

Interactive comment on “High levels of primary biogenic organic aerosols in the atmosphere in summer are driven by only a few microbial taxa from the leaves of surrounding plants” by Abdoulaye Samaké et al.

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The manuscript describes the contribution of primary biogenic organic aerosols (PBOAs) to PM₁₀, which were collected during the summer of 2017 (June–August) in a rural area of France. The quartz fiber filters (24-hour samples) were collected using high-volume sampling systems. The collected samples were analyzed for detailed chemical composition (inorganic ions, OCs and ECs, sugars and sugar alcohols) and for biological constituencies (DNA sequencing and analysis). Soil and vegetation samples were also collected and analyzed with the same techniques for comparison. The

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goal of this study was to investigate the association between the chemical composition of PM₁₀ (especially sugar compounds or SCs) and the identified PBOAs. This study is scientifically very important because very little is known about the contribution of bioaerosols to atmospheric particulate matter and what kinds of markers can be used for the quantitative analysis of bioaerosols in particular fungi and bacteria. The manuscript is well written and organized. I have several major comments.

We thank the anonymous referee for taking the time to evaluate this manuscript, and for all the suggestions for modifications and comments that helped us improve the quality of this work. We have taken all the comments into account and have made a point by point revision. Detailed responses to the comments are given below, point by point, in blue, including changes made directly to the manuscript, in red.

Major Comments:

(1) The title of the manuscript doesn't represent the research of this paper (it shows some of the results, but not the overall scope of this study)

We thank the reviewer for this remark, which was also suggested by anonymous referee 1. The title has been changed as follows “High levels of primary biogenic organic aerosols are driven by only a few plant-associated microbial taxa” in the main text.

(2) The author performed a comparison between SC concentrations and collected bioaerosols, assuming that SCs in the atmospheric aerosols are mainly due to PBOAs. Other sources (e.g., biomass burning, the ocean) can also emit SCs (sugars and sugar alcohols). The author is missing the entire discussion of these possible sources. Therefore, without a proper comparison of SC emissions from different sources, the statement regarding “suitable markers” (line 77) should be carefully used. A discussion on other SC sources is needed.

The reviewer is correct that other sources, including biomass burning and the ocean, have sometimes been proposed as potential emitters of SCs (Yang et al., 2012). How-

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ever, recent studies conducted at several sites across France have shown a weak correlation between daily concentrations of SC and levoglucosan in PM_{2.5} and PM₁₀ collected throughout the year (Golly et al., 2018; Samaké et al., 2019a). In the present study, there was no significant correlation between primary sugar species and levoglucosan, a tracer of biomass burning, in our PM₁₀ time series. In addition, primary sugar compounds were not significantly related to two typical marine ions (e.g. Na⁺ or Cl⁻) or methanesulfonic acid, a tracer of marine biogenic activity (Zhu et al., 2015). It therefore seems unlikely that sources of SC in PM₁₀ from biomass burning or ocean were significant at this site.

As suggested by the reviewer, a discussion of other potential sources proposed in a few previous studies has been added in the main text as follows:

Lines 68-73: "SC species are emitted from biologically derived sources (Medeiros et al., 2006, Verma et al., 2018) and have sometimes been detected in aerosols taken from air masses influenced by smoke from biomass burning (Fu et al., 2012; Yang et al., 2012). However, recent studies conducted at several sites across France revealed a weak correlation between daily concentrations of SC and levoglucosan in PM_{2.5} and PM₁₀ collected throughout the year (Golly et al., 2018; Samaké et al., 2019a). This suggests that open burning of biomass is not a significant source of SC in the environments studied here".

Lines 490-497: "However, in our study, SC species are not correlated ($R = -0.09$, $p = 0.46$; Fig. S7) with levoglucosan during the campaign period, confirming that biomass burning is not an important source of airborne microbial taxa associated with SCs in our PM₁₀ series. Bubble bursting associated with sea spray could also potentially be a source of Bacteria, Fungi and water-soluble organic species, along with sea salts, to PM₁₀ (Prather et al., 2013; Zhu et al., 2015). However, SC species were not found to be significantly related to Cl⁻ ($R = -0.14$, $p = 0.28$) or Na⁺ ($R = -0.18$, $p = 0.16$), which are two inorganic tracers typical of marine sources; nor correlated with methanesulfonic acid ($R = -0.05$, $p = 0.69$), a well-known tracer of biogenic marine activity (Arndt et al.,

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2017; Gaston et al., 2010). It therefore seems unlikely that the sources of SCs from marine environments were significant at this site".

(3) Pollen can be a huge contributor to atmospheric PBOAs. Why were only bacteria and fungi collected and analyzed?

The reviewer is correct that pollen can be a significant contributor to atmospheric PBOAs. However, this abundance is expected to also depend on the PM size range considered. Individual airborne pollen grains generally range about 10-100 μm (Manninen et al., 2014; Yoo et al., 2017), while fungal spores are much smaller, 1-30 μm , and most often < 10 μm (Després et al., 2012; Manninen et al., 2014). Similarly, the diameter of airborne Bacteria is generally between 0.25 and about 8 μm (Yoo et al., 2017). Our study therefore focused on Bacteria and Fungi as they are generally the dominant biological component of ambient aerosols in the size range of 2-10 μm (Zhang et al., 2010), discussed in this study.

(4) What standard deviations represented in this paper (e.g., lines 270-282)? It is unclear how they were calculated.

The standard deviations presented in this section measure the amount for dispersion of each SC species measured relative to its mean value. They represent SD standard deviations. This is now clarified in the main text.

(5) Figure 5 is not readable.

We are a bit surprised by this comment as heatmaps are commonly used in microbiology studies to facilitate the visual presentation and exploration of complex correlation patterns. If the comment is about the quality of the figure, we will provide the reviewer with a new figure with a much higher resolution for final publication.

(6) In lines 148-149, based on which factors (literature data etc.) was the OM/OC conversion factor of 1.8 used? This choice has to be well explained.

This value of 1.8 for the OM/OC ratio was chosen on the basis of previous studies

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carried out in France, for the purpose of spatial comparison. In a recent study, we performed a mass balance between PM10 chemistry and TEOM measurements where the conversion factor of 1.8 was found to be consistent with a correct reconstruction of the PM10 mass (Favez et al., 2010; Golly et al., 2018; Petit et al., 2015).

Our choice, as suggested by the reviewer, is now explained in the main text as follows:

Lines 158-159 : “This value of 1.8 for the OM/OC ratio was chosen on the basis of previous studies carried out in France (Samaké et al., 2019b, and reference therein)”.

Some (not all) minor comments:

(7) Line 16. It is should be “on rural area of France”

This has been changed in the main text (see line 16).

(8) Lines 37. References are missing.

Missing references have been added in the main text (see line 40-41).

(9) Line 119. Why PM10 cut was selected for sampling? Some PBOA have a larger size.

In European countries, including France, health alerts on particulate matter are based on measurements of particles less than 10 μm in diameter. More details on European Union particulate matter standards can be found elsewhere (Priemus and Schutte-Postma, 2009). Therefore, our study focused on understanding the PBOA in the PM10 fraction.

(10) Line 122. How the collected filters were stored prior analyses? (It has to be added to the experimental section).

This information has been added (see lines 131-132)

(11) Lines 143, 170, 172. Company’s city (state, country) is missing.

This information has been added (see lines 155, 183, 185)

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(12) Line 207 (and everywhere in the text). Words “bacteria” and “fungi” should not be capitalized

We thank the reviewer for this suggestion. However, according to the conventions of microorganisms nomenclature, the name of the microbiome phylum should generally begin with a capital letter.

(13) Figure 6. Explain what black diamonds represent in this figure.

Thank you for this suggestion. A detailed explanation has now been added (see line 432-434).

(14) Line 439. The space should be removed before "."

This extra space has been removed.

(15) Line 445. Use OM instead of “organic matter”.

This has been changed in the main text.

(16) Line 477. Use SC not “sugar compounds”

This has been changed in the main text.

(17) Line 518. Remove extra “.”

This has been changed in the main text.

(18) Line 531. Use “strongly” instead of “highly”.

This has been changed in the main text.

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