Changes in biodiversity impact atmospheric chemistry and climate through

plant volatiles and particles

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Abstract

Climate extremes in tandem with biodiversity change affect emissions of biogenic volatile organic compounds (BVOCs) from plants and, as a result, the formation of biogenic secondary organic aerosols (BSOA). The resulting BSOA can have a wide variety of impacts, such as on Earth's radiative balance or cloud- and precipitation formation. However, at present, it is unclear how changing biodiversity will lead to changes in BVOC emissions, BSOA formation and their corresponding effects. We present a conceptual framework of the relationships between biodiversity and BVOC emissions based on our current mechanistic understanding and combining knowledge from the fields of biology and atmospheric chemistry. Parts of this framework are tested in a case study using a tree diversity experiment with adjunct BVOC and BSOA characterisation. The relative differences in tree monocultures and mixtures show that the overall concentration of BVOCs decreases with increasing biodiversity ($p < 0.01$), but results for BSOA compounds are mixed and overall non-significant ($p = 0.40$). We suggest future studies should follow a multidisciplinary approach where the fields of biology, atmospheric chemistry and climate research interact.

Keywords: Biodiversity, biogenic aerosol, BSOA, biosphere-atmosphere interactions, biotic and abiotic stress, BVOCs, climate change, volatile organic compounds

Introduction

Biosphere and atmosphere are tightly interconnected [1,2], making the understanding of their interactions critically important as human life depends on them [3,4]. On the one hand, a suite of atmospheric drivers can affect the biosphere in different ways [2], including detrimental impacts of rising temperatures on ecosystem integrity and biodiversity [5]. On the other hand, the biosphere can exert feedbacks to the atmosphere, e.g. by releasing various biogenic volatile organic compounds (BVOCs) through plants [6,7] or soil [8]. Following their emission, the atmospheric oxidation of plant-emitted BVOCs leads to gas-phase products, which can either condense on already existing particles or form new ones [9], resulting in biogenic secondary organic aerosols (BSOA) [10], which are important particles for the radiative balance of the Earth [11]. Climate extremes as well as biodiversity change affect emissions of BVOCs from plants [8,12] and, consequently, BSOA formation [13]. However, studies using concerted measurements of BVOCs and BSOA along diversity gradients of emitting plant species are largely missing [14]. This lack of knowledge is particularly worrying given that human activities have traditionally favoured monoculture plantations [15], and scientific guidance is urgently needed for major reforestation efforts, especially in the light of current global climate change [16].

Tree species release a large variety of hydrocarbons, especially isoprene for deciduous trees and monoterpenes for coniferous trees. However, the composition and magnitude of emitted BVOCs are highly species-specific [17,18]. These BVOCs are produced primarily via the leaf surface as a result of metabolic processes and as a reaction to abiotic and biotic stress [19,20]. Plant-released BVOCs determine important biotic interactions such as intra- and interspecific communication [21] as well as herbivore and pathogen defences [22]. BVOCs can also protect plants against abiotic stress such as heat, drought, and high radiation [20,23].

In the forest, individual trees are not isolated but compete with their con- and heterospecific neighbours for resources, such as light, water, and nutrients [24]. Resource availability directly affects eco-physiological processes such as photosynthetic rates [25–27] and thus allocation to growth and potential interaction strength. In monocultures or low diverse forest stands, neighbouring trees share largely identical ecological niches and thus strongly compete for available resources. In contrast, mixed forests enable resource partitioning, e.g. via differences in tree architecture [28] or rooting depth [29,30], often leading to higher stand productivity [31] and leaf area index [32]. Likewise, forests with diverse leaf chemical traits can enhance soil nutrient availability through diverse plant inputs of leaf and root litter and through soil microbial communities and activities [33,34], thereby increasing forest stand

productivity. As forest biomass production and BVOC emissions are highly correlated [35,36],

diverse forests should thus emit higher amounts of BVOC (Figure 1, Hypothesis 1).

Yet, there are at least two counteracting mechanisms which should lead to decreased

BVOC emissions with increasing tree diversity. First, more diverse plant communities

increase facilitation and thereby might mitigate abiotic stress. It has been shown that

functionally diverse plant communities can increase water use efficiencies through variation

in root architecture as a result of spatiotemporal complementary resource use [29] and

microclimate amelioration, particularly in dry seasons [37]. In addition, complexity in the

vertical and horizontal stratification of diverse stands can relieve thermal stresses [38,39] and

alleviate the risks of drought stress [40]. Indeed, decreasing heat and drought stress in mixed

communities is likely to result in lower leaf temperature, which could result in lower BVOC

emissions [41,42]. Given that abiotic stress, but particularly heat and drought are known to be

important drivers of BVOC emissions, we hypothesize a decrease in BVOC concentrations in

mixtures where abiotic facilitation mediates abiotic stress reduction (Figure 1, Hypothesis 2).

Second, more diverse plant communities show decreased per capita herbivory damage

and pathogen infection [43,44]. Although multi-plant diets may be beneficial for different

herbivores *per se* [45,46], plant natural enemies are also more abundant in more diverse plant

communities, which could reduce herbivory pressures as well as insect outbreaks (top-down

regulation), as supposed by the "enemies hypothesis" [44,47]. Furthermore, the heterogeneity of

plant nutritive traits in diverse stands suppresses herbivore abundance, resulting in declining

herbivore performance [48], likely due to non-host species reducing the accessibility of host

species [49]. Overall, reduced biotic stress through herbivores and pathogens in mixtures should

reduce the amount of BVOC emissions (Figure 1, Hypothesis 3) but might increase the

diversity of emitted compounds. However, irrespective of the change in the amount of BVOC

emissions with increasing tree diversity – which might increase or decrease depending on the

primary mechanism – we would expect that the diversity of emitted compounds increases in

mixed forests given that BVOC emissions are known to be highly species-specific [50].

Once entering the atmosphere, plant-derived VOCs react quickly with ambient

hydroxyl radicals (OH), Ozone (O_3), and nitrate radicals (NO_3) resulting in the formation of

BVOC oxidation products that are more oxidized than the compounds originally emitted.

These compounds, including highly oxygenated molecules (HOMs) or, as more recently

called, oxygenated organic molecules (OOMs) [51,52], have lower vapor pressures and better

water solubility and can hence contribute to BSOA as atmospheric particles [53]. Changing

BSOA fractions in organic particle composition could have different effects. First, it could

influence radiative effects of the particles themselves ('direct effect') and hence couple to climate. Secondly, such compositional changes could influence the particles' ability and effectiveness to act as cloud condensation nuclei (CCN), thus influencing cloud formation with all its respective consequences for radiation through aerosol-cloud interactions ('indirect effects') that also affect precipitation formation rates [54–56]. However, the conversion of BVOC to BSOA is strongly determined by the local atmospheric composition and oxidation regime, temperature, the kinetics and mechanism of the emitted BVOCs and their respective reactions in the gas phase including their phase transfer characteristics and their particle-phase chemistry [57–59]. High BVOC emissions, e.g. due to rising temperature, increase BSOA formation which, to some extent, could trigger a heating by direct particle absorption but it could also lead to a cooling due to the effect of BSOA on cloud formation and reflectivity of clouds [11,13]. Recent findings suggest that biotic and abiotic stress-induced BVOC emissions even accelerate climate-relevant BSOA formation [60–62].

Our mechanistic understanding and existing knowledge suggest that increasing biodiversity could predictably influence the amount of BVOC released. So far, studies have primarily focused on single tree species though, with no studies available at the time using concerted measurements of BVOCs and BSOA in tree stands or along experimental or natural tree diversity gradients (Figure 1). We argue that this is a critical knowledge gap, as BVOC composition has been recognized as a powerful stress indicator and an important feedback mechanism of climate change [63]. Addressing this knowledge gap is relevant to understand the causal connections between biodiversity and climate change and to overcome these coupled crises as two of the most urgent challenges facing humanity [64]. To support our claim that biodiversity might play a significant role in this context and to inspire future research, we present the first data from a case study in the MyDiv tree diversity experiment site in Germany [65]. Here, we simultaneously measured the magnitude and variability of BVOC and BSOA compounds in ten plots differing in tree diversity. Using this case study, we aim to put forward the link between biodiversity and BVOC/ BSOA and provide a first test of principle in support of the conceptual framework. Our results show that the amount of BVOCs tends to decrease with biodiversity in most cases, while mixed results were found for BSOA compounds. Our findings highlight the need for multidisciplinary work at the interface between the biosphere and the atmosphere to better understand the reciprocal effects of biodiversity and climate change [4].

Results and discussion

BVOC emissions across biodiversity

We quantified nine different BVOCs from the investigated plots in the MyDiv experiment, i.e. α-pinene, camphene, β-pinene, △3-carene, p-cymene, limonene, α-terpinene, isophorone, and acetophenone. Our results show that most of the observed BVOC compounds were not significantly different between the individual tree species in monocultures and their corresponding mixtures, with some notable exceptions (Figure 2). In particular, we find some significant differences in the emission of limonene and acetophenone (ANOVA: $p = 0.01$ and $p < 0.01$, respectively; Figure 2A,I) which was higher in monoculture plots, particularly in *Sorbus aucuparia* L. (Figure 2A,I), highlighting the species-specific dependency of BVOC emission rates [17,18].

The comparison between the observed and expected amounts of each BVOC compound in mixture plots (two- and four-species mixtures) shows that there were no significant differences; however, we found a few exceptions (Figure 3). In four-species mixtures, some BVOC compounds differed significantly between the observed and expected amounts (Figure 3). As such, the observed amounts of limonene and acetophenone were significantly lower than the expected values (t-test: $p = 0.04$ and $p = 0.01$, respectively; Figure 3C,I). This indicates that four-species mixtures produced lower amounts of limonene and acetophenone than expected.

The relative differences in monocultures and mixtures based on standardized mean difference analysis across diversity levels show that increasing tree diversity significantly decreased the overall concentration of BVOCs (the overall standardized mean difference: $p < 0.01$, Figure 4). Particularly, the relative differences in the means of observed and expected amounts of limonene were significant (the standardized mean difference: $p < 0.01$, Figure 4), and p-cymene and α-terpinene were marginally significant (the standardized mean difference: $p = 0.09$ and $p = 0.07$, respectively; Figure 4), meaning that mixtures produced significantly less than expected amounts of these compounds compared to monocultures. The relative differences in monoculture and mixtures for two-and four-species mixtures also showed the same decreasing trend of BVOC emissions with increasing tree diversity and were significant (the standardized mean difference, $p < 0.01$, Supplementary Figure 1). As such, the relative differences in the means of observed and expected amounts of limonene were marginally significant in two-species mixtures (the standardized mean difference: $p = 0.06$, Supplementary Figure 1A), while acetophenone and α-terpinene showed strong and marginal significance in four-species mixtures, respectively (the standardized mean difference: $p = 0.02$

and $p = 0.09$, respectively; Supplementary Figure 1B). The overall results indicate that tree

diversity significantly reduced BVOC emissions, supporting our second and third hypotheses

(H2 and H3).

Consistent with our second hypothesis of abiotic stress amelioration (H2), the decline

in BVOC emissions with increasing biodiversity could partly be due to a reduced competition

for resources (e.g., light and water) because the emission of BVOCs acts as a defence strategy

during plant competition [66]. Besides, mixed plots can alleviate thermal and water stress [38,67]

through spatial and structural complexity of canopy cover [67] and also differences in rooting

depth [67]. In this context, microclimate buffering is important, as at a global scale, an increase

of 2–3 °C in the mean global temperature due to global warming has been shown to increase

global BVOC emissions by 30–45% [68]. Decreasing BVOC emissions with increasing

biodiversity also supports our third hypothesis of reduced biotic stress (H3), and can be

explained by multiple mechanisms: First, a higher percentage of leaf area eaten by herbivores,

which may occur in monoculture due to a higher probability of herbivores[44], can increase

BVOC emissions [69]. Second, increasing plant diversity decreases herbivore per capita effects

on plants while simultaneously benefiting predators [43,44]. Third, heterogeneity in plant

strategies in highly diverse communities reduces herbivore performance [70]. Fourth, plant-

herbivore interactions also highly depend on the heterogeneity of available nutrients. For

instance, herbivore abundance and performance decrease with plant nutrient variability [47,48],

indicating that diverse stands are expected to have reduced herbivory damage [71], resulting in

lower BVOC emissions. Altogether, biotic and abiotic patterns may result in lower emission

rates of BVOC in mixtures compared to monocultures, stressing the potential importance of

both, biotic and abiotic factors on BVOC emissions[23,72], yet further experiments are urgently

needed to properly test these hypotheses.

BSOA formation across biodiversity

We quantified a total of fifteen BSOA compounds on the investigated plots throughout the

2021 sampling period. These analytes were diaterpenylic acid acetate [DTAA], 3-methyl-

1,2,3-butanetricarboxylic acid [MBTCA], norpinonic acid, pinonic acid, terebic acid,

terpenylic acid, pinic acid, adipic acid, pimelic acid, azelaic acid, suberic acid, succinic acid,

glutaric acid, salicylic acid, and sebacic acid. Although the observed BSOA compounds did

not significantly differ between individual tree species and their corresponding mixtures

(Figure 5), succinic and sebacic acid were tendentially higher in monoculture compared to

mixture plots (Figure 5L,O); by contrast, pinonic and terebic acid were tendentially higher in

mixture plots compared to monoculture plots (Figure 5D,E).

While we found no significant differences between observed and expected amounts in
BSOA compounds, individual mixtures produced tendentially more or less BSOA compounds
than expected. In particular, a mixture of *Acer pseudoplatanus* L. and *Fraxinus excelsior* L.
and also a mixture of *Prunus avium* (L.) L. and *Sorbus aucuparia* L. produced a larger than
expected amount of norpinonic acid (Figure 6C). Similarly, the mixture of *Acer*
pseudoplatanus L. and *Fraxinus excelsior* L. produced a larger than expected amount of
glutaric acid (Figure 6M).

Similarly, the relative differences analysis shows that different compounds responded
differently to increasing tree diversity, showing that some BSOA compounds (like 3-methyl-
1,2,3-butanetricarboxylic acid [MBTCA], pinonic acid, pinic acid, and terpenylic acid)
increase relative to what is expected from monocultures but others (like norpinonic acid,
adipic acid, suberic acid, and azelaic acid) decrease; however, the overall results were mixed
and non-significant (the standardized mean difference: $p = 0.40$, Figure 7). Although the
relative differences analysis pattern in two-species and four-species mixtures revealed mixed
results, the overall results were significant for two-species mixtures (the standardized mean
difference: $p = 0.05$, Supplementary Figure 2A), but non-significant for four-species mixtures
(the standardized mean difference: $p = 0.71$, Supplementary Figure 2B).

The mixed results might be due to the influence of regional air masses, as the
atmospheric conversions from BVOC to BSOA require some time corresponding to spatial
spread and thus preventing the identification of a local BSOA occurrence patterns coupled to
the measured BVOC. However, in certain cases, we observed indications of local linkage
between atmospheric radicals and α-pinene across individual tree species and mixtures
(Supplementary Figure 3). Although our knowledge about biodiversity effects on BSOA
formation is limited, it seems likely and it can well be expected that biotic and abiotic stresses
also matter indirectly for BSOA formation [60–62]. For instance, using a compact chamber,
temperature dependence of BSOA formation from α- pinene has been reported [73], which
might partly be related to physical or abiotic differentiation in mixtures versus monocultures.
Earlier studies observed a higher formation rate of OH radicals in humid conditions [74], which
may increase the formation of some BSOA compounds even if BVOCs are low. In addition,
the complex formation patterns of BSOA reflect the potential that source precursor and
atmospheric oxidants can also matter for oxidation products [57,58]. Specifically, the differential
response of BSOA compounds to diversity gradients could partly be explained by the
difference in the gas-phase mechanisms related to the oxidation of α- and β-pinene [75,76]. For
example, MBTCA, which is formed from the oxidation (OH radical) of α-pinene, is a stronger

functionalized compound than pinic and pinonic acid [77]. Overall, we acknowledge that testing

our hypothesis needs further experimental work to improve our understanding of how the

magnitude and composition of BVOC and BSOA will change across tree diversity gradients.

Conclusions and research perspectives

In light of significant human-induced changes in biodiversity and climate, the link between

atmospheric and biological measurements is crucial to improve our understanding of

atmosphere-biosphere feedbacks. However, to what extent changes in atmospheric chemistry

and climate change are related to biodiversity is largely unknown. Here we show that the

general relationship between biodiversity and BVOC emissions works in principle. The aim

of this study is not to properly test our hypotheses, but rather to put forward the relationship

between biodiversity and BVOC or BSOA. So, with this first and limited dataset any

generalization would be premature. Based on our mixed results, we argue, that measuring

BVOC emissions and BSOA formation across biodiversity gradients alone is not enough to

fully understand their magnitude and composition. A deeper understanding requires in-depth

investigations of microclimate conditions, above- and below-ground herbivores and

pathogens, soil microbial communities, accurate monitoring of biotic and abiotic stress, and

manipulating biotic and abiotic stress across long-term biodiversity experiments. In addition,

we need local and regional-scale models, combined with field and chamber measurements, to

improve our understanding of biosphere-atmosphere interactions. These experimental

platforms along with the expertise exist in each of the separate disciplines. Therefore, a

multidisciplinary approach at the biosphere-atmosphere interface would extend our

understanding of the reciprocal effects of biodiversity and climate change [4] and help to

unravel the extent to which emissions of isoprenoids and aerosol formation are related to

biodiversity. This will open a new area of research where the fields of biology, climate

science, and atmospheric chemistry may interact.

Methods

We tested the effect of tree diversity on BVOC emission and BSOA formation by varying tree

species richness, including monocultures, two- and four-species mixtures at the MyDiv

experimental site located in Saxony-Anhalt, Germany [65], between September 21 and October

13, 2021 (13 days in total). The MyDiv experiment comprises eighty 11×11 m plots that are

located 2 m apart. We used ten of these plots in our case study. The monoculture plots

consisted of *Acer pseudoplatanus* L., *Fraxinus excelsior* L., *Prunus avium* (L.) L., and *Sorbus*

aucuparia L. The two-species mixture plots consisted of three sets: a mixture of *A.*

pseudoplatanus and *F. excelsior*; a mixture of *A. pseudoplatanus* and *P. avium*; and a mixture
of *P. avium* and *S. aucuparia*. The four-species mixture plot consisted of all four species. Two
monoculture plots, one two-species mixture plot, and one four-species mixture plot were
sampled per day for at least four consecutive days (except rainy days). BVOC emissions and
BSOA compounds were captured simultaneously using samplers installed at the top of the
canopy of each plot. The aluminum sampling box was custom-made and equipped with gas
phase (for BVOC) and particulate matter (for BSOA) sampling systems. The gas phase
sampling system consisted of a 2-cm-diameter tube with a glass-fritted disc to prevent insect
intrusion. The tube was connected to Carbotrap 300 and Tenax cartridges (Supelco, Bad
Homburg, Germany) and a GilAir Plus pump (Sensidyne, USA) operating at 150 ml/min
using flexible PVC tubes. The particulate matter (PM) system consisted of a PM10 PEM
impactor (SKC, USA) inlet connected to a Gillian 12 pump (Sysidyne, USA) operating at 10
l/min. We used an off-line method, where the applied adsorbent cartridges for capturing
BVOC and quartz filters for capturing BSOA lasted for four hours (10:00 AM to 2:00 PM).
After four hours of sampling, the cartridges and filters were transferred to the laboratory for
further analysis. After each day of sampling, the filters were eluted with 3 ml of acetonitrile.

Quantification of BVOC compounds trapped in adsorbent cartridges was then
conducted by gas chromatography mass spectrometry (GC/MS, Agilent, Waldbronn,
Germany) combined with thermodesorption (Perkin Elmer, Rodgau, Germany). To do so, the
BVOCs were first desorbed from the cartridges with a heating rate of 7 ˚C/min and a flow rate
of 50 ml/min to a final temperature of 220 ˚C. Then the BVOCs were cryofocused at -30 ˚C
on a cooling trap. The trap was heated to 220 ˚C (hold time: 3 min) with 1 ml/min and the
BVOCs were transferred to the Zebron ZB-5ms (30m; 0,25mm; 0,25µm) capillary column.
The initial temperature of the column was 35 ˚C (hold time: 5 min) and then increased
stepwise to 120 ˚C (ramp: 5 ˚C / min) and finally to 320 ˚C (ramp: 20 ˚C / min) and hold for
12 min. Finally, the observed BVOCs were identified using known calibration standards. As a
result of this, the following compounds were identified: α-pinene, β-pinene, camphene, △3-
carene, p-cymene, limonene, α-terpinene, isophorone, and acetophenone.

To analyse BSOA constituents every filter was cut into small pieces and transferred to
an extraction vial. 1.0 mL of a water acetonitrile mixture (1:1) was used as an extraction
solvent. All samples were shaken for 30 min at 1000 min^{-1} and then transferred into vials with
a syringe filter (pore size: 0.2 µm, wwPTFE membrane). The analysis of the extracts was
done with ultrahigh-performance LC system equipped with a C_{18} column (Acquity HSST3,
1.8 mm, 2.1 × 100 mm, Waters). The eluents were ultrapure water with 0.1% formic acid

(eluent A) and acetonitrile with 0.1% formic acid (eluent B). With a flow rate of 0.3 mL

min^{-1}, the following gradient was used: 5% B at 0 min, 5% B at 1.0 min, 100% B at 16.0 min,

100% B at 18.0 min, 5% B at 18.1 min, 5% B at 21.0 min. The LC effluent was connected to

a high resolution Orbitrap mass spectrometer (Q Exactive Plus, Thermo Fisher Scientific)

equipped with an ESI source operated in the negative mode (3.5 kV). Each sample was

measured three times in full scan mode (m/z 50−750, R = 70k at m/z 200). Finally, the

following BSOA compounds were quantified via an external calibration: diaterpenylic acid

acetate [DTAA], 3-methyl-1,2,3-butanetricarboxylic acid [MBTCA], norpinonic acid, pinonic

acid, terebic acid, terpenylic acid, pinic acid, adipic acid, pimelic acid, azelaic acid, suberic

acid, succinic acid, glutaric acid, salicylic acid, and sebacic acid.

We first compared the amounts of observed emitted BVOC and BSOA compounds across

biodiversity gradients using a one-way analysis of variance (ANOVA) test, and where the

ANOVA test was significant, we used Tukey's HSD for pairwise multiple comparisons. Due

to the highly reactive nature of α- and β-pinene with atmospheric oxidants such as ozone and

OH radical [78,79], we performed the Pearson's correlation coefficients between the products

(reacted between α- and β-pinene and ozone and ultraviolet radiation) and the observed

BSOA compounds. We analysed the correlation coefficients separately for each measurement

week to account for weekly changes of abiotic conditions.

Next, we calculated the expected amount of each BVOC or BSOA compound in the

mixture from their respective amounts of observed compounds in monoculture based on the

species richness of the mixture plot:

Expected amount of compounds in mixture plot $= \dfrac{\sum \text{Observed amount of compound in monoculture}}{\text{Plot species richness}}$

We then compared the observed and expected amounts of each BVOC or BSOA

compound using a two-sample t-test.

After that, we used a commonly known Cohen's d effect size[80], the standardized mean

difference (SMD), to quantify the magnitude and direction of the biodiversity effects on

BVOC emission or BSOA formation in monocultures and mixtures. For this, we calculated

the SMD for each BVOC compound or BSOA formation and their overall differences

between monoculture and mixture using the observed and expected values of each BVOC or

BSOA compound [81] to see the differences in BVOC emission or BSOA formation between

monoculture and mixtures.

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$$\text{SMD} = \frac{\text{Mean}_{\text{expected}} - \text{Mean}_{\text{observed}}}{\sqrt{(\text{standard deviation}^2_{\text{expected}} + \text{standard deviation}^2_{\text{observed}})/2}}$$

This measure calculates the effect size for each BVOC or BSOA compound, and also for
the overall by combining the observed and expected values of all quantified BVOC or BSOA
compounds. This approach also assigns a weight to each SMD using the inverse of the
variance [81] (within-compound variance), which requires the mean, standard deviation as well
as the number of observations [81] for the observed and expected compounds. To infer how tree
diversity levels contribute to BVOC emission or BSOA formation, we performed three
separate SMD analyses for BVOC or BSOA compounds for the overall mixtures (a
combination of two-and four-species mixtures), two-species mixtures, and four-species
mixtures. We finally visualized the results using a forest plot, which shows the effect size of
each individual BVOC or BSOA compound and the overall estimates. As such, positive
values indicate that the mixture produced a higher amount of compounds than expected based
on the monoculture data, and vice versa.

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Data availability

We will store the dataset in the public repository once the paper is accepted.

Competing interests
The authors declare no competing interests.

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Figure captions

Figure 1. The figure outlines (A) the processes and mechanisms, and (B) the general

hypotheses related to biogenic volatile organic compound (BVOC) emissions across

biodiversity gradients (monoculture vs. mixtures), biogenic secondary organic aerosol

(BSOA), and cloud condensation nuclei (CCN) formation. The hypothesized positive and

negative relationships between BVOC emissions and biodiversity are represented by blue and

red lines, respectively. The hypothesized positive relationship between BVOC emissions and

BSOA formation is indicated in black.

Figure 2. Changes in observed emitted BVOC compounds across individual tree species and

mixtures. Boxes indicate interquartile ranges (bottom and top parts of the box), and median

lines (horizontal lines within boxes) and whiskers indicate the minimum and maximum of the

measurement. *p*-values are shown. Different letters above boxes represent significant

differences among tree species ($p < 0.05$; Tukey's test). Abbreviations: So, *Sorbus aucuparia*

L.; Pr, *Prunus avium* (L.) L.; Ac, *Acer pseudoplatanus* L., and Fr, *Fraxinus excelsior* L.

Figure 3. Changes in observed and expected BVOC compounds across tree mixtures (two-

and four-species mixtures). Boxes indicate interquartile ranges (bottom and top parts of the

box), and median lines (horizontal lines within boxes) and whiskers indicate the minimum

and maximum of the measurement. *p*-values are shown (t-test). Abbreviations: So, *Sorbus*

aucuparia L.; Pr, *Prunus avium* (L.) L.; Ac, *Acer pseudoplatanus* L., and Fr, *Fraxinus*

excelsior L.

Figure 4. A forest plot summarizing the results of the standardized mean difference between

monocultures and mixtures for changes in BVOCs as a result of diversity change. The

squares' sizes and widths represent each compound's weight, while the diamond (red colour)

represents the overall effect estimate of the analysis. Effect sizes were calculated using the

standardized mean difference. The horizontal lines of the squares show 95% confidence

intervals (CI). *p*-values are given for each compound and for the overall effect.

Figure 5. Changes in BSOA compounds across individual tree species and mixtures. Boxes

indicate interquartile ranges (bottom and top parts of the box), and median lines (horizontal

lines within boxes) and whiskers indicate the minimum and maximum of the measurement. *p*-

values are shown. Abbreviations: So, *Sorbus aucuparia* L.; Pr, *Prunus avium* (L.) L.; Ac,

Acer pseudoplatanus L.; Fr, *Fraxinus excelsior*; DTAA, diaterpenylic acid acetate, and

MBTCA, 3-methyl-1,2,3-butanetricarboxylic acid.

Figure 6. Changes in observed and expected BSOA compounds across tree mixtures (two- and four-species mixtures). Boxes indicate interquartile ranges (bottom and top parts of the box), and median lines (horizontal lines within boxes) and whiskers indicate the minimum and maximum of the measurement. Abbreviations: So, *Sorbus aucuparia* L.; Pr, *Prunus avium* (L.) L.; Ac, *Acer pseudoplatanus* L.*;* Fr*, Fraxinus excelsior*; DTAA, diaterpenylic acid acetate, and MBTCA, 3-methyl-1,2,3-butanetricarboxylic acid.

Figure 7. A forest plot summarizing the results of the standardized mean differences between monocultures and mixtures for changes in BSOA compounds as a result of diversity change. The squares' sizes and widths represent each compound's weight, while the diamond (red colour) represents the overall effect estimate of the analysis. Effect sizes were calculated using the standardized mean diffrenece. The horizontal lines of the squares show 95% confidence intervals (CI). *p*-values are given for each compound and for the overall effect. Abbreviations: DTAA, Diaterpenylic acid acetate, and MBTCA, 3-methyl-1,2,3-butanetricarboxylic acid.

Figure 1.

676   **Figure 2.**

Figure 3.

700 **Figure 4.**

Test for overall effect: $z = -4.77$ ($p < 0.01$)

Figure 5.

Figure 6.

750 **Figure 7.**

Test for overall effect: z = -0.85 (p = 0.40)

751