

Response to comments by editor

Dear Zhe-Xuan Zhang, I have received two independent reviews of your submitted manuscript, and have evaluated your response to their comments. In general, your replies to the reviewers' comments are satisfactory.

I still advise to address the reviewer comments better by doing the following:

1) Apply the BigMac model to your dataset, instead of doing an independent analysis. This will allow to i) test the BigMac model along an interesting environmental gradient, ii) have an additional line of evidence to assign provenance to your downstream estuary samples (L 409: soil or in-situ?), that is not skewed by your localized soil sampling scheme. This is needed for both aims of the paper 'environmental controls' and 'development of salinity proxy'.

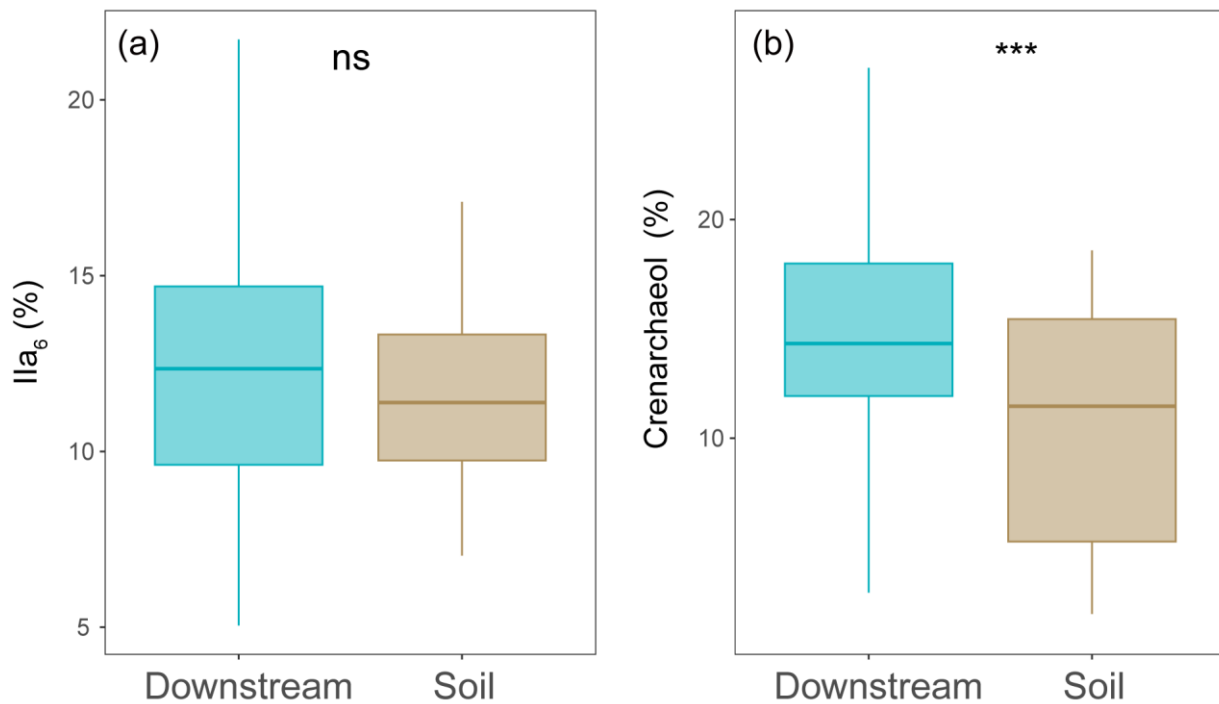
We would like to thank the editor for her valuable and constructive comments. A point-by-point reply to the comments is provided below and is colored blue. The text has been added into the revised manuscript is shown in orange italics. The line numbers correspond to those of the manuscript with tracked changes.

We have now applied the BigMac model to our GDGT dataset (including both isoGDGTs and brGDGTs). A figure and additional discussion have been added to the revised manuscript as follows (lines 529-557):

“In order to further assess whether downstream estuarine samples could be distinguished from soils, we applied the machine learning model (BigMac) developed by Martínez-Sosa et al. (2023) to our dataset with isoGDGT and brGDGT data as input. Most of our samples (SPM, sediments, and soils) were predicted as lake-type, with only one soil sample (soil6) collected at site B predicted as soil-type. This model suggests that, when considered altogether, the isoGDGT and brGDGT distributions are similar in aquatic and soil samples from the Seine estuary and differ from the soil-type samples described by Martínez-Sosa et al. (2023). Since the BigMac model does not include a river-type or estuary-type category (Martínez-Sosa et al., 2023), further inclusion of both isoGDGT and brGDGT data from global riverine/estuarine samples in the BigMac model may help enhance predictions for river-type or estuary-type SPM and sediment samples.

The BigMac model distinguishes the type of samples using $I\alpha_6$ and crenarchaeol as the two most important predicting variables. When accounting for both isoGDGTs and brGDGTs in the Seine River basin, the fractional abundance of crenarchaeol vs. total GDGTs (i.e. isoGDGTs + brGDGTs) varies significantly, whereas the one of $I\alpha_6$ does not differ significantly between the downstream estuary and soils (Fig. S8). Hence, the inclusion of isoGDGTs in the model may highly reduce the differences between sample types, as we observe significant differences in the fractional abundance of $I\alpha_6$ when calculated vs. total brGDGTs only (Fig. 3). As the BigMac model relies on both isoGDGT and brGDGT distribution, with no option of using brGDGTs alone, we chose to perform an independent analysis to assess the similarity in brGDGT relative abundance between downstream SPM and sediment samples on the one hand and soils from the Seine River basin on the other hand. This model was developed using the same algorithm (random forest) as Martínez-Sosa et al. (2023). Binary classification (downstream estuary vs. soils) was performed based on fractional abundances of brGDGTs. The trained model (Fig. S9) indicated distinguishable brGDGT distributions between downstream estuary (SPM and sediments) and soil samples, supporting the in situ production of brGDGTs in the downstream estuary. Although most of our soil samples were collected from the downstream estuary and showed similarity with the downstream

45 *SPM and sediment samples through PCA and comparison of fractional abundances, we were able to*
46 *distinguish their brGDGT compositions using machine learning.”*
47



48
49 *Fig. S8. Relative abundance of Ila₆ (a) and crenarchaeol (b) over 19 GDGTs (GDGT-0, GDGT-1,*
50 *GDGT-2, GDGT-3, Crenarchaeol, Crenarchaeol', IIIa₅, IIIa₆, IIIb₅, IIIb₆, IIa₅, IIa₆, IIb₅, IIb₆, IIc₅,*
51 *IIc₆, Ia, Ib, and Ic) used in the BigMac model. Boxes show the upper and lower quartiles of the data,*
52 *and whiskers show the range of the data, which are color-coded based on the sample type (downstream*
53 *estuary in blue and soil in brown). The center-line in the boxes indicates the median value of the dataset.*
54 *Statistical testing was performed by a Wilcoxon test (**p < 0.01; ***p < 0.001; ns, not significant, p > 0.05).*
55

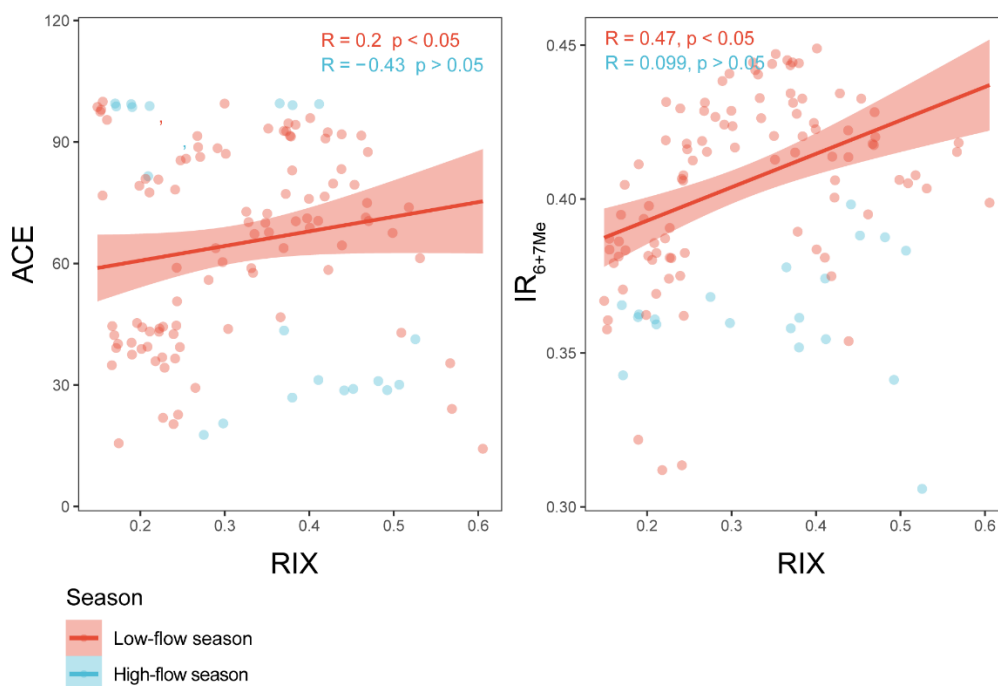
56
57 I agree with reviewer 2 that the similarity in distribution of brGDGTs present in soils and downstream
58 estuary sediments can not be determined based on the PCA (L 296: your PCA is done correctly, but only
59 reflects a part of the variance, as thus does not allow a straightforward comparison). Please compared
60 fractional abundances or a set of ratios that summarizes the GDGT variability.

61 We agree with the editor that the PCA alone cannot determine the similarity of brGDGT distribution
62 between soils and downstream samples. We have also compared the brGDGT fractional abundances, and
63 included additional discussion in the revised manuscript as follows (lines 519-523):

64 *“Additionally, no significant differences were observed in the fractional abundances of several brGDGTs*
65 *(IIIb₆, IIb₆, IIc₆, IIIa₇, IIa₇, 1050d, IIIa₅, IIIb₅, IIIb₇, IIIc₅, IIc₆, and Ia) between soils and downstream*
66 *samples (Fig. 3 and Fig. S4). This similarity in brGDGT distributions may be due to the influx of brGDGTs*
67 *from the downstream soils into the downstream estuary, as 82% of the soils were collected downstream*
68 *(Fig. 1a and Table 1).”*
69

70 2) Please compare your RIX values directly with the IR6+7ME and the ACE values (not just ACE and
71 IR6+7ME values vs salinity).

72 Thank you for this suggestion. We added a figure in Supplementary material (Fig. S13) showing the
73 correlations between the RIX and ACE on the one hand and RIX and IR_{6+7Me} on the other hand:



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75 *Fig. S13. RIX plotted versus ACE and IR_{6+7Me} through the linear regression. Shaded area represents*
76 *95% confidence intervals. Dataset is composed of SPM.*

78 Additional discussion was added to the revised manuscript as follows (lines 701-715):

79 “Since the other salinity proxies (i.e. ACE and IR_{6+7Me}) have shown positive correlations with salinity in
80 previous studies (Turich and Freeman, 2011; Wang et al., 2021), they were expected to be positively
81 correlated with salinity and negatively correlated with RIX in the Seine River basin. However, the ACE
82 index (Turich and Freeman, 2011) and IR_{6+7Me} (Wang et al., 2021) do not show significant correlations
83 with salinity in the Seine River basin ($p > 0.05$, Wilcoxon test; Fig. S10) and show weak but significant
84 relations with the RIX (Fig. S13). This could be attributed to the influence of other factors than salinity on
85 these indices (i.e. ACE and IR_{6+7Me}). Indeed, while ACE has been successfully applied in hypersaline
86 systems (Turich and Freeman, 2011), it performs less effectively in certain saline settings due to the
87 complex sources of archaeol and GDGTs (Huguet et al., 2015) and/or distinct ionization efficiencies
88 between these compounds (He et al., 2020; Wang et al., 2021). Similarly, IR_{6+7Me} may be influenced by
89 the preferential production of 6-methyl brGDGTs related with nitrogen nutrient loadings in a specific
90 region of the estuary (KP 255.6-337), as discussed in 4.1.2. Consequently, only RIX successfully tracks
91 salinity variations in this basin, while ACE and IR_{6+7Me} show relative insensitivity.”

92
93 3) Comment on the potential to do a quantitative reconstruction of salinity based on the RIX index, and
94 whether the Seine estuary is a good location to develop this index. An additional few words on the
95 potential impact of soil-derived brGDGTs would be beneficial (L601).

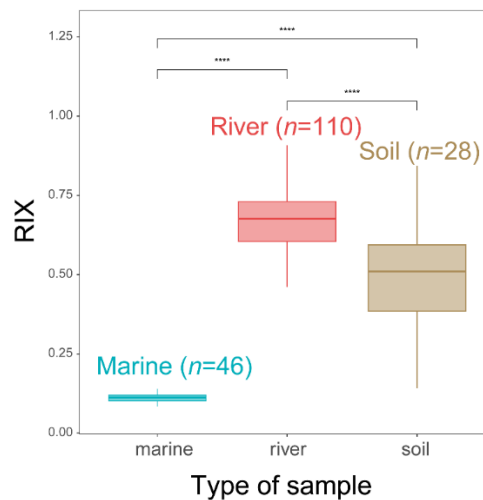
96 Thank you for this comment. In the current manuscript, we introduced RIX as a proxy for riverine runoff
97 (river-derived organic matter). We have indeed observed significant correlations between salinity and
98 various brGMGTs, with different isomers showing distinct behaviors in response to salinity changes. This
99 observation forms the basis and rationale for our RIX index. However, a quantitative reconstruction of

100 salinity using this index needs further exploration. Specific suggestions for this future work and an
101 assessment of whether the Seine Estuary is a suitable location for salinity calibration have been included
102 in the revised manuscript as follows (lines 715-717):

103 *“However, quantitatively reconstructing salinity with RIX is an important step forward that warrants*
104 *further investigation. This requires comparing brGMGT distributions from various aquatic*
105 *environments (e.g. estuaries and lakes) across salinity gradients.”*

106
107 Additionally, we assume that the editor referred to soil-derived brGMGTs (not brGDGTs) here (L601).
108 The RIX values in soils were compared with those from river, upstream estuarine, and downstream
109 estuarine samples. Our findings indicate that RIX values in soils are close to those in downstream
110 estuarine samples and are lower than those in river and upstream estuarine samples. This suggests that
111 potential soil contributions would dilute the riverine brGMGT signal, further decreasing RIX. Such
112 potential soil effects (dilution of riverine signal) in the Seine River basin are also observed in the Bay
113 of Bengal. However, given the differences in distributions between soil and aquatic samples, as well as
114 the lower brGMGT concentration in soils, soils may have only a limited influence on the value of RIX.

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116 Fig. 9 was modified to include soils from the Godavari basin:



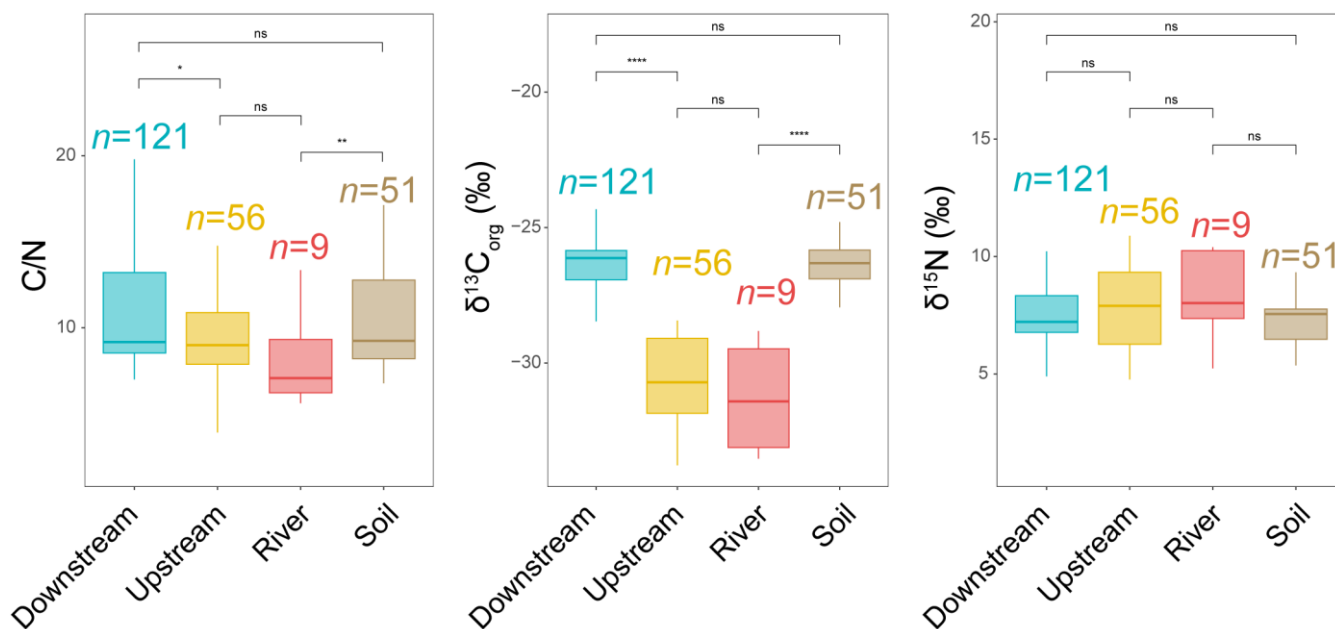
117 *Figure 9: RIX in the soils, SPM and sediment samples from Godavari River basin (India) and Bay of*
118 *Bengal sediments (data from Kirkels et al. (2022a)). Statistical testing was performed by a Wilcoxon*
119 *test. Boxes show the upper and lower quartiles of the data, and whiskers show the range of the data,*
120 *which are color-coded based on the sample type (river in red, marine in blue, and soil in brown). The*
121 *center-line in the boxes indicates the median value of the dataset.*

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123
124 Additional discussion was included in the revised manuscript as follows (lines 794-798):

125 *“RIX values in soils (0.49 ± 0.16) around the Godavari River basin are significantly lower than those of*
126 *the river samples ($p < 0.05$, Wilcoxon test; Fig. 9). Therefore, the potential soil contribution would dilute*
127 *the riverine brGMGT signal, further decreasing RIX in marine sediments. This is consistent with the*
128 *observations in the Seine River basin. However, given the distinct distributions between soil and aquatic*
129 *samples and the lower brGMGT concentration in soils (Kirkels et al., 2022a), the influence of soil-derived*
130 *brGMGTs on riverine RIX values may be limited.”*

131
132 My own comments:

133 1) Based on the study at the Seine River, do the authors propose that this proxy traces salinity, terrestrial
 134 organic matter or river-derived organic matter?
 135 Thank you for this comment. We have addressed it above. Please kindly refer to our earlier response.
 136
 137 2) Could you use C/N instead of TOC and TN as a commonly used geochemical proxy for soil input?
 138 In the revised manuscript, we have replaced the boxplot of TOC and TN by a boxplot of C/N:



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 140 *Figure 2: Distribution of bulk parameters (C/N, $\delta^{13}C_{org}$ and $\delta^{15}N$) from soils (surficial soils and mudflat*
 141 *sediments) as well as river, upstream estuary and downstream estuary samples across the Seine River*
 142 *basin. Box plots of upstream and downstream estuary samples are based on SPM and sediments, whereas*
 143 *those of river samples are based only on SPM. Boxes show the upper and lower quartiles of the data, and*
 144 *whiskers show the range of the data, which are color-coded based on the sample type (river in red,*
 145 *upstream estuary in yellow, and downstream estuary in blue). The center-line in the boxes indicates the*
 146 *median value of the dataset. Statistical testing was performed by a Wilcoxon test (* $p < 0.05$; ** $p < 0.01$;*
 147 **** $p < 0.001$; **** $p < 0.0001$; ns, not significant, $p > 0.05$).*

148
 149 These data are described in the result section of revised manuscript (lines 339-341):
 150 “Lower C/N values were observed in the river (8.04 ± 4.31 , based on SPM) and upstream estuary
 151 (9.42 ± 3.67 , based on SPM and sediments) compared to the downstream estuary (10.73 ± 3.59 , based on
 152 SPM and sediments) and soils (11.59 ± 4.79 , based on surficial soils and mudflat sediments) (Fig. 2).”

153
 154 Additional discussion is provided as follows (lines 810-820):
 155 “It is worth noting that another terrestrial proxy (C/N) was not included because it may be ineffective in
 156 tracing terrestrial OM in this anthropogenic estuary. The C/N ratio is commonly used as a bulk indicator
 157 of terrestrial OM, with higher values indicating a greater terrestrial contribution than marine sources
 158 (Bianchi and Canuel, 2011). However, other parameters such as decomposition processes,
 159 remineralization, and distinct sources could complicate its application (Lamb et al., 2006). In the Seine
 160 River basin, C/N values were significantly lower in the river and upstream estuary than in downstream
 161 samples (Fig. 2). Given that anthropogenic OM contains more nitrogen than natural OM, an increase in

162 *anthropogenic sources would result in a decrease in C/N values (Van Den Hende et al., 2011; Liu et al.,*
163 *2020). As a result, the observed decrease in C/N values in river and upstream estuarine samples could be*
164 *attributed to a higher contribution of nitrogen from anthropogenic sources, possibly due to intense*
165 *agricultural activities as discussed in 4.1.2. As BIT, RIX, and $\delta^{13}\text{C}_{\text{org}}$ provide similar information, they*
166 *may be more reliable tracers of terrestrial OM compared to C/N in the Seine River basin.”*

167
168 3) Can you constrain for the Bay of Bengal at all what are the RIX values of the soils are? How would a
169 change in soil input impact the RIX values, and does this complicate the interpretation of the RIX index
170 as a salinity tracer? Even if it is a minor process in the Seine estuary, it might be an important driver of
171 downcore changes in the Bay of Bengal?

172 Thank you very much for your comment, which was addressed above.

173
174 Minor comments:

175 Fig. 3: Please include what compound the name “1036d” refers to. 7-methyl brGDGTs are not included
176 in the Fig.S1, please update. Specify that for some compounds the structure has not been described yet.

177 Thank you for your comment. To date, the structures of 1050d and 1036d have not been described.
178 Therefore, we tentatively refer to these compounds as 1050d and 1036d. We have added related
179 information into the caption of Fig. 3 (which has now been moved to Fig. S2 as suggested by reviewer
180 #2) as follows (lines 944-945):

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182 *“1050d and 1036d represent compounds eluting later than IIIa₇ and IIa₇, respectively.”*

183
184 Additionally, the structures of 7-methyl brGDGTs (i.e., IIIa₇, IIa₇, IIIb₇) have been included in Fig. S1.
185 We have also specified that, for some compounds eluting later than 7-methyl brGDGTs, their structures
186 have not been described yet. The relevant information has been added to the caption of Fig. S1 as follows
187 (lines 939-940):

188
189 *“Note that the structures of brGMGTs and compounds eluting later than 7-Methyl brGDGTs (1050d and*
190 *1036d) have not been described.”*

191
192 L 340: Were brGMGTs present in all samples, or did some compounds fall below detection level in a
193 certain samples type? Please include this description.

194 Some of the brGMGTs, especially H1034a, are below detection level in most of the SPM and sediment
195 samples. We have now included related information in the revised manuscript as follows (lines 425-426):

196
197 *“H1034a is the least abundant isomer and is below detection limit for most of the SPM and sediment*
198 *samples in the Seine River basin.”*

In this manuscript, Zhang et al propose a new proxy to reconstruct fluvial organic matter inputs to coastal marine settings. They suggest that brGMGTs are produced in-situ in rivers and estuaries and that the distribution of brGMGTs is principally controlled by salinity. Based on these facts they generate a new Riverine Index (RIX) using the fractional abundances of H1020c and H1034b versus H1020a and H1020b. To validate the RIX in deep time they compare RIX values to the BIT index and terrestrial pollen/spore deposits deposited during the PETM from IODP Expedition 302 Hole 4A. They report a closer relationship between RIX and terrestrial pollen abundance than BIT and terrestrial pollen abundance, indicating that at least in this site RIX outperforms BIT in accurately reconstructing riverine inputs. In all, this is an interesting study that will likely be of interest to BG readers. I have a number of comments that aim to strengthen the manuscript.

We would like to thank the reviewer for all the constructive comments and the positive assessment on the significance of our manuscript. A point-by-point reply to all the reviewer's comment is provided below and is colored blue. The text has been added into the revised manuscript is shown in orange italics. The line numbers correspond to those of the manuscript with tracked changes.

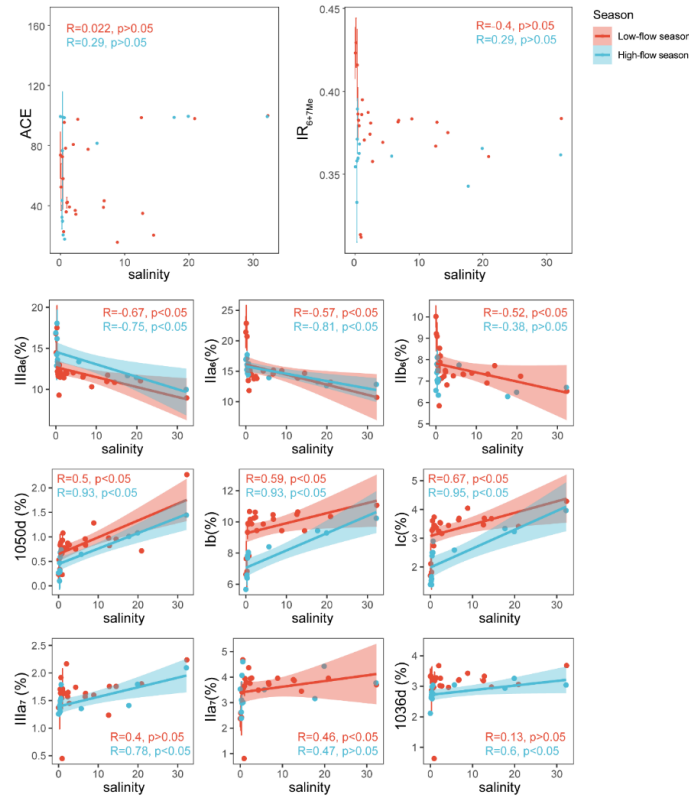
General comments:

In some sections (see specific comments) the use of English in the paper is poor and obfuscates the meaning of the text. I suggest that the authors carefully read through the manuscript to catch all typos and grammatical errors. Likewise figure quality varies considerably. In some cross plots, it is impossible to see the data as the marker size is so small (see specific comments). Characters that should be superscripted/subscripted are left as regular text (see specific comments). Lines of best fit are drawn through data but there is no information as to how these lines were constructed (see specific comments). As such, this manuscript would benefit greatly from more attention to detail from the authors.

We have carefully checked the manuscript for any typos and grammatical errors to improve its readability and clarity. Regarding the figure quality, we have addressed all the issues highlighted by the reviewer, including marker size, superscripting/subscripting, and provided detailed information (e.g. lines of best fit) in the figure caption.

Additionally, as the authors are proposing a new GDGT salinity index, I would like them to calculate and report previously formulated salinity indices from their samples. Specifically, the ACE index (Turich and Freeman 2011) and the IR_{6+7me} (Wang et al 2021) both have been calibrated against water salinity in marine saline ponds and hypersaline lakes respectively. I know the author's brief touched on comparing IR_{6me} from this study to values from Wang et al (2021) in the text but a more thorough examination of prior GDGT-derived water salinity reconstructions would strengthen the manuscript. Readers will be interested to see how these indices compare against RIX in reconstructing salinity from an estuarine environment.

We agree with the suggestion by the reviewer. We have calculated the ACE and IR_{6+7me} indices and have presented them in a supplementary figure (presented below), allowing the comparison of the corresponding values with those of the RIX. The ACE and IR_{6+7me} indices do not correlate with salinity in the Seine River basin.



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Fig. S10. Salinity plotted versus ACE, IR_{6+7Me} , relative abundance of 6-methyl and 7-methyl brGDGTs ($IIIa_6$, Ila_6 , Iib_6 , $IIIa_7$ and Ila_7) as well as compounds 1050d, 1036d, Ib, and Ic through the linear regression. Shaded area represents 95% confidence intervals. Vertical error bars indicate mean \pm s.d for samples with the same salinity. Dataset is composed of SPM.

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We have added the following text in the revised manuscript (lines 701-715):

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“Since the other salinity proxies (i.e. ACE and IR_{6+7Me}) have shown positive correlations with salinity in previous studies (Turich and Freeman, 2011; Wang et al., 2021), they were expected to be positively correlated with salinity and negatively correlated with RIX in the Seine River basin. However, the ACE index (Turich and Freeman, 2011) and IR_{6+7Me} (Wang et al., 2021) do not show significant correlations with salinity in the Seine River basin ($p > 0.05$, Wilcoxon test; Fig. S10) and show weak but significant relations with the RIX (Fig. S13). This could be attributed to the influence of other factors than salinity on these indices (i.e. ACE and IR_{6+7Me}). Indeed, while ACE has been successfully applied in hypersaline systems (Turich and Freeman, 2011), it performs less effectively in certain saline settings due to the complex sources of archaeol and GDGTs (Huguet et al., 2015) and/or distinct ionization efficiencies between these compounds (He et al., 2020; Wang et al., 2021). Similarly, the IR_{6+7Me} may be influenced by the preferential production of 6-methyl brGDGTs related with nitrogen nutrient loadings in a specific region of the estuary (KP 255.6-337), as discussed in 4.1.2. Consequently, only the RIX successfully tracks salinity variations in this basin, while ACE and IR_{6+7Me} show relative insensitivity.”

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Additionally, the evidence for in situ production of brGDGTs and brGMGTs in downstream estuary sites is pretty weak. This is demonstrated by Fig 2 where we see that distributions of $d_{13}C_{org}$ and $d_{15}N$ in soils and downstream estuary samples are very similar in addition to Fig 5 where your PCA on sample brGDGT and brGMGT distributions cannot separate out soils from downstream estuary samples. Yes you

269 see (on average) higher concentrations of brGDGTs and brGMGTs in downstream estuary samples than
270 in soils but the actual distributions of brGDGT and brGMGT abundance in soils are pretty large, indicating
271 that some soils have pretty substantial quantities of these compounds. A great way to add more clarity to
272 this sourcing issue is to train a random forest model using a similar method to Martínez-Sosa et al (2023)
273 on your brGDGT and brGMGT samples (and isoGDGTs as these should be available to you). If the
274 random forest model can accurately separate out soils from downstream estuary samples then you can be
275 pretty sure that your downstream estuary samples were produced in situ. This won't require much
276 additional work and can be implemented easily using python (<https://scikit-learn.org/>) or another language
277 of your choice.

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279 We thank the reviewer for this suggestion. As suggested by the editor, we have applied first the BigMac
280 model (based on isoGDGTs and brGDGTs). However, the inclusion of isoGDGTs in the Seine River
281 basin may highly reduce differences between sample types. Consequently, we used independent models
282 for brGDGTs and brGMGTs, respectively. Please see our response to the editor (lines 16-54 in this
283 response letter).

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285 The application of this model to brGDGT and brGMGT datasets accurately separates downstream (SPM
286 and sediment) estuarine samples from soil ones, indeed supporting *in situ* production of these lipids in
287 downstream Seine Estuary.

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289 In the material and methods, we added a new machine-learning section describing the model, as follows
290 (Lines 307-319):

291 *“The BigMac model, developed by Martínez-Sosa et al. (2023) based on brGDGT and isoGDGT*
292 *distribution, was applied. Subsequently, using the same algorithm (random forest), we developed our own*
293 *model based on either brGDGTs or brGMGTs.*

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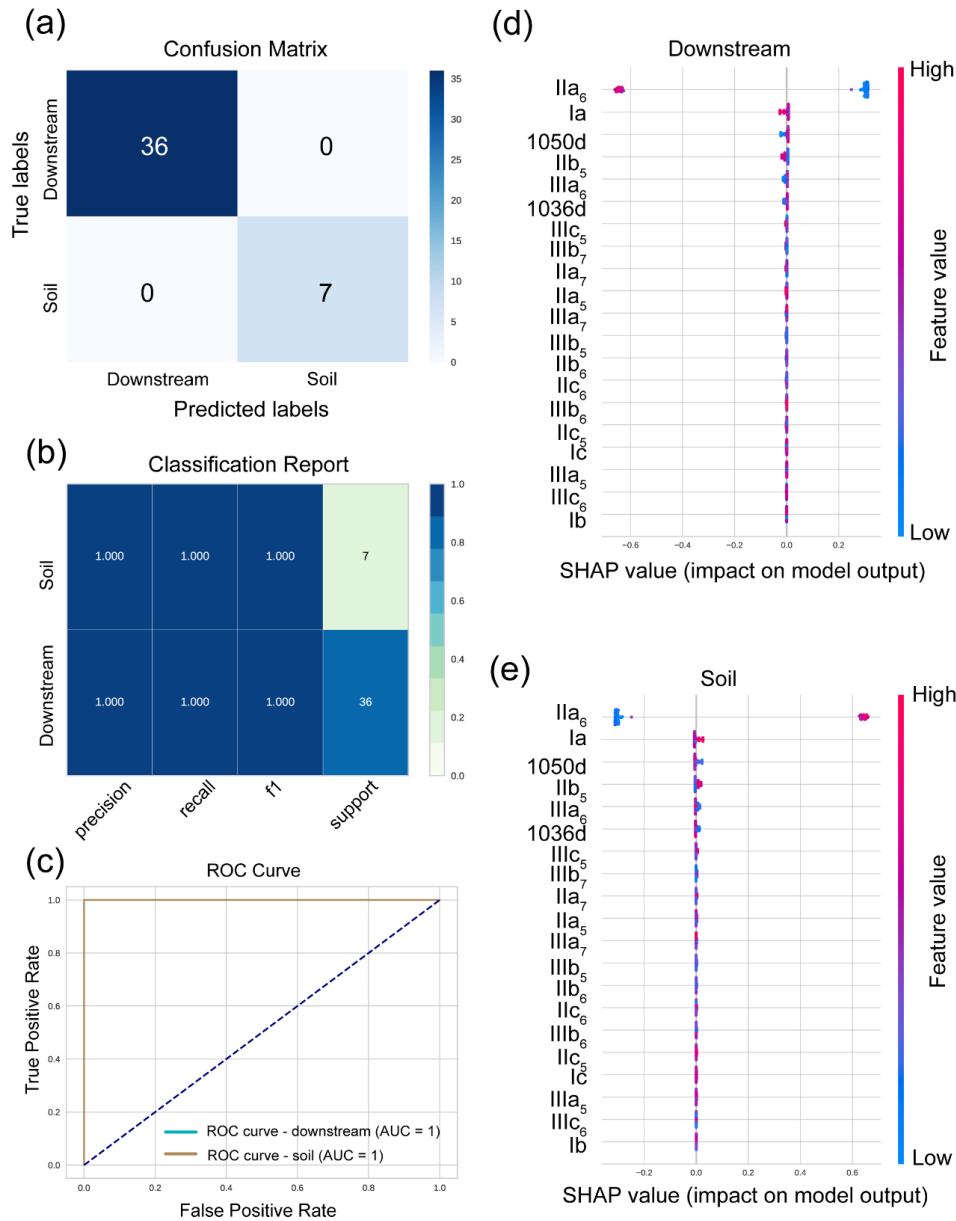
295 *For independent models, our lipid dataset was split into a training set (75%) and a test set (25%). We*
296 *then used a supervised machine-learning algorithm (random forest) to train models. This algorithm was*
297 *applied to classify the downstream estuary and soil samples based on brGDGTs or brGMGTs as input,*
298 *implemented using the scikit-learn library (<https://github.com/scikit-learn/>) (Pedregosa et al., 2011) in*
299 *Python (version 3.10.12). Hyperparameter tuning was conducted using a randomized search approach*
300 *implemented through the RandomizedSearchCV function in scikit-learn.*

301

302 *SHapley Additive exPlanations (SHAP) is a game-theoretical method used to interpret machine learning*
303 *models (Lundberg et al., 2020). SHAP analysis was applied to identify which compounds were important*
304 *for the classifications, implemented by the SHAP library in Python. A higher SHAP value indicates a*
305 *more substantial contribution of the feature (brGDGTs or brGMGTs) to the predicted outcome*
306 *(downstream estuary or soils).”*

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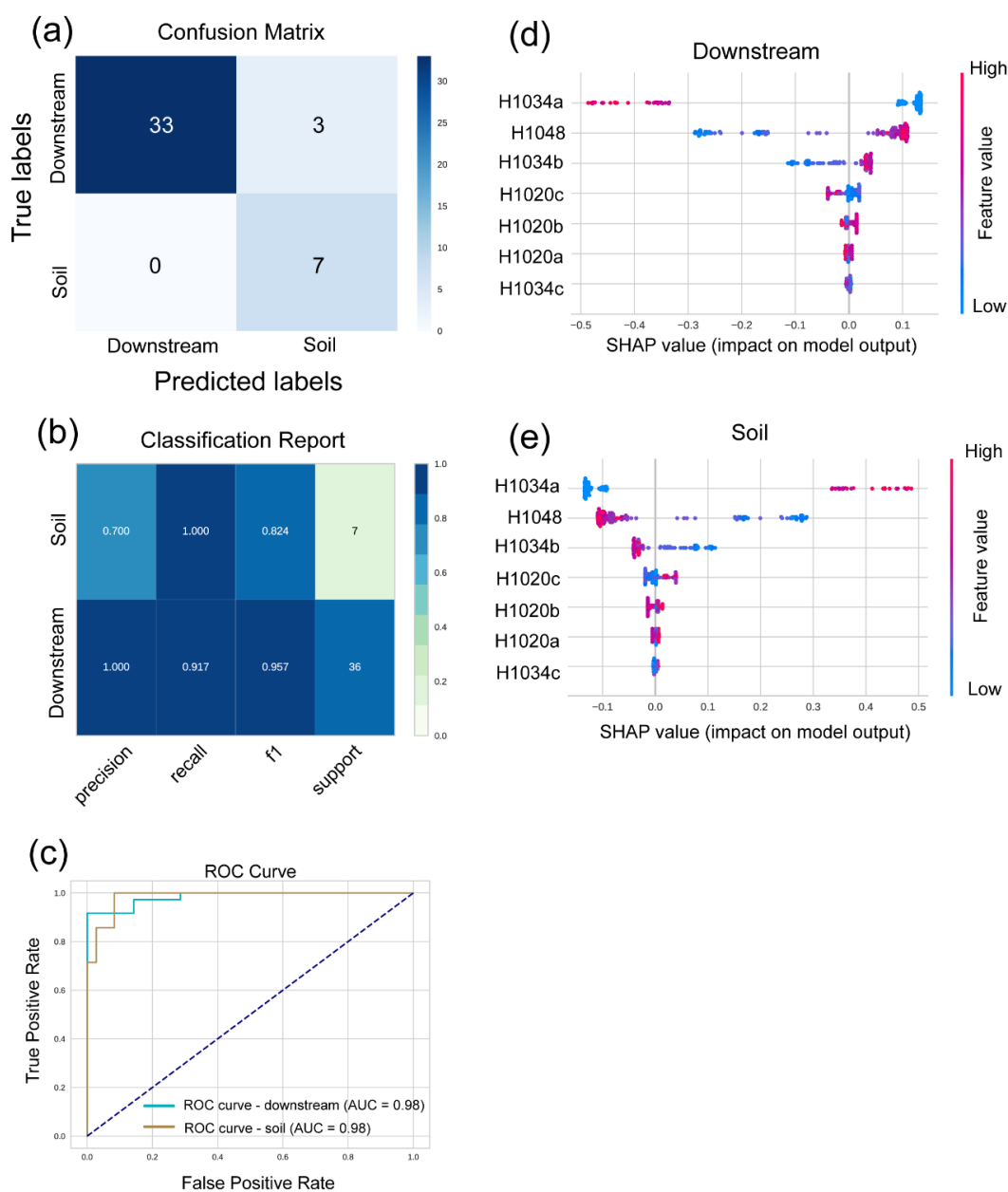
308 Two figures (one for brGDGTs, another for brGMGTs) showing the performance of the model have been
309 added to the supplement:



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311 *Fig. S9 (brGDGTs). Evaluation of the random forest model based on brGDGTs through the confusion*
 312 *matrix (a), classification report (b), and receiver operating characteristic (ROC) curve (c). SHAP*
 313 *summary plots (d-e) show the feature importance obtained from the random forest algorithm and the*
 314 *SHAP library. Each bullet in the plot represents a single sample in the training set, with the color*
 315 *indicating the feature value (fractional abundance of the brGDGTs) from low (blue) to high (pink). The*
 316 *bullets positioned on the right side of the SHAP summary plot correspond to positive SHAP values,*
 317 *indicating a positive effect on the model output (downstream estuary or soils). The bullets on the left side*
 318 *of the plot indicate negative SHAP values, suggesting a negative effect on the model output. The variables*
 319 *(brGDGTs) with higher impact on the model performance are shown at higher positions. Training sets*
 320 *include downstream SPM and sediment samples (d) and soils (e).*

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Fig. S12 (brGMGTs). Evaluation of the random forest model based on brGMGTs through the confusion matrix (a), classification report (b), and receiver operating characteristic (ROC) curve (c). SHAP summary plots (d-e) show the feature importance obtained from the random forest algorithm and the SHAP library. Each bullet represents a single sample within the training set, with the color representing the feature value (fractional abundance of the brGMGTs) ranging from low (blue) to high (pink). The bullets positioned on the right side of the SHAP summary plot correspond to positive SHAP values, indicating a positive effect on the model output (downstream estuary or soils). The bullets on the left side of the plot indicate negative SHAP values, suggesting a negative effect on the model output. The variables (brGDGTs) with higher impact on the model performance are shown at higher positions. The training sets include downstream SPM and sediment samples (d) as well as soils (e).

We added the following text in a revised manuscript to describe and discuss the results related to the application (i) of the Bigmac model to the brGDGT and isoGDGT dataset and (ii) our independent model

337 applied to the brGDGT dataset (lines 529-557):

338 *“In order to further assess whether downstream estuarine samples could be distinguished from soils, we*
339 *applied the machine learning model (BigMac) developed by Martínez-Sosa et al. (2023) to our dataset*
340 *with isoGDGT and brGDGT data as input. Most of our samples (SPM, sediments, and soils) were*
341 *predicted as lake-type, with only one soil sample (soil6) collected at site B predicted as soil-type. This*
342 *model suggests that, when considered altogether, the isoGDGT and brGDGT distributions are similar in*
343 *aquatic and soil samples from the Seine estuary and differ from the soil-type samples described by*
344 *Martínez-Sosa et al. (2023). Since the BigMac model does not include a river-type or estuary-type*
345 *category (Martínez-Sosa et al., 2023), further inclusion of both isoGDGT and brGDGT data from global*
346 *riverine/estuarine samples in the BigMac model may help enhance predictions for river-type or estuary-*
347 *type SPM and sediment samples.*

348
349 *The BigMac model distinguishes the type of samples using Ila₆ and crenarchaeol as the two most*
350 *important predicting variables. When accounting for both isoGDGTs and brGDGTs in the Seine River*
351 *basin, the fractional abundance of crenarchaeol vs. total GDGTs (i.e. isoGDGTs + brGDGTs) varies*
352 *significantly, whereas the one of Ila₆ does not differ significantly between the downstream estuary and*
353 *soils (Fig. S8). Hence, the inclusion of isoGDGTs in the model may highly reduce the differences between*
354 *sample types, as we observe significant differences in the fractional abundance of Ila₆ when calculated*
355 *vs. total brGDGTs only (Fig. 3). As the BigMac model relies on both isoGDGT and brGDGT distribution,*
356 *with no option of using brGDGTs alone, we chose to perform an independent analysis to assess the*
357 *similarity in brGDGT relative abundance between downstream SPM and sediment samples on the one*
358 *hand and soils from the Seine River basin on the other hand. This model was developed using the same*
359 *algorithm (random forest) as Martínez-Sosa et al. (2023). Binary classification (downstream estuary vs.*
360 *soils) was performed based on fractional abundances of brGDGTs. The trained model (Fig. S9) indicated*
361 *distinguishable brGDGT distributions between downstream estuary (SPM and sediments) and soil*
362 *samples, supporting the in situ production of brGDGTs in the downstream estuary. Although most of our*
363 *soil samples were collected from the downstream estuary and showed similarity with the downstream*
364 *SPM and sediment samples through PCA and comparison of fractional abundances, we were able to*
365 *distinguish their brGDGT compositions using machine learning.”*

366
367 We also added the following text to describe and discuss the results related to the application of the model
368 to the brGMGT dataset (lines 728-734):

369 *“As with brGDGTs, we applied a random forest algorithm to distinguish brGMGT distributions between*
370 *downstream estuary and soil samples. This trained model accurately distinguishes soils from downstream*
371 *estuarine samples (Fig. S12), indicating in situ production of brGMGTs in the downstream estuary. Given*
372 *the significantly low brGMGT concentrations in soils ($p < 0.05$, Wilcoxon test; Fig. S5b) and the distinct*
373 *distributions between brGMGT in soils and aquatic settings identified through PCA (Fig. 4) and machine*
374 *learning (Fig. S12), it can be assumed that the impact of soil-derived brGMGTs on the observed RIX*
375 *signal in the water column of the Seine basin is low.”*

376

377

378 **Specific comments**

379

380 **Line 35: This complicates paleoenvironmental interpretations in SOME aquatic settings not ALL aquatic**

381 settings

382 We agree with this suggestion, this has been corrected.

383

384 Line 37: “all along this basin, from land to sea” awkward phrasing

385 We have rephrased this sentence as follows (lines 37-38):

386

387 *“BrGDGTs and brGMGTs were analyzed in soils, Suspended Particulate Matter (SPM), and sediments*
388 *(n=237) collected along the land-sea continuum of the Seine basin.”*

389

390 Line 40: “Redundancy analysis further shows that both salinity and nitrogen loadings dominantly control
391 the brGDGT distributions.” No, the loadings indicate that SALINITY (not salinity loadings) controls the
392 brGDGT distribution.

393 This has been corrected.

394

395 Line 40-43: “Furthermore, the relative abundance of 6- methyl vs. 5-methyl brGDGTs (IR6Me ratio),
396 Total Nitrogen (TN), $\delta^{15}\text{N}$ and chlorophyll a concentration co-vary in a specific zone with low salinity”
397 Is this zone geographical, in your redundancy analysis, or something else?

398 This zone is geographical. We have added the following sentence in a revised manuscript (lines 43-46):

399 *“Furthermore, the relative abundance of 6-methyl vs. 5-methyl brGDGTs (IR_{6Me} ratio), Total Nitrogen*
400 *(TN), $\delta^{15}\text{N}$ and chlorophyll a concentration co-vary in a specific geographical zone with low salinity,*
401 *suggesting that 6-methyl brGDGTs are preferentially produced under low-salinity and high-productivity*
402 *conditions.”*

403

404 Line 44-45: “Salinity is positively correlated with homologs H1020a and H1020b, 45 and negatively
405 correlated with compounds H1020c and H1034b.” Is this in soils, sediments or SPM?

406 This correlation was found in SPM. This has been specified as follows (lines 47-48):

407 *“Salinity is positively correlated with homologues H1020a and H1020b, and negatively correlated with*
408 *compounds H1020c and H1034b in SPM.”*

409

410 Line 45: “This suggests that bacteria thriving...” thriving is not the correct word (carries implications of
411 a value judgment) replace with “living”.

412 This has been corrected.

413

414 Line 45-47: It seems like you aren’t mentioning results from soils and sediments, only from SPM? Or
415 maybe all your sediment samples are exclusively from rivers? The reader is unclear on this.

416 Correlation between salinity and lipid distribution is based on SPM samples. We have modified the
417 abstract as follows (lines 40-42):

418 *“Both types of compounds (i.e. brGDGTs and brGMGTs) are shown to be produced in situ, in freshwater*
419 *and saltwater, based on their high concentrations and distinct distributions in aquatic settings (SPM and*
420 *sediments) vs. soils.”*

421

422 Line 51-52: “a paleorecord across the upper Paleocene and lower Eocene,” You should name this record
423 and say where it is.

424 We have added the name (Arctic Coring Expedition) and location (Lomonosov Ridge) of this record.

425

426 Lines 51: “showing its potential applicability in both modern samples and in paleorecords.” Perhaps you
427 could evaluate its usage in both these cases e.g. - we successfully/unsuccessfully applied RIX in ...

428 We have rephrased this sentence as follows (lines 53-55):

429 *“We successfully applied RIX to the Godavari River basin (India) and a paleorecord across the upper*
430 *Paleocene and lower Eocene from the Arctic Coring Expedition at Lomonosov Ridge, showing its*
431 *potential applicability in both modern samples and in paleorecords.”*

432

433 Line 55: “, although some of them were attributed to the phylum Acidobacteria” Imprecise language.

434 Thank you for the comment. We have rephrased this sentence into (lines 58-60):

435 *“Branched glycerol dialkyl glycerol tetraethers (brGDGTs) are membrane lipids produced by bacteria,*
436 *some of them belonging to the phylum Acidobacteria.”*

437

438 Line 57-58: “The distribution of brGDGTs (number of cyclopentane moieties and methyl groups; cf.
439 structures in Fig. S1) was empirically linked with pH and Mean Annual Air Temperature” Again,
440 imprecise language. The phrase “empirically linked” doesn’t convey much useful information.

441 We have replaced “empirically” by “has been” (lines 64-66):

442 *“The distribution of brGDGTs (number of cyclopentane moieties and methyl groups; cf. structures in Fig.*
443 *S1) has been linked with pH and Mean Annual Air Temperature (MAAT) in soils.”*

444

445 Line 60: Should really cite some earlier lake GDGT papers in addition to Martinez-Sosa et al., 2021.

446 Thank you for this suggestion. We have cited some earlier work: Tierney et al., 2010 GCA and Russell
447 et al., 2018 OG.

448

449 Lines 60-61: “The brGDGT-based proxies (i.e. MBT’₅ME and CBT’) have been largely applied to
450 reconstruct MAAT and pH from sedimentary archives (Coffinet et al., 2018; Harning et al., 2020; Wang
451 et al., 2020).” Not quite true - Martinez-Sosa et al (2021) and Dearing Crampton-Flood et al (2020)
452 generated Bayesian linear regressions between the Mean temperature of months above freezing and
453 MBT’₅me. These BayMBT models have been used widely in the community since their publication.

454 We agree that the new models were not applied to records available before their publication. We have
455 modified this sentence as follows (lines 64-66):

456 *“The brGDGT-based proxies (i.e. MBT’₅ME and CBT’) have been largely applied to reconstruct*
457 *paleoclimate from sedimentary archives (Coffinet et al., 2018; Harning et al., 2020; Wang et al., 2020).”*

458

459 Line 62-63: “In aquatic settings, brGDGTs were initially suggested to be predominantly derived from
460 watershed soils and transported by erosion in the sediments (Hopmans et al., 2004).” Maybe you mean
461 “transported by erosion to the sediments”?

462 We agree and have rephrased this sentence as follows (line 68):

463 *“In aquatic settings, brGDGTs were initially suggested to be transported by erosion to the sediments.”*

464

465 Lines 63-78: The use of English throughout this paragraph is poor and hard to follow. Needs copyediting.

466 We have rephrased this paragraph as follows (lines 68-89):

467 *“In aquatic settings, brGDGTs were initially suggested to be transported by erosion to the sediments*
468 *(Hopmans et al., 2004). Based on this assumption, the Branched and Isoprenoid Tetraethers (BIT) index*

469 was defined as the abundance ratio of the major brGDGTs to crenarchaeol (isoprenoid GDGT mainly
470 produced by marine Nitrososphaerota). The BIT index ranges between 0 and 1, with high BIT values
471 (around 1) reflecting a higher contribution of terrestrial organic matter compared to marine organic
472 matter (Hopmans et al., 2004).

473 Over the last few years, the BIT index has been broadly used to quantify the relative contribution of
474 terrestrial organic matter in aquatic systems (Xu et al., 2020; Yedema et al., 2023) and to evaluate the
475 reliability of the TEX₈₆ palaeothermometer (Cramwinckel et al., 2018). However, several studies have
476 shown that brGDGTs can also be produced in situ in aquatic settings, including rivers (e.g., De Jonge et
477 al., 2015; Freymond et al., 2017; Kim et al., 2015; Zell et al., 2014, 2013), lakes (Tierney and Russell,
478 2009), and marine environments (Dearing Crampton-Flood et al., 2019; Zeng et al., 2023). This adds
479 complexity to the identification of brGDGT sources in aquatic ecosystems and to the application of
480 brGDGTs as (paleo)environmental proxies, including the BIT index.

481 The BIT values have all the more to be carefully interpreted, especially considering the potential influence
482 of the selective degradation of branched vs. isoprenoid GDGTs (Smith et al., 2012). Thus, complementary
483 molecular proxies for quantifying the input of terrestrial organic matter to aquatic settings are still
484 needed. These proxies may cross-validate other available terrestrial proxies, such as the $\delta^{13}\text{C}$ of organic
485 carbon (Lamb et al., 2006), heterocyst glycolipids (Kang et al., 2023), and long-chain diols (Lattaud et
486 al., 2017).”

487
488 Lines 63-78: You should read and cite Martinez-Sosa et al (2023) here for their work on a Random Forest
489 approach to classify GDGT sources (i.e. Marine, Soil, Lake etc).

490 *Thank you for this suggestion. We have referred to the work by Martínez-Sosa et al. (2023) as follows*
491 *(lines 88-93):*

492 *“Recently, a machine-learning approach (BIGMaC model) was proposed to infer the origin of*
493 *environmental samples (e.g. soil, peat, marine and lake settings) based on their GDGT distribution*
494 *(Martínez-Sosa et al., 2023). While such an approach shows potential for differentiating distinct sources*
495 *of GDGTs, its application to aquatic systems has not yet been extensively explored.”*

496
497 Line 80-83: “The improvement of analytical methods allowed the separation and quantification of 5-, 6-
498 and 7-methyl brGDGTs (methyl groups at the fifth, sixth, and seventh positions; Fig. S1), that in previous
499 chromatographic protocols co-eluted (De Jonge et al., 2013, 2014; Ding et al., 2016).” No real link
500 between the previous paragraph and this one. Also, which methods? How did they improve?

501 *This comment has been taken into account as follows (lines 96-102):*

502 *“The improvement of chromatographic methods allowed the separation and quantification of 5-, 6- and*
503 *7-methyl brGDGTs (methyl groups at the fifth, sixth, and seventh positions; Fig. S1) that previously co-*
504 *eluted (De Jonge et al., 2013, 2014; Ding et al., 2016). This led to the development of new brGDGT-*
505 *based proxies based on these specific brGDGT isomers (De Jonge et al., 2014).”*

506
507 Lines 86-87: “In addition to temperature and pH, other environmental factors may influence brGDGT
508 distributions in terrestrial and aquatic settings and hence the application and interpretation of brGDGT-
509 derived proxies” This is a repetition from earlier in the introduction.

510 *This sentence has been removed from the revised manuscript.*

511
512 Lines 91-99: You should mention that brGMGTs have previously been called H-brGDGTs in the literature.

513 We have mentioned this as follows (lines 111-112):
514 *“Compared with brGDGTs, the branched glycerol monoalkyl glycerol tetraethers (brGMGTs, also*
515 *referred as H-brGDGTs) are a much less studied group of lipids.”*
516
517 Lines 91-111: This paragraph was very well written and is an example of the standard the entire
518 manuscript should meet.
519 As said above, we have carefully checked the English quality of our revised manuscript.
520
521 Lines 117-123: You go from talking about the hypothesis you aim to test in the paper to talking about the
522 aims of the paper. Surely your aim is to test the hypothesis you have just laid out - why do we need to talk
523 about more aims here?
524 Thank you very much for this comment. We consider as appropriate to transition from stating the
525 hypothesis to clearly discussing the aims of the paper in this paragraph.
526
527 Line 125-126: “by high population density”. High population density of what?
528 Population density refers to the number of human beings who live in a given region. We have removed
529 this as it is redundant with the next half sentence.
530
531 Line 127: “macrotidal”. Please define this term.
532 Macrotidal means large tidal range, as specified in the sentence where it is used.
533
534 Figure 1: I really like this figure - it nicely summarizes your water sampling campaign.
535 Thank you for this comment.
536
537 Line 167: “Both decarbonated and non-decarbonated samples (~6 mg for SPM and ~20 mg for soils) were
538 enclosed in a tin capsule” You should mention that you will split your samples and decarbonate one aliquot.
539 Otherwise, the reader is confused as to where your non-decarbonated samples are coming from.
540 Thank you for the suggestion. We have added the following sentence (lines 202-203):
541 *“The samples were split, and one aliquot was decarbonated.”*
542
543 Line 172-174: “The isotopic composition ($\delta^{13}\text{C}$ or $\delta^{15}\text{N}$) was expressed as the relative difference
544 between isotopic ratios in samples and in standards (Vienna Pee Dee Belemnite for carbon or atmospheric
545 N_2 for nitrogen)” Should be “and atmospheric N_2 ...”.
546 This has been corrected.
547
548 Line 176: What were these “additional...analyses”? Do you mean the same analyses aforementioned but
549 on different samples, or different analyses on different samples?
550 The elemental and isotopic data of the SPM and sediments collected in 2015 and 2016 ($n=84$) were
551 published by Thibault et al. (2019). To avoid confusion, we have rephrased this sentence as follows (lines
552 212-214):
553 *“Additional elemental and isotopic data based on SPM and sediments collected in 2015 and 2016 ($n=84$)*
554 *were obtained from Thibault et al. (2019).”*
555
556 Line 177: “(4-20g, $n=51$)” Looks to me like you’ve used the minus sign, not the en dash here. If so use

557 the en dash.

558 This has been corrected.

559

560 Line 180-183: “The total lipid extracts were then separated into fractions of increasing polarity on an
561 activated silica gel column, using (i) 30 mL of heptane, (ii) 30 mL of heptane:DCM (1/4, v/v), and (iii)
562 30 mL of DCM/MeOH (1/1, v/v) as eluents.” That seems like a nonstandard amount of solvent. Are you
563 using very large columns here? If so state how many g of silica gel were used.

564 We use glass columns with a total volume of ca. 10 mL to separate the lipids. The amount of solvent is
565 three times the column volume. We have modified the text as follows (lines 219-221):

566 *“The total lipid extracts were then separated into fractions of increasing polarity on an activated silica
567 gel column (ca. 10 mg), using (i) 30 mL of heptane, (ii) 30 mL of heptane:DCM (1/4, v/v), and (iii) 30 mL
568 of DCM/MeOH (1/1, v/v) as eluents.”*

569

570 Line 233: Vegan should be vegan. No capital V.

571 This has been corrected.

572

573 Lines 240-243: I don’t think you effectively explain how your hierarchical partitioning method actually
574 works here. As some readers won’t be familiar with this method, more details are needed.

575 We agree with the reviewer and have added more information about the hierarchical partitioning method
576 as follows (lines 294-300):

577 *“Briefly, this approach suggests that shared variance can be decomposed into equal components based
578 on the number of involved predictors (environmental factors), allowing for the estimation of the relative
579 importance of each predictor by adding its partial R^2 to the sum of all allocated average shared R^2 . While
580 most selection procedures, such as forward selection, use predictor ordering to assess variable
581 importance, hierarchical partitioning calculates individual importance (the sum of unique and total
582 average shared effects) from all subset models, generating an unordered assessment of variable
583 importance (Lai et al., 2022).”*

584

585 Figure 2. I really don’t like how the axis of these plots has been extended to include chart labels. The top
586 left panel scale is completely distorted by the addition of these labels. You should also define the features
587 of your “boxes” in your box plot. These comments apply to all boxplots in the manuscript.

588 We agree with the reviewer and have modified the plots (notably by changing the scales) and captions of
589 the figures accordingly.

590

591 Line 268: “The different brGDGTs were detected in all studied samples” Which brGDGTs?

592 We have specified all the brGDGTs which were detected in the following sentence (lines 347-348):

593 *“The different brGDGTs (IIIa5, IIIb5, IIIc5, IIa5, IIb5, IIc5, IIIa6, IIIb6, IIIc6, IIa6, IIb6, IIc6, IIIa7, IIIb7,
594 IIa7, Ia, Ib, Ic, 1050d, and 1036d) were detected in all studied samples.”*

595

596 Line 275: “The relative abundances of the brGDGTs were determined all along the Seine River basin” I
597 feel like this sentence should be at the start of this section not in the second paragraph.

598 We prefer to keep the current arrangement. We still prefer placing the description of chromatogram at
599 the beginning of the section and then fractional abundances, which emphasizes the logical flow of
600 information.

601
602 Line 290: “which explained 40.9% of the variance in two dimensions” Which two dimensions are these?
603 Thank you for the comment. These are the first two dimensions. We have corrected a typo here and
604 modify this paragraph as follows (lines 368-371):

605 *“A Principal Component Analysis (PCA) was performed to statistically compare the fractional*
606 *abundances of brGDGTs from different location (river, upstream and downstream estuary, based on SPM*
607 *and sediments collected in the river channel), which explained 54.1% of the variance in the first two*
608 *dimensions (Fig. 4a). The first axis (PC1) explained 40.9% of the variance, with negative loadings for*
609 *most of the 6-methyl brGDGTs and positive loadings for the remaining brGDGTs (Fig. 4a).”*

610
611 Line 291: “Samples from the downstream estuary clustered well” Colloquial language, you should
612 describe the data using words that don’t convey a value judgment.
613 We have removed the word “well” from this sentence.

614
615 Line 314-315: “It allowed to explain 39.79% of the variability through two dimensions.” Doesn’t make
616 sense - please proofread your manuscript.
617 We have removed this sentence.

618
619 I feel like you have just randomly placed the figures in the text. You should line up the first in-text citation
620 of a figure with the location of the figure in the manuscript. Currently, the text and the figures are out of
621 sequence which makes reading this document a challenge.

622 We have carefully checked our figures and in-text citations. They were appropriately positioned. We
623 present the PCA (RDA) plots for brGDGTs and brGMGTs together to avoid adding too many figures to
624 the manuscript. In the text, we describe the results related to brGDGTs first and then those related to
625 brGMGTs. The in-text citations indeed correspond to the order of the figures.

626
627 Figure 5: Visually this figure is quite busy. I don’t think having the brGDGT names in blue (the same
628 colour used for the downstream bubble) helps. I would use black for these names and also the arrows.
629 We agree with the reviewer and have changed the color of these names and arrows into black.

630
631 Line 336: “The brGMGTs identified in previous studies” Which brGMGTs and which studies? This lack
632 of precise usage of language is present throughout the text.
633 This has been corrected in the following sentence (lines 424-425):

634 *“The brGMGTs (H1020a, H1020b, H1020c, H1034a, H1034b, H1034c, and H1048) identified by Baxter*
635 *et al. (2019) were detected in the samples collected across the Seine River basin.”*

636
637 Line 343-345: “In SPM and river channel sediments, the total brGMGT concentration was observed to
638 be slightly higher in the riverine part ($0.26 \pm 0.24 \mu\text{g/g C}_{\text{org}}$) than in downstream ($0.20 \pm 0.13 \mu\text{g/g C}_{\text{org}}$)
639 and upstream estuary samples ($0.17 \pm 0.18 \mu\text{g/g C}_{\text{org}}$; Fig. S4b).” Slightly higher but not significantly
640 higher. If it’s not significant you should say so.

641 The difference in brGMGT concentration along the estuary is not significant. This has been acknowledged
642 as follows (lines 431-433):

643 *“In SPM and river channel sediments, the total brGMGT concentration was observed to be slightly (but*
644 *not significantly) higher in the riverine part ($0.26 \pm 0.24 \mu\text{g/g C}_{\text{org}}$) than in downstream estuary ($0.20 \pm$*

645 *0.13 µg/g C_{org}) and upstream estuary samples (0.17 ± 0.18 µg/g C_{org}; Fig. S5b). The total brGMGT*
646 *concentrations were the lowest in soils (surficial soils and mudflat sediments) all over the basin (0.07 ±*
647 *0.09 µg/g C_{org}; Fig. S5b)."*

648

649 Line 346: "The PCA analysis based on the brGMGT relative abundances (Fig. 5b) explained 70 % of the
650 variance". I'm unsure what the authors are trying to say here but I think they mean that the first two PCs
651 sum to 70%. The second half of the sentence "which allows to observe that samples from the different
652 parts of the basin clustered well apart from each other." doesn't make sense and I'm unsure what the
653 authors are trying to say.

654 Yes, the first two PCs sum to 70%. To clarify this point, the sentence has been rephrased as follows (lines
655 436-437):

656 *"The PCA analysis based on the brGMGT relative abundances (Fig. 4b) explained 70 % of the variance*
657 *in the first two dimensions, which separate samples from different parts of the basin."*

658

659 Line 357: "allows to explain" This phrase doesn't make sense in this context - please remove all uses of
660 it from the manuscript.

661 This has been corrected.

662

663 Lines 406-408: "The similarity in distributions between soils and downstream samples may be due to the
664 overrepresentation of downstream soil samples, as 82% of the soils were collected downstream (Fig. 1a
665 and Table 1)." I don't understand your point here. Are you saying that the similarity between downstream
666 estuary brGDGT distributions and soil brGDGTs is because the downstream estuary predominantly
667 receives brGDGTs from downstream soils?

668 We thank the reviewer for this comment. We have rephrased the sentence as follows to clarify this point
669 (lines 521-523):

670 *"This similarity in brGDGT distributions may be due to the influx of brGDGTs from the downstream soils*
671 *into the downstream estuary, as 82% of the soils were collected downstream (Fig. 1a and Table 1)"*

672

673 Lines 409-412: "Nevertheless, the soil-derived brGDGT contribution to the downstream samples is
674 expected to be much lower than the autochthonous one, as the average brGDGT concentration in soils
675 was ca. 3 times lower than the one in downstream (i.e. SPM and river channel sediment) samples (Fig.
676 S4a)." Right, but it's curious that the distributions are so similar between brGDGTs in soils and
677 downstream estuaries. To bring more clarity to this point it would be interesting to see you attempt a
678 machine learning approach (see Martinez-Sosa et al 2023, PP) to investigate whether (or not) a random
679 forest model can distinguish soil samples from downstream estuary samples.

680 As previously said, we applied a machine learning approach, similar to that of Martinez-Sosa et al. (2023),
681 to our dataset. Additional figures have been added to the supplementary material, as well as text to the
682 discussion (cf. reply to main comments above).

683

684 Lines 426-429: It would be great to see you calculate and report IR_{6+7me} following Wang et al (2021)
685 to determine if these indices correlate to salinity in an estuarine location.

686 We have calculated IR_{6+7me} as suggested by the reviewer. We have modified the figures and main text
687 accordingly (cf. reply to main comments above) and notably added the following sentence (lines 583-
688 586):

689 *“The salinity proxy (IR_{6+7me}) proposed by Wang et al. (2021) does not show significant correlation with*
690 *salinity in this study ($p > 0.05$, Wilcoxon test; Fig. S10). This suggests that IR_{6+7me} is relatively insensitive*
691 *in the Seine Estuary, potentially due to the preferential production of 6-methyl brGDGTs in specific*
692 *estuarine regions (i.e. KP 255.6-337)”*

693
694 433-436: “The distinct behavior of 6-methyl brGDGTs between lakes and the Seine river-sea continuum
695 might be due to the lower salinity range in the Seine River basin (0-32 psu) vs. the lakes (0-376 psu) 435
696 investigated by Wang et al. (2021). This suggests that the limited range of salinity variation in the Seine
697 River basin might be insufficient to trigger significant 6-methyl brGDGT production, as observed in
698 hypersaline lakes.” This is actually incorrect. Wang et al 2021 report that IR_{6me} is sensitive to salinity in
699 the range of 5-1000 (mg/L) but relatively insensitive beyond this range.

700 We agree with the reviewer and have modified the text accordingly by removing the reference to the
701 publication by Wang et al. (2021) here (lines 596-598):

702 *“Indeed, the significant negative correlations between the salinity and the relative abundance of 6-methyl*
703 *brGDGTs is observed in the Seine basin (Fig. S10), which suggests that the bacteria producing 6-methyl*
704 *brGDGTs are preferentially present in the low salinity area of the estuary.”*

705
706 458-460: “As the nutrient concentration is higher in the upstream part of the Seine estuary (Wei et al.,
707 2022), the substantial 6-methyl brGDGT production observed in the aforementioned zone ($260 < KP$
708 < 340 , Fig. 8)” Right but why would the nutrient runoff be higher for this specific section of the basin?
709 Do we see more agricultural activity here or something? It would be good to try and flesh out this point.
710 This specific region of the estuary is indeed characterized by intense agricultural activity, which could at
711 least partly explain the high nutrient concentration in this zone, especially during the low-flow season.
712 The text of the manuscript has been revised as follows (lines 616-619):

713 *“As the nutrient concentration is higher in the upstream part of the Seine estuary (Wei et al., 2022), and*
714 *this zone is characterized by high proportions of agricultural land use (Flipo et al., 2021), the substantial*
715 *production of 6-methyl brGDGT observed in the aforementioned zone ($260 < KP < 340$, Fig. 8) during*
716 *low flows could be attributed to elevated nutrient levels, particularly nitrogen, resulting from intense*
717 *agricultural activities.”*

718
719 Figure 8 and throughout: Make sure to superscript 15 in $d_{15}N$ and subscript 6 in IR_{6me} .

720 This has been corrected.

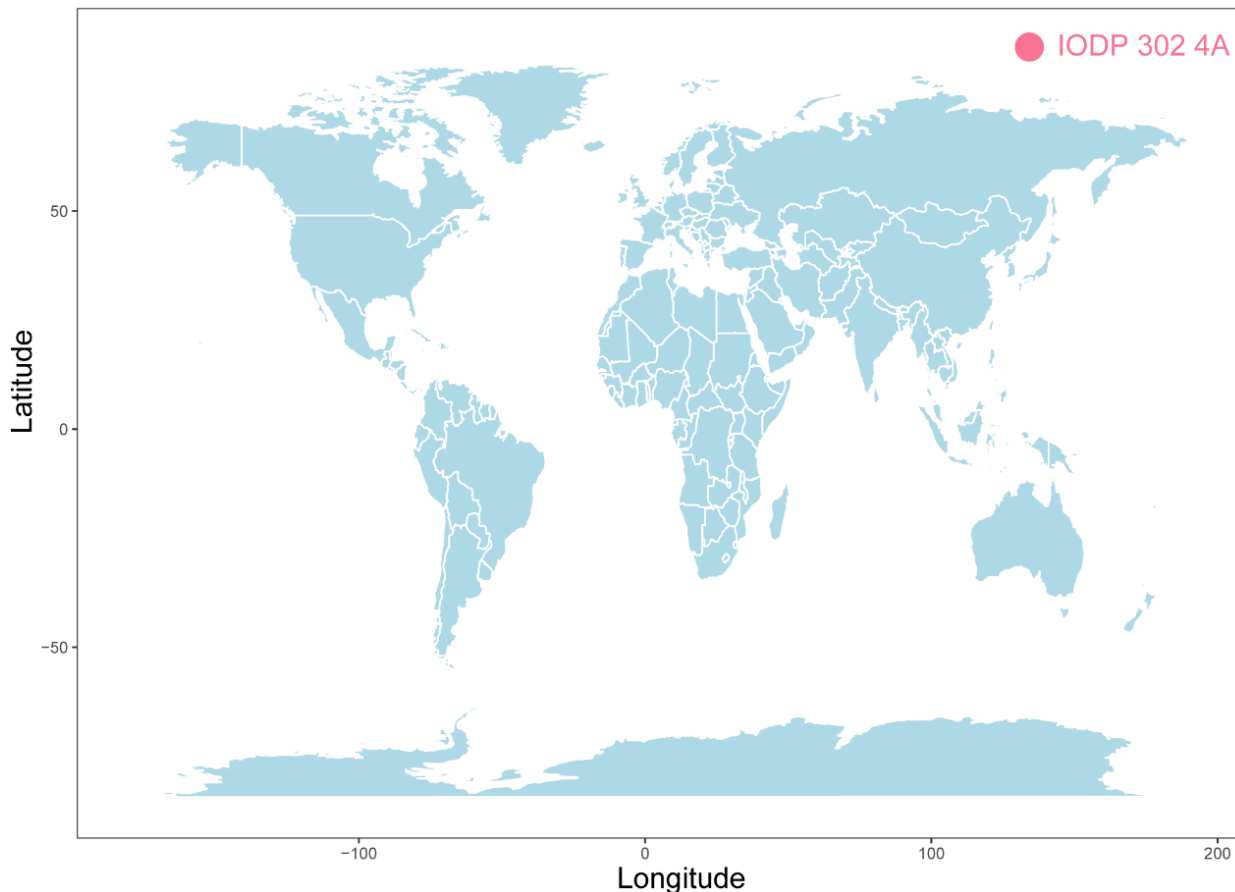
721
722 509-510: “The current knowledge on the parameters controlling the brGMGT distributions in the
723 terrestrial and marine realm is still limited.” Why is it limited? Be specific.

724 Thank you for the comment. This group of lipids (brGMGTs) has only recently gained attention.
725 Consequently, there are many aspects (e.g. controlling factors) still unknown for brGMGTs compared to
726 brGDGTs. To be more specific, we have rephrased our sentence as follows (lines 669-670):

727 *“The current knowledge on the parameters controlling the brGMGT distributions in the terrestrial and*
728 *marine realm is still limited, as there is little literature available (Kirkels et al., 2022a).”*

729
730 Fig 9: Almost impossible to see the data points on some of the figure panels (e.g. panel e). Make the
731 points bigger. Also, keep a consistent label text size to make the figure look neater. Also, you should say
732 in the caption how you constructed the straight lines drawn through the data in some panels (e for instance).

733 I'm assuming this is a linear regression but you have to inform the reader of your methods.
734 Thank you for this suggestion. To enhance visibility, we have increased the size of the data points in
735 figure panels, especially in panel e. Additionally, we have standardized the label text size across all figure
736 panels. Furthermore, we have provided more information in the figure captions.
737
738 557-558: “ However, the average concentrations of brGMGTs are an order of magnitude lower in the soils
739 than in the river channel sediments and SPM samples of the Seine basin (Fig. S4b).” Maybe it is, but
740 visually it doesn't look like that, so include the numbers in this sentence. You can also argue that the
741 brGMGT abundance within soils varies by an order of magnitude. Do you know what is driving such a
742 large variance in the soil brGMGT abundance?
743 We agree with the reviewer that the brGMGT concentration in soil samples shows large variance. This
744 highlights the need for further investigation into the environmental controls on brGMGT concentration
745 and distribution in soils. However, as shown by the boxplot, the upper and lower quartiles as well as the
746 median value of the soil brGMGT data are low compared to the river, upstream, and downstream samples.
747 In any case, downstream (SPM and sediment) samples and soils display distinct distribution and
748 concentrations, also captured by the application of the machine learning model to the brGMGT dataset
749 (cf. reply to the main comments above).
750 We have considered the comment of the reviewer in a revised manuscript through the following sentence
751 (lines 727-728):
752 *“A large variance in the soil brGMGT concentration was observed (Fig. S5b), suggesting that further*
753 *investigation is needed to better understand the environmental controls on the brGMGT production in*
754 *soils.”*
755
756 589: Missing the word “index” after BIT
757 This has been corrected.
758
759 You need a map showing the location of IODP 302 Hole 4A
760 We have added the following map showing the location of the core in the supplement:



761

762 Lines 605-607: “This core is considered proximal to the coast and has considerable changes in terrestrial
 763 inputs (i.e. continental spores and pollen) over time (Sluijs et al., 2009, 2006), making it a suitable
 764 paleorecord for testing runoff proxies.” Again would be great to have some specifics. The readers will be
 765 interested in how close this core site was to the coast around the PETM. You should also say why there
 766 was a considerable change in terrestrial inputs (I’m assuming large changes in sea level are responsible).
 767 We thank the reviewer for this comment. The changes of sea level are indeed responsible for the changing
 768 terrestrial inputs. We have rephrased this sentence as follows (lines 851-854):

769 *“This core is considered to record significant changes in terrestrial inputs (i.e. continental spores and
 770 pollen) due to sea level changes over time (Sluijs et al., 2009, 2006), making it a suitable paleorecord for
 771 testing runoff proxies.”*

772

773 Lines 616-617: “Such decreased runoff during the PETM body was previously attributed to a local sea
 774 level rise” Ah here is the explanation - this should have been in the previous paragraph. Also, be specific,
 775 are you saying there was decreased runoff during the PETM, OR did your sediment core record decreased
 776 runoff due to a change in sea level? These are two different things.

777 In addition to this core (Sluijs et al., 2008), a rise in sea level during the PETM has been recorded in many
 778 other sites worldwide (Speijer and Morsi, 2002; Harding et al., 2011; Sluijs et al., 2014). We have
 779 rephrased this sentence to clarify this point in a revised manuscript (lines 863-865):

780 *“Such decreased runoff during the PETM body was previously attributed to a rise in sea level (Sluijs et
 781 al., 2006), which has been recorded in many other sites worldwide (Speijer and Morsi, 2002; Harding et
 782 al., 2011; Sluijs et al., 2014).”*

783

784 References:

785

786 Martínez-Sosa, P., Tierney, J. E., Pérez-Angel, L. C., Stefanescu, I. C., Guo, J., Kirkels, F., ... & Reyes,
787 A. V. (2023). Development and application of the Branched and Isoprenoid GDGT Machine learning
788 Classification algorithm (BIGMaC) for paleoenvironmental reconstruction. *Paleoceanography and*
789 *Paleoclimatology*, 38(7), e2023PA004611.

790

791 Wang, H., Liu, W., He, Y., Zhou, A., Zhao, H., Liu, H., Cao, Y., Hu, J., Meng, B., Jiang, J., Kolpakova,
792 M., Krivonogov, S., and Liu, Z.: Salinity-controlled isomerization of lacustrine brGDGTs impacts the
793 associated MBT5ME' terrestrial temperature index, *Geochimica et Cosmochimica Acta*, 305, 33–48,
794 <https://doi.org/10.1016/j.gca.2021.05.004>, 2021.

795

796

797
798 Response to comments by reviewer #2
799

800 The authors analyzed brGDGTs and brGMGTs in soils, suspended particulate matter, and river sediments
801 in the Seine River basin to evaluate the environmental controls on and sources of these lipids. The basin
802 ranges from freshwater to estuarine, allowing the authors to evaluate the effects of salinity on the GDGT
803 compositions. The major motivation seems to be development of a new GMGT index, called “RIX”, to
804 detect terrestrial inputs of GMGTs to marine environments. The authors test this index through application
805 of Cenozoic sections of an IODP site.
806

807 There is now a relatively large literature on the environmental controls on GDGTs, though there is less
808 on GMGTs, and combining these across a riverine salinity gradient is a strength of the paper. Overall I
809 think the paper does provide some novel contributions and findings that merit publication. That said, there
810 are a number of technical problems that will require major revision before the paper can be published.
811 We thank the reviewer for his/her detailed comments and for recognizing the novelty and strength of our
812 work. A point-by-point reply to all the reviewer’s comment is provided below and is colored blue. The
813 text which has been added into the revised manuscript is shown in orange italics. The line numbers
814 correspond to those of the manuscript with tracked changes.
815

816 First, the Seine basin is complicated by the presence of a dam that separates sections of the river
817 influenced by tides (salinity) from sections upstream. The dam also presumably traps upstream sediment
818 and likely presents a barrier for transport of GDGTs (other than SPM). The authors also have relatively
819 few soil sampling sites – there are only 5 sites and the soil samples are dominated by downstream
820 estuarine soils. I don’t think these challenges are adequately discussed in the paper. The dam may be
821 a good thing for the study, since it establishes clear environmental boundaries, but it could be tricky to
822 apply a GDGT index from this environment to other sites/time periods.
823

824 The dam of Poses (cf. location on the revised map below) is the frontier between the Seine river and
825 estuary. It represents the upstream limit of the fluvial estuary and of the tidal propagation. It was built in
826 1887 to regulate the water level and to allow navigation of the ships up to Paris, whatever the season.
827 Indeed, the average water flow of the Seine River measured at Poses is $\sim 470 \text{ m}^3 \text{ s}^{-1}$, with marked intra-
828 annual differences between winter and summer flows ($\sim 250 \text{ m}^3 \text{ s}^{-1}$ in the summer and over $700 \text{ m}^3 \text{ s}^{-1}$ in
829 the winter). Whatever the period of the year, at least part of the water from the Seine river upstream Poses
830 flows to the estuary. Therefore, the dam should not prevent (part of) the riverine GDGTs associated to
831 SPM to arrive to the estuary. Nevertheless, it cannot be excluded that part of the riverine sediments are
832 trapped by the dam.
833

834 Regarding the estuary itself (downstream Poses), it comprises two major sections: the upstream,
835 freshwater section (from site 5 to 12) and the lower, downstream section influenced by salinity (from site
836 12 to the coastal zone). All our estuarine samples were (logically) collected downstream of the dam of
837 Poses. Therefore, the observed changes in brGDGT/brGMGT distribution and abundance all along the
838 estuary, with distinct signal in the upstream and downstream estuarine zones, are intrinsic to the
839 biogeochemical functioning of the Seine estuary and cannot be attributed specifically to this dam.
840 Corresponding details were added to the revised manuscript as follows (lines 483-490):

841
842 *“The decrease in the fractional abundance of 6-methyl brGDGTs from the upstream estuary to the*
843 *downstream estuary cannot be explained by the dam located at Poses (Fig. 1a). This dam separates the*
844 *riverine part of the Seine from the upstream estuarine section. Even during the low-flow season (Fig. 1b),*
845 *at least part of the water from the Seine River upstream of Poses flows into the estuary (Romero et al.,*
846 *2019). Thus, the dam should not prevent (part of) the riverine brGDGTs associated with SPM from*
847 *reaching the estuary. It cannot be excluded that part of the riverine sediments is trapped by this dam.*
848 *Nevertheless, all our estuarine samples were collected downstream of the dam, implying that the observed*
849 *changes in brGDGT abundance and distribution within the estuary are intrinsic to the biogeochemical*
850 *functioning of the Seine estuary and cannot be attributed to the dam.”*

851
852 Regarding the soils, we agree with the reviewer and acknowledge the limitations of our sampling strategy,
853 with a low number of sampling sites, mainly located downstream. We cannot exclude that the overlay in
854 brGDGT/brGMGT distribution between the soils and the downstream estuary SPM and sediment samples
855 is partly due to the sampling approach. This has been specified in a revised manuscript with the following
856 sentence (lines 521-523):

857 *“This similarity in brGDGT distributions may be due to the influx of brGDGTs from the downstream soils*
858 *into the downstream estuary, as 82% of the soils were collected downstream (Fig. 1a and Table 1).”*

859
860 Nevertheless, the comparison of the brGDGT/brGMGT concentrations and distributions between soils
861 and downstream estuary samples allows distinguishing the two types of samples, as captured by the
862 application of an independent machine learning approach to our brGMGT/brGDGT datasets.

863
864 Last, we kindly disagree with reviewer 2 when saying that “it could be tricky to apply a GDGT index
865 from this environment to other sites/time periods.” The RIX index was developed based on samples from
866 the Seine estuary. Nevertheless, it was successfully tested in both modern (Godavari River basin) and
867 past settings (marine sedimentary core IODP 302-4A), showing its potential general applicability.

868
869 Second, there are a lot of data / statistical difficulties with this paper, the details of which are discussed
870 below. At times the authors compare concentrations of GDGTs to evaluate in situ production, which is
871 generally not a good way to do this due to the effects of sediment transport from soils to river to estuaries
872 – concentrations may be higher in SPM than soils, for instance, as SPM contains less coarse-grained
873 particles. Although the writing is a bit unclear, the authors appear to compare results of two PCAs, one
874 on soils and one on aquatic samples, to differentiate these two sample types, which is not possible given
875 how PCA works.

876
877 In order to better evaluate the *in situ* production of brGDGTs/brGMGTs in the estuary, a machine learning
878 approach GDGT/GMGT datasets, as suggested by Reviewer #1. We have now several lines of evidence
879 supporting the *in situ* production of brGDGT/brGMGTs in the Seine estuary:

- 880 1) higher brGDGT/brGMGT concentrations in aquatic environments compared with soils.
- 881 2) distinct distributions between soils and aquatic settings (riverine and upstream estuarine samples)
882 identified by PCA.
- 883 3) the application of the machine learning approach, which allows distinguishing downstream estuary and
884 soil samples based on brGDGT/brGMGT distributions. This is addressing the overlap observed between

885 downstream estuary and soil samples in the PCA biplot based on brGDGT/brGMGT distributions.
886 As detailed in the reply to reviewer 1, additional discussion on the *in situ* production of
887 brGDGTs/brGMGTs (based on the above mentioned points) has been added to our revised manuscript.
888

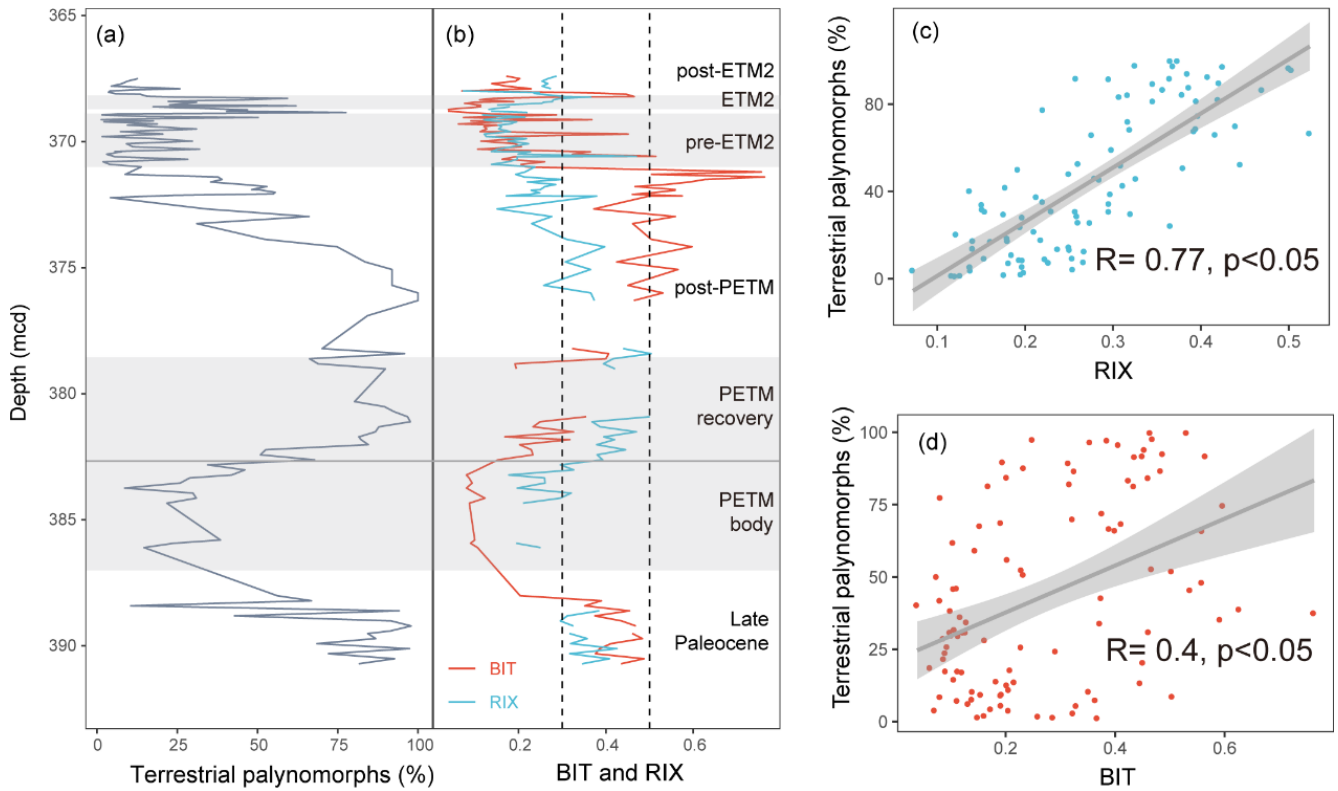
889 In addition, we would like to clarify that the PCAs of soils and aquatic samples were not done separately.
890 The biplots do not correspond to a simple overlay. Only active individuals (river, upstream, and
891 downstream estuarine samples) were used for principal component analysis. The coordinates of passive
892 individuals (also known as supplementary individuals) (i.e. soil samples) were just **predicted**/projected
893 using the **existing** PCA information obtained with active ones. This is actually a widely used approach
894 which can be implemented by the R package FactoMineR ([https://cran.r-](https://cran.r-project.org/web/packages/FactoMineR)
895 [project.org/web/packages/FactoMineR](https://cran.r-project.org/web/packages/FactoMineR)). It has also been used in a recent GDGT paper (Kirkels et al.,
896 2022 *Biogeosciences*), which aims to compare GDGT distributions in soils and aquatic settings. We prefer
897 this approach as it effectively delivers the key information: brGDGT/brGMGT distributions in riverine
898 and upstream estuarine samples are distinguishable from those in soils. However, in the PCAs based on
899 brGDGT/brGMGT distribution, soils partly overlay with downstream estuary samples. This similarity
900 may be at least partly attributed to our sampling strategy, given that most of the soils were collected
901 around the downstream estuary, as mentioned in the manuscript. Nevertheless, we can efficiently
902 distinguish brGDGT/brGMGT distributions in downstream estuarine samples from those in soils by using
903 an independent machine learning approach as said above.
904

905 In the revised manuscript, we have modified the figure caption of the PCAs to better illustrate our
906 methodology (lines 408-410):

907 *“The coordinates of soils (passive individuals) are added as an overlay and are predicted based on the*
908 *information provided by the existing PCA performed on SPM and sediments (active individuals).”*
909

910 Third, Section 4.4 compares the application of the RIX to IODP site 302 to results from other
911 measurements, such as the BIT and % terrestrial palynomorphs. The comparison is largely qualitative,
912 and it’s hard to tell from Figure 11 how well these compare in a statistical sense. Could the authors
913 provide correlation coefficients to show that the RIX captures terrestrial inputs?

914 We thank the reviewer for this comment. We have provided the correlation coefficients between RIX
915 and % terrestrial palynomorphs as well as between BIT and % terrestrial palynomorphs in our revised
916 manuscript. The corresponding figure is provided below:
917



918
 919 *Figure 10: Comparison between (a) terrestrial palynomorphs (%) and (b) BIT and RIX across the upper*
 920 *Paleocene and lower Eocene between 391 and 367 meters composite depth below sea floor (mcd) of*
 921 *IODP Expedition 302 Hole 4A. Terrestrial palynomorphs data are from Sluijs et al. (2006) and Sluijs et*
 922 *al. (2009). RIX and BIT were calculated using data from Sluijs et al. (2020). Grey shading represents*
 923 *Eocene Thermal Maximum 2 (ETM2), pre-ETM2 interval, and Paleocene-Eocene Thermal Maximum*
 924 *(PETM). Dotted line represents cutoff values of RIX (below 0.3 for marine contribution and above 0.5 for*
 925 *riverine contribution). Linear regression of the RIX (c) and BIT (d) against the terrestrial palynomorphs.*
 926 *Shaded area represents 95% confidence intervals.*

927
 928 We have also added the following sentence to our revised manuscript (lines 912-914):

929 *“This indicates that RIX performs better in this core compared with BIT, which is further supported by a*
 930 *higher correlation coefficient observed between RIX and terrestrial palynomorphs (0.77; Fig. 10c)*
 931 *compared with BIT and terrestrial palynomorphs (0.4; Fig. 10d)”*

932
 933 Detailed comments:

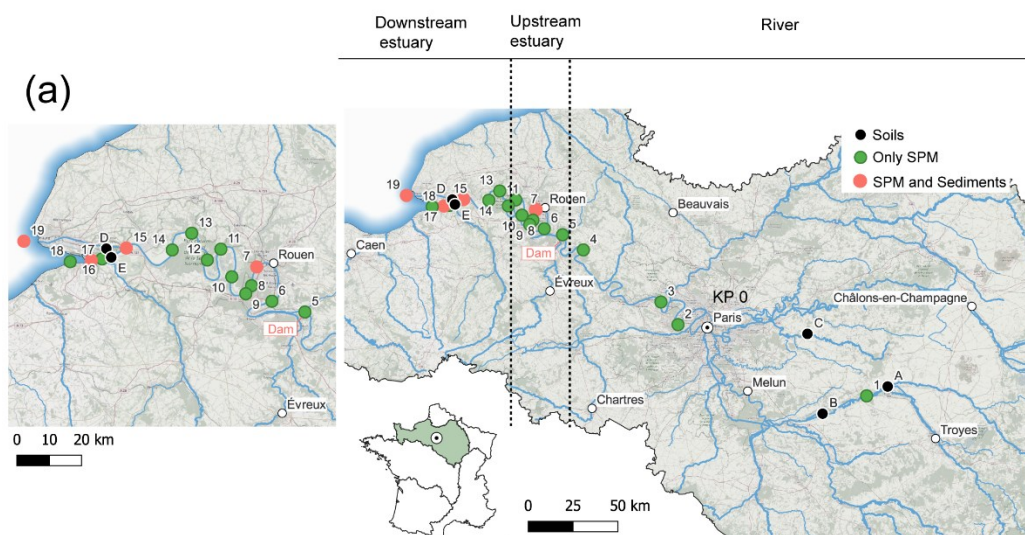
934
 935 Section 2.2. It is a bit hard to tell from this description and the table exactly what samples were collected
 936 and analyzed. I take it from the description that 1) subsurface SPM was collected from every green dot
 937 (correct?). 2) deeper water SPM was filtered from 5 sites (perhaps these could be indicated in the table),
 938 3) Sediment samples from 8 cores were collected. I cannot tell from the description what depth in the core
 939 these samples were taken from (10 cm?), nor how 8 cores yielded n = 68.

940
 941 Perhaps the dots could be color coded to indicate what types of samples exist (surface SPM, subsurface
 942 SPM, these + sediment). It might also be helpful to designate the environment type (river, upstream
 943 estuary, downstream estuary) on the map. It would be particularly helpful to indicate the city of

944 Poses/location of the dam on this map.

945

946 We agree with the reviewer here. We have changed the color of the dots in the map to indicate the different
947 sample types. Locations where only soils were collected were indicated in black; those where only SPM
948 were collected were indicated in green; those where both SPM and sediments were collected were
949 indicated in red. In addition, the location of the dam, as well as information about the environmental type
950 (river, upstream estuary, downstream estuary), have been added to the map. The revised map and caption
951 are shown below:



952

953 *Figure 1: Geographical locations of sampling sites in the Seine River Basin (KP: kilometric point, the*
954 *distance in kilometers from the city of Paris (KP 0)). The sampling sites from upstream estuary and*
955 *downstream estuary are shown in the zoom-in figure. Sub-surface SPM was collected for all sites from*
956 *site 1 to site 18, while both sub-surface and bottom SPM were collected at sites 4, 6, 10, 13, and 15.*

957

958 To maintain the readability of the map and avoid too many colors, additional details have been provided
959 in the caption as well as in Table 1. Table 1 allows distinguishing 5 categories of sites depending on the
960 type of samples collected: 1) only soils; 2) only subsurface SPM; 3) subsurface SPM and sediments; 4)
961 subsurface and bottom SPM as well as sediments; 5) subsurface and bottom SPM. We have differentiated
962 subsurface and bottom SPM samples in this table.

963

964 Regarding the sediment samples, they were collected from 7 cores (and not 8). This typo has been
965 corrected in the revised manuscript. We have added further details on the sampling strategy as follows
966 (lines 180-184):

967 *“Sediments (n=68) from 7 cores (10-cm depth) were collected in the river channel at the same sites as*
968 *these SPM samples in 2015 and 2016 using a UWITEC corer as described by Thibault et al. (2019) (Table*
969 *1). These sediments were further sliced (1-cm thickness) and freeze-dried. For each core, ten samples*
970 *were analyzed for brGDGTs and brGMGTs, except for the one collected at site 17 in April 2016, where*
971 *no lipids were detected between 4-5 and 5-6 m depth.”*

972

973 What differentiates “upstream estuary” and “downstream estuary”? Is this salinity? Or judgement?
974 The river and upstream estuary are differentiated by the dam located at Poses. The tide influences the

975 estuary up to Poses, where the dam prevents further tidal propagation. The upstream and downstream
976 estuary are differentiated based on spatiotemporal variations of salinity. The upstream estuary
977 corresponds to the freshwater tidal sector, whereas the downstream estuary is affected by a salinity
978 gradient (e.g. Romero et al., 2016, Environmental Science and Policy; Druine et al., 2018, Marine
979 Geology). This has been clearly specified in the revised manuscript as follows (lines 150-155):

980 *“The tide influences the estuary up to the city of Poses (site 5, KP 202 in Fig. 1a; KP represents kilometric*
981 *point and is defined as the distance in kilometers from the city of Paris), where a dam constitutes the*
982 *boundary between the river and the estuary. Based on spatiotemporal variations of salinity, the estuary*
983 *can be divided into two major parts. The upstream estuary corresponds to the freshwater tidal sector (KP*
984 *202 to KP 298, from site 5 to site 12; Fig. 1a and Table 1) and the downstream estuary is influenced by*
985 *a salinity gradient (starting at KP 298, from site 12 to the coastal area; Fig. 1a and Table 1) (Romero et*
986 *al., 2016; Druine et al., 2018).”*

987
988 Line 237: “correlations” here should be “relationships”. These are not correlations in the statistical sense.
989 This has been corrected.

990
991 Line 271? Should this be “decreased in the downstream estuary” samples (not just “downstream”)?
992 Having defined upstream estuary and downstream estuary it is good to stay with these terms.
993 The term “estuarine” has been added in this sentence as well as in other sentences throughout our
994 manuscript.

995
996 Line 290. “Negative loadings” is confusing. On which axis? Both? I suggest describing the results by
997 axis – first axis 1, then 2.

998 To clarify this point, this sentence has been revised as follows (lines 368-371):

999 *“A Principal Component Analysis (PCA) was performed to statistically compare the fractional*
1000 *abundances of brGDGTs from different location (river, upstream and downstream estuary, based on SPM*
1001 *and sediments collected in the river channel), which explained 54.1% of the variance in the first two*
1002 *dimensions (Fig. 4a). The first axis (PC1) explained 40.9% of the variance, with negative loadings for*
1003 *most of the 6-methyl brGDGTs and positive loadings for the remaining brGDGTs (Fig. 4a).”*

1004
1005 Figure 3 is not particularly helpful to the reader. If the authors wish to retain it, I suggest moving it to
1006 supplemental text.

1007 As suggested by the reviewer, we have moved this figure to the supplement (Fig. S2).

1008
1009 Results:

1010
1011 The results of the “bulk parameters” describes the elemental and bulk stable isotopic composition of the
1012 solid samples. Nowhere does the paper describe results of other environmental parameters –
1013 temperature, etc. It would be helpful to have at least a table indicating the mean and range of these. I
1014 expect, for instance, that there is a large range of salinities associated with these samples and a very
1015 narrow range of temperatures (they are all close to each other).

1016 We thank the reviewer for this comment. Our revised manuscript now includes a new supplementary table
1017 (shown below) to describe the available environmental parameters (temperature, salinity, water discharge,
1018 TOC and TN):

1019
1020

Table S1. Description of available environmental parameters

	River	Upstream estuary	Downstream estuary	Soil
Min temperature (°C)	20	8.49	6.4	n.a.
Max temperature (°C)	23.41	24.4	23.38	n.a.
Mean temperature (°C)	21.51	20.09	18.27	n.a.
Number of samples	6	44	62	n.a.
Min salinity	0	0	0.1	n.a.
Max salinity	0.3	0.32	32.3	n.a.
Mean salinity	0.2	0.22	3.77	n.a.
Number of samples	6	43	60	n.a.
Min discharge (m ³ /s)	99	99	99	n.a.
Max discharge (m ³ /s)	156	978	978	n.a.
Mean discharge (m ³ /s)	129.78	183.62	218.85	n.a.
Number of samples	9	48	62	n.a.
Min TOC (%)	0.82	0.75	0.11	0.22
Max TOC (%)	4.22	7.71	7.35	22.28
Mean TOC (%)	2.88	4.64	3.3	3.03
Number of samples	9	57	120	51
Min TN (%)	0.12	0.12	0.01	0.01
Max TN (%)	0.58	0.84	0.619	1.07
Mean TN (%)	0.37	0.51	0.31	0.24
Number of samples	9	57	120	51

n.a.= not applicable

1021
1022

1023 The treatment of the soils samples in the analysis and results is difficult to understand. It appears that a
1024 large number of soils (up to 34) was taken from some sampling sites, whereas at others 1 sample was
1025 taken. These data were then analyzed via PCA separately from the aquatic samples, and the PCA was
1026 overlaid onto the PCA of the aquatic samples. The authors conclude that the PCAs show that the GDGT
1027 distribution of soils overlap with the SPM and channel sediments. If the PCAs were done separately, one
1028 cannot simply overlay the biplots and conclude that they overlap – the PCAs may capture different
1029 variance structures such that the PCA axes are not the same. If the authors wish to compare the soils
1030 and aquatic samples, do a PCA on all the data together. It’s always possible to do a second PCA excluding
1031 the soils to evaluate the variance structure of the aquatic samples alone.

1032 [This comment was addressed above.](#)

1033

1034 Line 290: “explained 40.9% of the variance in two dimensions”. What is meant by this? Based on the
1035 plot, axis 1 captures 40.9% of the variance and axis 2 13.2%.

1036

1037 [We thank the reviewer 2 for this comment, which was also made by reviewer 1. We have rephrased this](#)

1038 paragraph as follows (lines 368-371):

1039 *“A Principal Component Analysis (PCA) was performed to statistically compare the fractional*
1040 *abundances of brGDGTs from different location (river, upstream and downstream estuary, based on SPM*
1041 *and sediments collected in the river channel), which explained 54.1% of the variance in the first two*
1042 *dimensions (Fig. 4a). The first axis (PC1) explained 40.9% of the variance, with negative loadings for*
1043 *most of the 6-methyl brGDGTs and positive loadings for the remaining brGDGTs (Fig. 4a).”*

1045 Line 346: Similar problem. I think the authors mean that axes 1 and 2 capture 71%. The PCA will
1046 capture more than this on axes 3 - ???

1047 We agree with the reviewer. The first two dimensions explain 70% of the brGMGT variations. The
1048 corresponding sentence has been rephrased as follows (lines 436-437):

1049 *“The PCA analysis based on the brGMGT relative abundances (Fig. 4b) explained 70 % of the variance*
1050 *in the first two dimensions, which separates samples from different parts of the basin.”*

1052 Similar problems exist in the description of the RDA, Section 3.3

1053 We have specified that 30.2% of the variance was captured from the first two axes in the revised
1054 manuscript.

1056 4.1.1. Why do the authors focus on the 6-methyl brGDGTs here?

1057 We start this section by discussing 6-methyl brGDGTs, as this group of compounds is typically produced
1058 in rivers. Nevertheless, this section is also mentioning and discussing the variations of the relative
1059 abundances of other types of brGDGTs, especially 7-methyl brGDGTs, across the salinity gradient.

1061 Line 390: The authors suggest that the higher abundances of 6-methyl brGDGTs in upstream vs.
1062 downstream samples may reflect degradation:

1063 “It may reflect the fact that riverine 6-methyl brGDGTs are more easily degraded than soil-derived
1064 homologues and only partially transferred downstream.”

1065 Why would 6-methyl brGDGTs produced in a river degrade faster than those produced elsewhere? The
1066 authors argue that this could reflect attachment to particles – but how do these particles differ in upstream
1067 vs. downstream river environments.

1068 It seems likely that production of the 6-methyl compounds is suppressed in downstream environments
1069 and the dam traps the upstream sediments (and lipids). Can the authors show that this is not the case?

1071 The decrease in the abundance of 6-methyl brGDGTs from the upstream estuary to the downstream
1072 estuary cannot be explained by the dam located at Poses, as the latter is separating the riverine part of the
1073 Seine and the upstream part of the estuary. There is no dam between the upstream and downstream parts
1074 of the estuary (cf. revised version of the map above). Therefore, we favor other hypotheses discussed in
1075 the manuscript to explain the changes in 6-methyl brGDGT abundances along the estuary, including 1)
1076 preferential degradation of labile (riverine) 6-methyl brGDGTs, as notably proposed by De Jonge et al.
1077 (2015) and 2) dilution by brGDGTs from other sources during downstream transport.

1079 Regarding the first hypothesis, the higher degradation of 6-methyl brGDGTs upstream could indeed be
1080 due to the different attachment to particles upstream vs. downstream. The median diameter of the SPM
1081 was monitored between February 2015 and June 2016 in the upstream (sites 7 and 10) and downstream

1082 (sites 15 and 17) parts of the Seine Estuary (Druine, 2018: <https://theses.hal.science/tel-01896520>).
1083 Upstream, the size of the particles showed only a slight dispersion (80-110 μm) whatever the hydrological
1084 conditions. The homogeneity of the size of the particles in the upstream estuary likely reflects their
1085 predominant continental origin (i.e. Seine river before the dam of Poses). In contrast, a large variability
1086 in the size of the SPM particles was observed in the downstream estuary (15-20 μm to 80-90 μm), related
1087 to the complex flocculation and defragmentation processes of the particles occurring in this part of the
1088 estuary (Druine, 2018). Therefore, the variability in the size of the SPM particles from upstream to
1089 downstream could have an influence on the brGDGT distribution in the Seine estuary. This point is now
1090 discussed in the revised manuscript using the aforementioned data (lines 500-509):

1091 *“Indeed, the higher degradation of 6-methyl brGDGTs upstream could be attributed to their different*
1092 *attachment to particles compared to downstream. The median diameter of the SPM was monitored*
1093 *between February 2015 and June 2016 in both the upstream (sites 7 and 10) and downstream (sites 15*
1094 *and 17) parts of the Seine Estuary (Druine, 2018). The particle size showed only slight dispersion (80-*
1095 *110 μm) under various hydrological conditions in the upstream estuarine section. The homogeneity in*
1096 *particle size in the upstream estuary likely reflects its predominantly continental origin (i.e. from the*
1097 *Seine River before the dam at Poses). In contrast, a large variability in the size of SPM particles was*
1098 *observed in the downstream estuary (15-20 μm to 80-90 μm), attributed to the complex flocculation and*
1099 *defragmentation processes of particles in this part of the estuary (Druine, 2018). Hence, the variability*
1100 *in the size of SPM particles from upstream to downstream could influence the distribution of brGDGTs*
1101 *in the Seine estuary.”*

1102
1103 Line 405. Here the authors suggest that the brGDGT distributions in estuarine soils is similar to that of
1104 the downstream samples, based on the PCA (see comment above about the PCA). In the next section
1105 (4.1.2), this is not discussed and instead production of the brGDGTs in saline environments is the primary
1106 factor accounting for compositional differences in upstream vs. downstream samples. Please coordinate
1107 these ideas.

1108 Since PCA alone does not allow distinguishing brGDGT distributions between soils and downstream
1109 estuary samples, we further applied a machine learning approach as suggested by Reviewer #1. This
1110 method supports the *in situ* production of brGDGTs by effectively distinguishing the brGDGT
1111 distributions between downstream estuary and soil samples. As brGDGTs are produced *in situ*, we can
1112 explore the compositional differences of these compounds from upstream to downstream and investigate
1113 the controlling factors. This has been discussed in a revised manuscript, as also detailed in the reply to
1114 reviewer 1.

1115
1116 Line 487, 559: One cannot conclude from concentrations alone that the GMGTs are produced in aquatic
1117 environments. Soils contain abundant coarse clastic material that may be lost in the fine SPM and river
1118 sediment. The distributions (relative abundances) of GMGTs are key to identifying *in situ* production.
1119 We fully agree with reviewer 2. The relative abundances of brGMGTs are essential to identify *in situ*
1120 production in the estuary, especially through machine learning approach. This comment was addressed
1121 above.