Ryu, C. (2013) Dynamic Chemical Communication between Plants and
- Bacteria through Airborne Signals: Induced Resistance by Bacterial Volatiles. J. Chem. Ecol. 38,
 1007-1018
- 136. Xu, Y.Y. et al. (2015) Effect of volatile organic compounds from bacteria on nematodes. Chem.
 Biodivers. 12, 1415–21
- 137. Davis, T.S. et al. (2013) Microbial volatile emissions as insect semiochemicals. J. Chem. Ecol. 39,
 840–59
- 138. Venu, I. et al. (2014) Social attraction mediated by fruit flies' microbiome. J. Exp. Biol. 217,
 1346–52
- 139. Wallingford, A.K. et al. (2016) Behavioral response of spotted-wing drosophila, *Drosophila suzukii* Matsumura, to aversive odors and a potential oviposition deterrent in the field. Pest.
 Manag. Sci. 72, 701–6
- 140. Dillon, R.J., Vennard, C.T. and Charnley, A.K. (2000) Exploitation of gut bacteria in the locust.
 Nature 403, 851–851
- 141. Farine, J.-P. et al. (2017) Maternally-transmitted microbiota affects odor emission and
 preference in *Drosophila* larva. Sci. Rep. 7, 6062

- 661 142. Leroy, P.D. et al. (2011) Microorganisms from aphid honeydew attract and enhance the efficacy
 662 of natural enemies. Nat. Commun. 2, 348
- 143. D'Alessandro, M. et al. (2014) Volatiles produced by soil-borne endophytic bacteria increase
 plant pathogen resistance and affect tritrophic interactions. Plant Cell Environ. 37, 813–26
- 144. Niu, Q. et al. (2010) A Trojan horse mechanism of bacterial pathogenesis against nematodes.
 PNAS 107, 16631–6
- 145. Hamby, K.A. and Becher, P.G. (2016) Current knowledge of interactions between *Drosophila suzukii* and microbes, and their potential utility for pest management. J. Pest. Sci. 89, 621–30
- 146. Colman, D.R., Toolson, E.C. and Takacs-Vesbach, C.D. (2012) Do diet and taxonomy influence
 insect gut bacterial communities? Mol. Ecol. 21, 5124–37
- 147. Hadapad, A.B. et al. (2016) Diversity of bacterial communities in the midgut of *Bactrocera cucurbitae* (Diptera: Tephritidae) populations and their potential use as attractants. Pest.
 Manag. Sci. 72, 1222–30
- 148. Cho, S.M. et al. (2008) 2R,3R-butanediol, a bacterial volatile produced by *Pseudomonas chlororaphis* O6, is involved in induction of systemic tolerance to drought in *Arabidopsis thaliana*. Mol. Plant-Microbe Interact. 21, 1067–75
- 149. Zhang, H. et al. (2010) Choline and osmotic-stress tolerance induced in Arabidopsis by the soil
 microbe *Bacillus subtilis* (GB03). Mol. Plant-Microbe Interact. 23, 1097–104
- 150. Cappellari, L.d.R. and Banchio, E. (2020) Microbial volatile organic compounds produced by
 Bacillus amyloliquefaciens GB03 ameliorate the effects of salt stress in *Mentha piperita* principally through acetoin emission. J. Plant Growth Regul. 39, 764–775
- 151. Zhang, H. et al. (2008) Soil bacteria confer plant salt tolerance by tissue-specific regulation of
 the sodium transporter HKT1. Mol. Plant-Microbe Interact. 21, 737–44
- 152. Vaishnav, A. et al. (2016) PGPR-mediated expression of salt tolerance gene in soybean through
 volatiles under sodium nitroprusside. J. Basic Microbiol. 56, 1274–88
- 153.Ledger, T. et al. (2016) Volatile-mediated effects predominate in *Paraburkholderia phytofirmans* growth promotion and salt stress tolerance of *Arabidopsis thaliana*. Front.
 Microbiol. 7, 1838
- 154. Lantz, A.T. et al. (2019) Isoprene: New insights into the control of emission and mediation of
 stress tolerance by gene expression. Plant Cell Environ. 42, 2808–2826
- 155. McGenity, T.J., Crombie, A.T. and Murrell, J.C. (2018) Microbial cycling of isoprene, the most
- abundantly produced biological volatile organic compound on Earth. ISME J 12, 931–941

- 693 156. Xue, J. and Ahring, B.K. (2011) Enhancing isoprene production by genetic modification of the 1-
- deoxy-D-xylulose-5-phosphate pathway in *Bacillus subtilis*. Appl Environ Microbiol. 77, 2399–
- 695 405

697 **Box 1. Low molecular weight volatile compounds**

Some volatile compounds, characterised by a small and simple chemical structure, interact directly with plant metabolism or signalling cascades. However, these molecules have received, in explorative studies, only marginal attention, because of technical limits, including the impossibility of a correct identification by GC-MS (due to short retention times and/or scarcely indicative spectral profiles), or the occurrence of artefacts (such as accumulation effects in sealed cuvets, or interference with growing media) linked with the experimental setting.

Carbon dioxide fuels photosynthesis and it is often the most important limiting factor of this process. 704 705 CO₂ regulates stomata opening and, consequently, photosynthesis and transpiration [115]. CO₂ may 706 contribute to the growth and resistance promoting effects observed in plants treated with bacterial 707 VOCs [116]. CO₂ produced by respiration of bacterial endophytic symbionts can re-enter the photosynthetic pathway, not being limited by stomata opening, and has been estimated to be able 708 709 to provide up to 57% of total CO₂ photo-assimilated by the plant [117]. In this view, plants colonised by endosymbionts may have a better water use efficiency and a higher availability of 710 photoassimilates for growth and defences. 711

Ethylene is a gaseous plant hormone playing a central role in plant development and resistance response to abiotic and biotic stresses [118], also interacting with salicylic acid- and jasmonic aciddependent signalling pathways. Ethylene and its precursor 1-aminocyclopropane-1-carboxylate (ACC) are subjected to sophisticated co-regulation by plants and associated microbes, thus shaping the plant microbiome [119,120]. In fact, bacteria can actively produce ethylene, or reduce its biosynthesis in plants subtracting ACC by ACC-deaminase activity, thus lowering plant ethylene levels and promoting plant growth.

Nitric oxide (NO) is a radical gas mediating a large variety of physiological responses in plants. Plantassociated bacteria can produce NO, as a result of denitrification, by enzymatic conversion of Larginine or by release from siderophores [121,122]. One important effect of bacterial NO, observed e.g. in *Azospirillum* spp. [123], is the enhancement of root branching, promoting both plant nutritional status and bacterial colonisation.

While commonly regarded as a toxic and/or defence compound, hydrogen sulfide (H₂S) also has a regulative role in plants by interacting with NO and thiols [124,125]. Plant-associated bacteria synthesise H₂S through cysteine desulfhydrylation or sulfite reduction.

Ethanol and methanol are common products of fermentation, originated by both plants and
bacteria under anoxic conditions, and have been implied in the activation of plant stress responses
[126].

730

731 Box 2. BVC-elicited induction of plant defences

Since the discovery of induced systemic resistance (ISR, see glossary) by 2,3-butanediol [127], the 732 733 potential application of BVCs for the elicitation of plant defences has drawn attention. Notably, 734 induction of plant defences is one of the very few measures that can be adopted against viral diseases [128]. ISR is often stimulated as the result of a specific symbiotic interaction between the 735 736 host plant and bacteria, which also promote plant nutrition and growth. Thus, the two aspects of 737 defence and growth promotion coexist in the same symbiotic relation and are somewhat difficult to tell apart (e.g. activation of stress responses in Figure 1). However, relevant details for plant defence 738 engineering, including mechanisms of signal perception and decoding (i.e., how relatively simple 739 740 molecules drive specific responses), remain obscure [88].

2,3-butanediol biosynthesis, for instance, was observed both in defence-eliciting (*Bacillus* and *Serratia* spp., *Pseudomonas chlororaphis*) and pathogenic (*Erwinia, Pectobacterium, Dickeya* spp.) bacteria [127,129-131]. In different pathosystems, the action of 2,3-butanediol and related compounds (acetoin, 2,3-butanedione) has been connected to different combinations of salicylic acid, jasmonate and/or ethylene signal cascades [127,128,132]. According to the relative stimulation of these pathways, specific subsets of plant defensive responses may be activated.

While 2,3-butanediol and related compounds are the best-studied example of defence-inducing BVCs, other molecules [43,133] were identified which could stimulate plant defences. In some cases, such as for DMDS and benzothiazole, direct antimicrobial and plant defence induction effects may coexist [134]. One advantage of these compounds is that, although acting through plant hormone signal cascades, they are less prone than hormones to cause drastic physiological reprogramming. For instance, ISR is expressed only after pathogen challenge, and is not generally associated to changes in plant phenotype or crop yield [40,135].

Synergism of BVCs in complex mixtures may also occur in natural conditions [88]. By modulating the
 simultaneous activation of several signal cascades (ethylene, jasmonate, salicylate, and other
 hormones), BVC mixtures could attain protection against a broader range of pathogens [135].

- 757
- 758

759 Box 3. Pest management by BVCs

Survival and replication rate of pests (including *Drosophila suzukii* and several nematodes) were reduced by means of bacterial volatile emissions [48,136]. However, the use of toxic BVCs seems impractical against most motile animal species, due to limitations in exposure time and concentration. Instead, microbial biocontrol agents releasing toxic compounds may act as effective biopesticides for soil-borne or sessile pests, with a significantly reduced environmental impact.

765 Insects and nematodes use a variety of semiochemicals to coordinate their life functions, including feeding, mating, oviposing and alarm behaviour. Pest- or plant-associated bacteria contribute to the 766 767 production of biologically active BVCs (Figure 1), and many cases of attraction to microbes 768 associated to the host have been observed [137]. Fruit flies (Drosophila spp.), for instance, are 769 attracted by BVCs from symbiotic *Lactobacillus* spp. acting as aggregation pheromones [138], and are repelled by the common BVCs, 1-octen-3-ol and geosmin [139]. Locusts use guaiacol derivatives, 770 produced by *Pantoea agglomerans* residing in their intestines, as an aggregation pheromone [140]. 771 Finally, the association with certain bacteria may determine the insect's preference for BVCs 772 emitted by those microbes, by effect of conditioning or learning [141]. Parasitoid recruitment can 773 774 be mediated by BVCs, either by direct attraction to microbes indicating a food source [142], or 775 indirectly by eliciting a more intense release of plant VOCs [143]. Concerning nematodes, 776 experiments on the model organism Caenorhabditis elegans showed that attractive BVCs produced by Bacillus nematocida, such as benzyl benzoate, benzaldehyde, 2-heptanone, and acetophenone, 777 778 stimulate bacterial swallowing by the host. Thus, the bacterium colonises the worm's intestines, 779 leading to its death [144].

780 BVCs may, therefore, find an application in pest management. VOC-based technologies, employing 781 attractants, deterrents and pheromone-like compounds have been applied to lure-and-kill, push-782 pull and sexual confusion control strategies. However, information concerning semiochemicals for 783 such technologies is currently restricted to a relatively small number of species (mostly Lepidoptera, Diptera and Coleoptera). New impulse, in this sense, may derive from metagenomic analyses 784 performed either on the insect or on its hosts [145]. With the exception of specialised symbiotic 785 relations, insect guts have often been demonstrated to host relatively simple microbiota, dominated 786 787 by Enterobacteriaceae (notably Enterobacter and Klebsiella spp.), to be highly influenced by the diet, and possibly transmitted by parents [141,146,147]. This body of knowledge may provide useful 788 789 information for the selection of effective and persistent biocontrol agents interacting with insects.

790

791 Box 4. Examples of abiotic stress tolerance induced by BVCs

Abiotic stresses elicit NO and ethylene production in plants, which exert several and multifaceted physiological effects. Plant-associated bacteria may indirectly influence plant NO and ethylene emission or produce these bioactive compounds (Box 1).

795 Water deficiency, osmotic stress and salt toxicity are partially interconnected and overlapping both in causes and in the induction of plant responses. 2,3-butanediol, or total BVCs from 2,3-butanediol-796 emitting bacteria Bacillus subtilis GB03 and Pseudomonas chlororaphis O6, increased Arabidopsis 797 tolerance to water deficiency and osmotic stress. Abscisic acid, salicylic acid, ethylene, and jasmonic 798 799 acid signaling pathways were implicated in *P. chlororaphis* O6- and 2,3-butanediol-induced stomata 800 closure, increasing tolerance to drought [148]. B. subtilis GB03 stimulates the biosynthesis of 801 osmoprotectants (choline, glycine-betaine) in the plant, enhancing its growth under water withholding and osmotic treatment [149]. 802

803 Several cases of improved plant tolerance to salt stress have been observed after interaction with BVC-releasing bacteria. B. subtilis GB03 and its main volatile, acetoin, enhance peppermint 804 805 tolerance to salt stress by stimulating SA biosynthesis and reducing ABA [150]. Another mechanism 806 of induction of Na⁺ stress tolerance in Arabidopsis consists of the tissue-specific modulation of HKT 807 ion transporters [151]. Such transporters are downregulated in roots to reduce Na⁺ uptake and 808 upregulated in shoots to promote internal recirculation. NO, produced by salt-stressed plants, was also implicated in the enhanced colonisation of the rhizosphere by Pseudomonas simiae strain AU, 809 which in turn elicits antioxidant defences, osmoprotection and expression of ion transporters in 810 811 soybean [152]. Other salt tolerance-inducing BVCs (namely, 2-undecanone, 1-heptanol and 3methyl-butanol) were identified from Parabulkholderia phytofirmans [153]. 812

813 Drought stress and high temperature promote isoprene emission by plants [154]. Isoprene is the most abundantly produced biogenic VOC on Earth, with an estimated emission of more than 2% of 814 815 all photoassimilates. Isoprene has a likely a role in protection from reactive oxygen and nitrogen species formed under diverse stress conditions [155]. Many Proteobacteria, Actinobacteria and 816 817 Firmicutes produce isoprene. Bacillus and related genera are among the terrestrial bacteria accounting for the highest production of isoprene. Interestingly, isoprene emission by Bacillus 818 819 subtilis is enhanced by supra-optimal temperature and salinity [156], suggesting that also plantassociated B. subitlis may mediate the plant reactions to these stresses. Soil and phyllosphere 820 821 bacteria may also directly influence the host isoprene metabolism by using plant-derived isoprene 822 as carbon source [155].

823 Table 1. Applications of VOC-based recognition of bacterial pathogens in different plant species

and organs, indicating the diagnostic techniques and the VOCs marker(s)

Crop species and conditions	Bacterial pathogen/contaminant	Methods	Distinctive features and remarks	Refs.
Apple, dormant plants	Erwinia amylovora Pseudomonas syringae pv. syringae	GC-MS E-NOSE PTR-MS	Multiple pathogen discrimination Dilution effects Markers: acetoin, 2,3-butanediol, 2-hexenal, phenylethanol (<i>E. amylovora</i>)	[15]
Bell pepper-derived medium	Leuconostoc gelidum ssp. gasicomitatum and Lactococcus piscium	GC-MS SIFT-MS	Control of spoilage off-odours by controlled atmosphere	[16]
Carrot, roots	Pectobacterium carotovorum	GC-MS	Multiple pathogen discrimination Markers: 3-methyl-butan-1-ol, 1-propanol, 2,3- butanedione	[17]
<i>Citrus sinensis</i> , asymptomatic plant	C. liberibacter	GS-MS FAIMS	Correct identification of PCR-false negatives Severity-dependent markers: methyl-salicylate (severe), geranyl acetone, linalool (mild)	[18]
Ficus benjamina and Spathiphyllum wallisii, in vitro cultures	Escherichia coli contamination	SIFT-MS		[19]
Grapefruit, leaves	Xanthomonas axonopodis pv. citri	GC-MS	Marker: phenylacetaldehyde O-methyloxime	[20]
Grapevine, rootstock cuts	Agrobacterium vitis	GC-MS E-NOSE	Marker: styrene	[21]
Kiwifruit <i>, in vitro</i> explants	Pseudomonas syringae pv. actinidiae	GC-MS E-NOSE PTR-MS	Marker: 1-undecene	[22]
Lettuce	Resident microflora (mainly Pseudomonas spp.)	GC-MS SIFT-MS	Control of spoilage off-odours by packaging	[23]
Onion, bulbs	Burkholderia cepacia	GC-MS E-NOSE	Markers: 2-nonanone, 2-octyl-5-methyl-3(2H)- furanone	[24]
Onion, bulbs	Pectobacterium carotovorum	GC-MS	Multiple pathogen interaction Marker: 3-bromo-furan	[25]
Onion, bulbs	Burkholderia cepacia	FAIMS		[26,27]
Poplar, wood	Bacterial wetwood (non- determined species)	E-NOSE		[28]
Potato, tubers	Clavibacter michiganensis ssp. sepedonicus	GC-MS PTR-MS	Marker: 2-propanol	[29]
Potato, tubers	Pectobacterium carotovorum Bacillus polymyxa Arthrobacter sp.	GC-MS	Markers: 2-propenal, DMDS, 1-alkenes, branched alkanes, octanal, naphtalene, butanoic acid (<i>P.</i> <i>carotovorum</i>); N,N-dimethylformamide, 1- pentadecene (<i>B. polymyxa</i>); 2,3-dihydrofuran (<i>Arthrobacter</i> sp.)	[30]
Potato, tubers	Pectobacterium carotovorum	GC	Multiple pathogen discrimination	[31]
Potato, tubers	Ralstonia solanacearum, Clavibacter michiganensis ssp. sepedonicus	E-NOSE	Lab- to real scale Threshold of disease severity for recognition	[32]
Potato, tubers	Pectobacterium carotovorum	E-NOSE	Pre-symptomatic recognition	[33]
Potato, tubers	Pectobacterium carotovorum	FAIMS	Early detection (1 d post inoculation)	[34]
Potato, tubers	Pectobacterium carotovorum	FAIMS		[26]

825

827 Table 2. Overview of analytical techniques employed for VOC-based plant diagnosis

Analytical technique	Working principle	Operative features	Diagnostic principle
GC-MS	Differential retention time and	+ High analytical power (identification of	Recognition of markers
	fragmentation profile of VOCs	single compounds)	Multivariate statistical analysis
		+ May use sample concentration on sorbents	
		(e.g. SPME)	
PTR-MS	Fragmentation profile of VOCs	+ Partial analytical power	Partial recognition of markers
SIFT-MS		+ Quick response	Multivariate statistical analysis
Electronic nose	Electric properties of the overall VOC	+ Simple operation, portability	Multivariate statistical analysis
	mixture	+ Quick response	Neural network machine learning
		+ May adjust sensitivity by regulating flow	
		- Interference by water	
		- No analytical power	
		- Instrumental drift	
FAIMS	Differential mobility of ion fragments in	+ Partial analytical power	Partial recognition of markers
	electric field	+ Portability	Multivariate statistical analysis

830 Table 3. Volatile organic compounds exerting direct toxicity against plant pathogens

Compound(s)	Emitting species	Target organism(s)	Refs.
Hydrogen cyanide Ammonia 1-Undecene	Pseudomonas spp. Bacillus spp. others	Phytophthora infestans Rhizoctonia solani Helminthosporium solani Fusarium oxysporum Dickeya dianthicola	[49]
2-(2'-heptyl)-3-methyl-4-quinolone	Burkholderia cepacia	Aspergillus niger and other fungi	[44]
Alkylated benzene derivatives Phenol derivatives Naphthalene derivatives Benzothiazole 2-Ethyl-1-hexanol 2-Undecanol 2-Nonanone 2-Decanone 2-Undecanone Nonanal Decanal	Bacillus amyloliquefaciens NJN-6	Fusarium oxysporum f. sp. cubense	[45]
N,N-dimethyl- hexadecanamine	Arthrobacter agilis UMCV2	Botrytis cinerea Phytophthora cinnammomi	[46]
2-methyl-isoborneol	Streptomyces alboflavus TD-1	Fusarium moniliforme	[47]
1-Undecene 2-Nonanone 2-Undecanone	Pseudomonas chlororaphis	Agrobacterium tumefaciens Synechococcus spp. Rhizoctonia solani	[48]
Dimethyl disulfide 2-Heptanone	Serratia proteamaculans	Agrobacterium tumefaciens Synechococcus spp. Rhizoctonia solani	[48]
3-hexanone 1-dodecene isovaleric acid 5-methyl-butanethioate 5-methyl-methanethiosulfonate furfuryl alcohol acetophenone phenylpropanedione 2- acetylthiazole nitropentane	Pseudomonas spp.	Phytophthora infestans	[49]
2-(2-Methylpropyl)-3-(1-methylethyl) pyrazine 2- Isopropylpyrazine 2- methyl-1-butanol Hexadecanal Isoamyl acetate	Paenibacillus polymyxa Sb3-1	Verticillium longisporum	[51]
Phenylethyl alcohol Methyl salicylate Ethyl phenylacetate Methyl anthranilate α-Copaene Caryophyllene 4-Ethylphenol Humulene	Streptomyces fimicarius BWL-H1	Peronophythora litchii	[52]
2,3,5-Trimethylpyrazine 2-Nonanone 2-Decanone 2-Dodecanone Dimethyl disulfide Dimethyl trisulfide	Bacillus spp. Pseudomonas spp.	Fusarium spp. Colletotrichum gloeosporioides	[53]
Hexanedioic acid, bis(2-ethylhexyl) ester Octadecane 1-Hexadecanol Docosane Chloroacetic acid, tetradecyl ester	Bacillus atrophaeus HAB-5	Colletotrichum gloeosporioides	[54]

2-methyl-1-butanol ethyl hexanoate 3-methyl-1-butanol ethyl octanoate phenylethyl acetate phenylethyl alcohol	Pseudomonas chlororaphis subsp. aureofaciens SPS-41	Ceratocystis fimbriata	[55]
2-methylbutyrate 2-phenylethanol	Streptomyces yanglinensis 3-10	Aspergillus flavus A. parasiticus	[56]
Isooctanol Linalool 3-Octanone 2-Naphthalene methanol 3-Undecanone 2-Tridecanone	Corrallococcus sp. EGB	Fusarium oxysporum f. sp. cucumerinum Penicillium digitatum	[57]
Dimethyl disulfide	Pseudomonas fluorescens B-4117 P. fluorescens Q8r1-96 Serratia plymuthica IC1270	Agrobacterium tumefaciens A. vitis	[58]
2,4-diacetylphloroglucinol Hydrogen cyanide	Pseudomonas sp. LBUM300	Clavibacter michiganensis ssp. michiganensis	[59]
Benzaldehyde Nonanal Benzothiazole Acetophenone	Bacillus subtilis FA26	Clavibacter michiganensis ssp. sepedonicus	[60]
3,5,5-trimethylhexanol Decyl alcohol	Bacillus cereus D13	Xanthomonas oryzae pv. oryzae	[61]
Toluene Ethyl benzene m-xylene Benzothiazole 2-decanol 2-tridecanol 1-undecanol Dimethyl disulfide Benzaldehyde 1-Methyl naphthalene	Pseudomonas fluorescens WR-1	Ralstonia solanacearum	[62]

833 Table 4. Examples of bacterial interactions with crops influencing quality parameters

Crop plant	Bacterial species	Quality parameter(s)	Mechanism of interaction	Refs.
Sweet basil	Bacillus subtilis GB03	Increased production of essential oils		[92]
Peppermint	Pseudomonas fluorescens WCS417r, Bacillus subtilis GB03, Azospirillum brasilense SP7	Increased production of essential oils		[93]
Atractylodes lancea	Pseudomonas fluorescens ALEB7B	Increased production of essential oils	Benzaldehyde mediates the effect	[94]
Strawberry	Methylobacter spp.	Production of aromatic compounds (furanones)	Bacterial alcohol dehydrogenase	[95,96]
Raspberry	Methylobacter spp., Bacillus spp.	Production of aromatic compounds (frambinone)		[109,110]
Strawberry	Bacillus megaterium	Production of aromatic compounds (2,3- dialkylacroleins)	Conversion of linear aldehydes	[111]
Basmati rice	Acinetobacter spp.	Production of aromatic compounds (2-acetyl-1- pyrroline)		[112]
Citrus, mango, cherry, litchi, peach	Bacillus spp.	Protection from spoliage	Antifungal action of cedrol and 2-pentylfuran	[97-102]
Strawberry, citrus, tomato, chili	Streptomyces spp.	Protection from spoliage	Antifungal action of acetophenone	[103- 107]
Grapevine	Paenibacillus spp.	Production of aromatic compounds in wine production		[108]

Figure 1. Summary of VOC-mediated biological functions of plant-associated bacteria.

836 Biological effects of plant-associated bacteria and their mechanisms of interaction with the host 837 plant and the environment can be exploited in the agricultural practice. The resident bacteria may increase the availability of certain mineral nutrients, or stimulate plant growth and stress responses 838 839 by means of hormones or other signalling compounds. As a result, a better nutritional status and a better ability to cope with stresses is achieved in the host plant. In their interaction with pests and 840 pathogens, plant-associated bacteria may act as direct competitors and/or predators with a biocidal 841 action, or exert a disturbance in long-range signalling, possibly influencing the pest's behaviour, its 842 recognition by natural enemies, and the expression of «social» phenotypes related to virulence in 843 844 pathogens.

845

846 Glossary

BCA: Biological Control Agent, an organism exerting directly (e.g. by killing or predating) or indirectly
(by competition for resources, or through the action of other organisms) a limiting effect on the
population of a pest or pathogen.

BVCs: Bacterial Volatile Compounds, including organic (i.e. carbon-containing) and inorganic (e.g.
H₂S, nitrogen oxides) compounds.

E-NOSE: Electronic nose, a device including an array of electric sensors with differential affinity for different chemical classes, and variating their electric conductance upon interaction with the components of a gas blend. Used to compare gas samples, has good portability and ease of operation, allows real-time analysis, but not chemical identification.

FAIMS: Field Asymmetric Ion Mobility Spectrometry, analytical method based on the separation of
ions in an oscillating electric field. Allows real-time analysis of gas profiles with good portability, and
can be coupled to GC-MS for chemical identification.

GC-MS: Gas Chromatography-Mass Spectrometry, analytical technique based on separation of molecules in a gas mixture according to affinity to a chromatographic column, followed by their fragmentation to yield a typical spectrum. Most used technique for identification of volatile compounds.

Gnotobiotics: study of test organisms, in which the resident microbial community is artificial,
 controlled and/or completely characterised.

Holobiont: the complex formed by a host organism and its associated microflora.

ISR: Induced Systemic Resistance, condition of increased and generalised plant resistance to
 potential pathogens and pests, activated after interactions with microbes (including beneficial
 symbionts).

Metagenomics: study of the complex of genomes associated in one super-organism, such as a plant
with its associated microflora.

PTR-MS: Proton Transfer Reaction-Mass Spectrometry, analytical technique based on
fragmentation of gas compounds in an electric field. Allows highly sensitive real-time detection and
tentative identification of compounds.

QS: Quorum Sensing, bacterial communication system allowing the coordination of 'social'
 phenotypes (motility, biofilm formation, etc.) according to population density.

876 **SIFT-MS**: Selected Ion Flow Tube-Mass Spectrometry, analytical technique based on fragmentation 877 of gas compounds in an air flow. Allows real-time detection and tentative identification of 878 compounds.

VOCs: Volatile organic compounds, organic molecules characterised by low vapour pressure, high
lipophylicity and low molecular weight, normally found in the gas phase in standard conditions.

881 Volatilome: the complete set of volatile compounds originating from an organism or biological882 system.

Outstanding questions

- Bacteria can release complex mixture of BVCs, including simple volatile compounds (NO, ethylene, H₂S, CO₂), acting directly on plant metabolism, signalling and ecological interactions. Simple BVCs often elude explorative studies. What is the actual contribution of simple molecules to plant growth, crop yield and stress defence *in situ*?
- Because VOCs may be released from diverse sources, in a variety of contrasting conditions and rates, their specificity as signals should be low. How do plants perceive environmental VOCs, and integrate VOC signals to produce specific and dose-dependent responses?
- Beyond effects of single compounds, the effects on plant physiology and resistance of complex VOCs mixtures as a whole have been overlooked. What are the interactions among VOCs in complex mixtures? Could VOC ratios, rather than single VOCs, carry specific information? What are the interactions between VOCs and other environmental parameters (e.g. oxygen, UV light, humidity)?
- Do production rates and persistence of antimicrobial volatile compounds support a role in bacterial competition and/or elicitation of plant responses in field conditions?
- Biological and analytical parameters should be standardised for a sensitive, reliable and robust VOC-based pathogen detection.
- Could innovative formulations facilitate the application of BVC-releasing beneficial bacteria and improve their efficacy in field?
- Is BVC-enhanced plant growth effective under oligotrophic/stress conditions?
- The establishment of stable populations of BVC-emitting beneficial bacteria is a requisite for
 a successful application in agriculture. Research on bacterial consortia, formulation as
 biostimulant products and plant breeding for ability to form symbiotic relations could
 facilitate the colonisation and resilience of beneficial bacteria.

