

1 **Anonymous Referee #1**

2 Received and published: 1 June 2018

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4 Note regarding document formatting: black text shows original referee comment, blue text shows  
5 author response, and red text shows quoted manuscript text. Changes to manuscript text are  
6 shown as *italicized and underlined*. Bracketed comment numbers (e.g. [R1.1]) were added for  
7 clarity. All line numbers refer to discussion/review manuscript.

8  
9 [R1.0] This paper builds on existing literature examining unsupervised learning techniques to improve the  
10 interpretation and classification of data obtained with WIBS UV-LIF spectrometers. As shown in  
11 previous publications, Hierarchical Agglomerative Clustering (HAC) can serve as a robust data analysis  
12 method for classification/interpretation of bioaerosol data but the accuracy of technique is highly sensitive  
13 to the choice of clustering linkage and data pre-treatment (e.g., Crawford et al., 2015); this is further  
14 explored in this paper which elucidates how data pre-treatment choices such as choice of fluorescent  
15 threshold and log normalising data may influence clustering accuracy using laboratory samples of known  
16 particle types (Savage et al., 2017) in various synthetic mixtures, and thus the authors present tentative  
17 recommendations of data pretreatment regimes depending on the analysis goals. Overall the paper is well  
18 written and the computational experiments well thought out. The findings here are useful and further  
19 validate the usefulness of Hierarchical Agglomerative Clustering for interpretation of WIBS data. The  
20 results also provide a useful framework for testing Hierarchical Agglomerative Clustering data pre-  
21 treatment regimes for other atmospheric science data problems and neatly demonstrate the potential  
22 pitfalls of not rigorously performing such tests. I recommend publication after the following comments  
23 have been addressed.

24  
25 [A1.0] Author response: We thank the referee for her/his positive summary of the manuscript and  
26 recommendation to publish after comments are addressed.

27  
28 Specific comments

29 [R1.1] L73-77: The authors have conflated some of the terminology relating to unsupervised and  
30 supervised leaning methods. I'm uncomfortable with the use of the term clustering when discussing  
31 supervised methods as clustering specifically relates to cluster analysis. I suggest replacing "clustering  
32 techniques" with "classification algorithms" and "(trains) the clustering algorithm" with "(trains) the  
33 classification algorithm".

34  
35 [A1.1] The referee raises a good point. We changed terminology on page 2 according the referee  
36 suggestions, as listed below:

- 37 - L68: "*Classification algorithms, including several clustering techniques in particular, have*  
38 *shown successful results ...*"  
39 - L73: "~~Clustering techniques~~ *Classification algorithms* can be divided ..."  
40 - L76: "This type of method enhances (trains) the ~~clustering-classification~~ *algorithm* in that the  
41 output ~~cluster classes~~ *groups* are predetermined ..."

42  
43 [R1.2] L120: Please state the bands and what they relate to.

44  
45 [A1.2] Additional text was added, as shown below:

46 "The WIBS collects 3 channels of fluorescence intensity information (FL1, FL2, and FL3),  
47 particle size, and particle asymmetry for each interrogated particle. *The bands of excitation and*  
48 *fluorescence emission are: FL1 ( $\lambda_{ex} = 280 \text{ nm}$ ,  $\lambda_{em} = 310 - 400 \text{ nm}$ ), FL2 ( $\lambda_{ex} = 280 \text{ nm}$ ,  $\lambda_{em} = 420$   
49  $- 650 \text{ nm}$ ), and FL3 ( $\lambda_{ex} = 370 \text{ nm}$ ,  $\lambda_{em} = 420 - 650 \text{ nm}$ ). The excitation and emission  
50 wavelengths chosen for each of the 3 fluorescence channels were designed to maximize the*

51 information gained about key biological fluorophores present in a broad range of bioparticles  
52 (Kaye et al., 2005; Pöhlker et al., 2012). *Early generations of UV-LIF bioaerosol spectrometers*  
53 *were often interpreted to be able to detect proteins via channels similar to FL1 and products of*  
54 *active cellular metabolism (i.e. riboflavin and NAD(P)H) via channels similar to FL3, but these*  
55 *approximations are gross simplifications that confound more detailed investigation of particle*  
56 *types.”*  
57

58 [R1.3] L198: Can the authors please clarify why they have used log spaced bins. Do you mean that you  
59 have taken a log of the data and it is binned naturally by the discrete nature of the detector resolution (i.e.,  
60 fine bins) or have you binned the data into specific (coarse) log bins? If it is the latter can you please state  
61 what the bins are and can you comment on how forcing the data to in bins may influence the clustering?  
62 My concern here is that too coarsely binning the data may create artificial hotspots due to reduced  
63 resolution and bias the clustering, reducing the capacity to differentiate between particles with similar  
64 properties. Can the authors comment on this and demonstrate the effect this may have by providing an  
65 example for comparison where the data is converted to log space and not binned. I also wonder if the bins  
66 should be normalised by the bin width to further complicate matters.  
67

68 [A1.3] Aspects of this discussion are presented in L209-212. To summarize in different words,  
69 the data values from a given channel were either used as recorded (i.e. “value”) or as  
70 logarithmically transformed (i.e. “log(value)”), depending on the Scenario. The values were not  
71 forced into specific bins, but rather input into the cluster algorithm using the exact value in either  
72 of these forms. The reason that logged values can provide different results by HAC is that the  
73 distance between points is different in linear space or log space, because the cluster process does  
74 not independently take into account whether a value is as recorded or as log(value). Because  
75 many real-world particle variables can present normal distributions only in log space (i.e.  
76 lognormal size distributions), we explored inputting values in both raw and log forms.  
77

78 The following sentence was added to the manuscript at L211 for clarity:

79 *“By this process, data values were input into the algorithm as log(value), but without additional*  
80 *binning.”*  
81

82 [R1.4] L254: Can the authors comment on the environmental applicability of the chosen ratios. I would  
83 suspect that in an urban environment you may expect something closer to a ratio of 1:99 fungal to diesel  
84 particles with the converse being true in a forest environment. How does the clustering perform under  
85 such extreme mismatches?  
86

87 [A1.4] We originally explored three different ratios of particle concentrations (80:20, 50:50, and  
88 20:80) for each of the three match-ups discussed in Figure 3 in order to show that input ratio can  
89 be important to how the algorithm responds. This was certainly not intended to be exhaustive, and  
90 one could additionally explore more extreme ratios. So to limit the scope of the analysis here, we  
91 chose to present evidence only that the ratio matters, without trying in all cases to predict ratios  
92 that could be relevant to a wider range of ambient environments.  
93

94 The question the referee brings up is interesting, however, and so we explored 1:99 ratios of each  
95 of the three particle type combinations presented in Figure 3, where the bioparticle is the minority  
96 concentration in each experiment. The results are shown below in a plot/table form identical to  
97 how they are presented in Figure 5. The Bacteria:Diesel and Fungi:Dust separations still  
98 performed quite well (6.6% and 13.5% misclassification, respectively), even with the extreme  
99 mismatch of input concentrations. The Fungi:Diesel separation was poor, however, in a 2-factor  
100 solution, because the Diesel particles split into both clusters, and the Fungi particles were likely

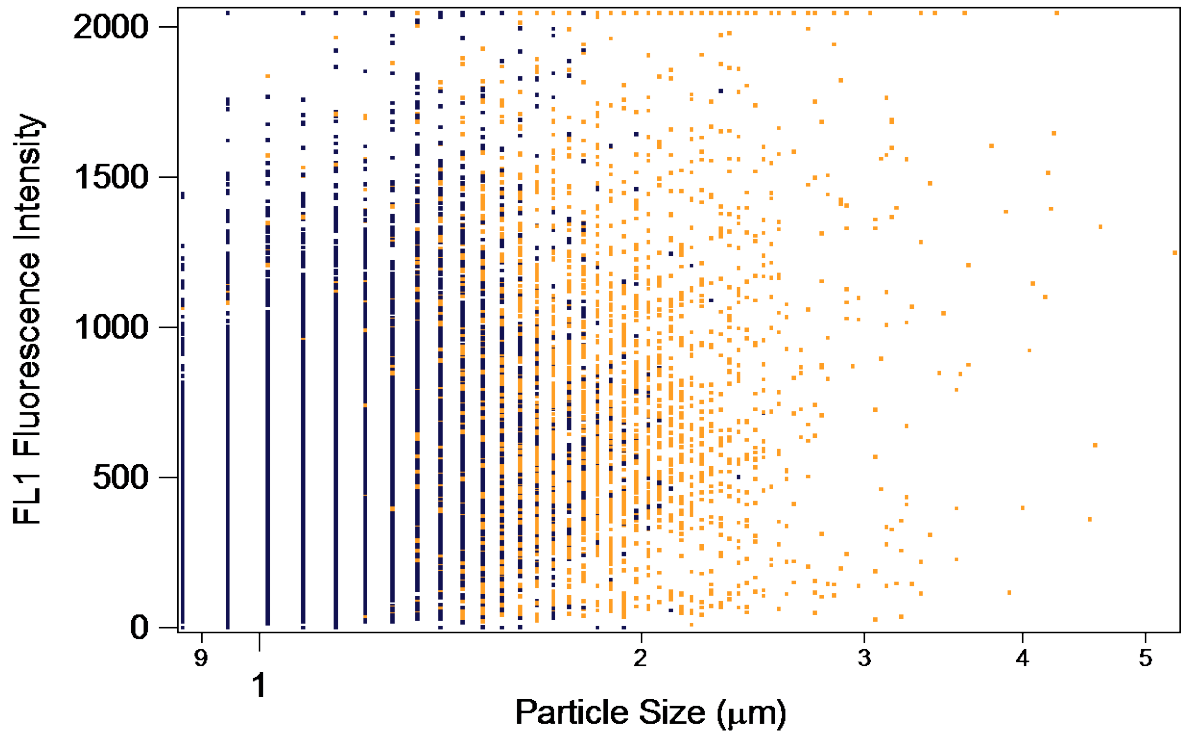
101 too low in concentration to influence the cluster properties. We added text including a summary  
 102 of these new experiments to the manuscript at L304:  
 103 *“To extend the investigation of particle input ratio, the three match-ups presented in Figure 3*  
 104 *were investigated using Scenario B with 1% bioparticles and 99% non-bioparticles in each*  
 105 *respective case. In these experiments the Bacteria:Diesel and Fungi:Dust particles separated*  
 106 *relatively well (6.6% and 13.5% misclassification, respectively). The Fungi:Diesel separation*  
 107 *was poor, however, because the Diesel particles were nearly evenly split into both clusters, and*  
 108 *the Fungi particles were too low in concentration to influence the cluster properties. More*  
 109 *investigation is needed to explore how extreme disparities in particle ratio could negatively*  
 110 *influence cluster quality in real-world settings.”*  
 111

		<b>Part A: Individual Clusters</b>					<b>Part B: Grouped Clusters</b>			<b>Part C: Summary</b>		
		(Particle Number)					(Particle Number)			(Cluster Quality)		
Fungi : Diesel	Cluster	B3	F2	S4	D12	Cluster	Bio	Non-bio	Total P.	Miscl.	Cat.	
	1	-	37	2588	-	1	37	2588	2625	98.6%	Fungi	
	2	-	0	1111	-	2	0	1111	1111	0.0%	Diesel	
Bacteria : Diesel	Cluster	B3	F2	S4	D12	Cluster	Bio	Non-bio	Total P.	Miscl.	Cat.	
	1	57	-	4	-	1	57	4	61	6.6%	Bacteria	
	2	0	-	5653	-	2	0	5653	5653	0.0%	Diesel	
Fungi : Dust	Cluster	B3	F2	S4	D12	Cluster	Bio	Non-bio	Total P.	Miscl.	Cat.	
	1	-	45	-	7	1	45	7	52	13.5%	Fungi	
	2	-	12	-	5650	2	12	5650	5662	0.2%	Dust	

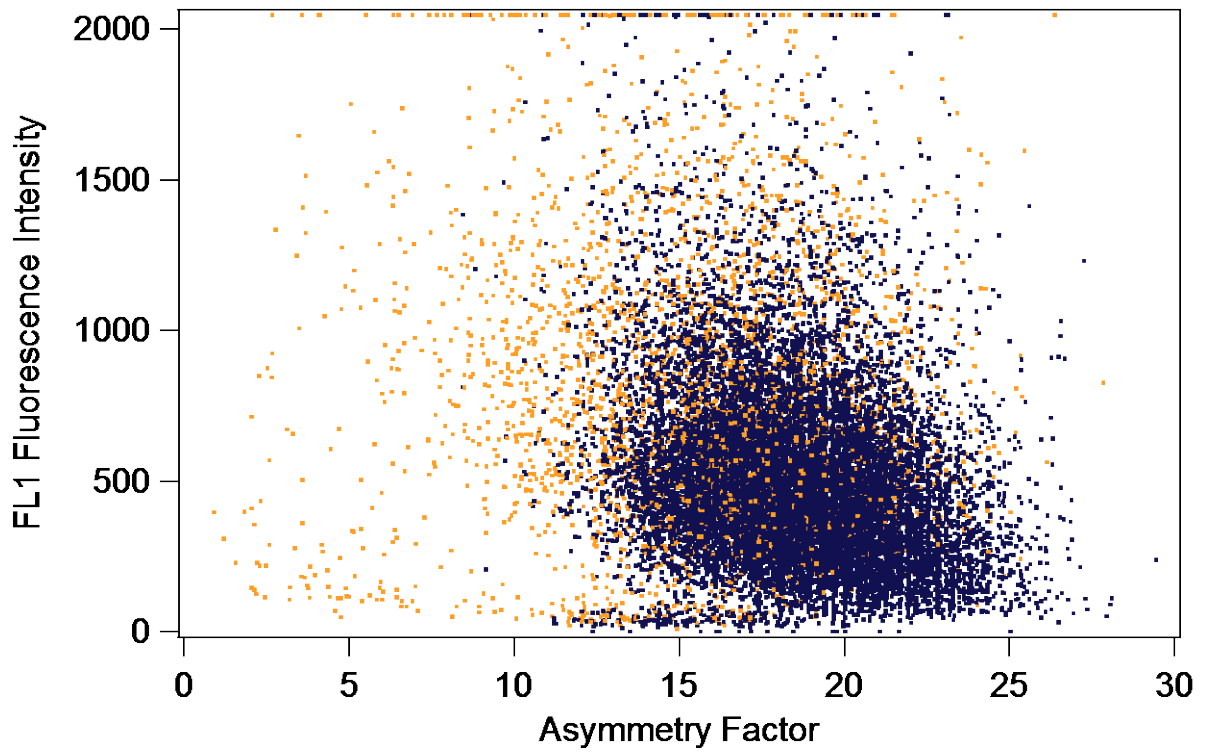
112  
 113  
 114 [R1.5] L238: Would it be possible to show examples of the cluster centroids for a case where there is  
 115 significant misclassification? This may illuminate why the algorithm is failing to correctly attribute  
 116 particles. It may also be useful to examine the fluorescence/AF characteristics of each cluster as a  
 117 function of size here. A 2D histogram or color density plot could show distinct hot spots that haven't been  
 118 separated correctly and could provide a basis for manual separation based on sensible thresholds.  
 119

120 [A1.5] To address the referee's suggestion, we included an additional set of plots here as  
 121 suggested. The results below correspond to the match-up between Bacteria 1 and Bacteria 3 using  
 122 Scenario B and the 3-sigma threshold, which corresponds to Experiment 22 from Table 2 (65%  
 123 misclassification). The two colors of dots in the plots represent clusters 1 and 2. In this case it is  
 124 still unclear how to utilize a single threshold to separate between the two particle types here.  
 125

126 In the process of analyzing results of this study we produced countless plots and tables, each of  
 127 which showed slightly different angles of the same story. We chose to simplify the results in  
 128 many cases to make the manuscript shorter and more manageably readable. We find that the table  
 129 of fluorescence intensity and AF median values (Table 2 from original data published in Savage  
 130 et al., 2017) often summarizes the differences in the particle types rather well and so were rarely  
 131 able to separate using 2D histograms as the referee suggests. One example of these two additional  
 132 plots is included here for reference, however.



133  
134



135  
136

137 [R1.6] L312-315: Can you describe the method for producing the soot as they seem rather large as  
138 compared to that in the study of Toprak and Schnaiter (2013) which were also coincidentally found to be  
139 weakly fluorescent in FL1. Perhaps the soot used in this study is larger and more fluorescent than we may

140 expect of ambient/urban soot which may cause some of the difficulty in correctly attributing in in some  
141 cases?  
142

143 [A1.6] The method for aerosolization of particle types discussed was presented in Section 3.2 of  
144 the associated Savage et al., 2017. Specifically, the aerosolization details related to soot are  
145 copied here:  
146

147 From Page 4284, Section 3.2.3 of Savage et al., 2017: “Dry powders were aerosolized by  
148 mechanically agitating material by one of several methods mentioned below and passing  
149 filtered air across a vial containing the powder. For each method, approximately 2.5–5.0  
150 g of sample was placed in a 10 mL glass vial. For most samples (method P1), a stir bar  
151 was added, and the vial was placed on a magnetic stir plate. Two tubes were connected  
152 through the lid of the vial. The first tube connected a filter, allowing particle-free air to  
153 enter the vessel. The second tube connected the vial through approximately 33 cm of  
154 conductive tubing (0.25 in. inner diam.) to the WBS for sample collection.”  
155

156 The referee is correct that the method of producing/aerosolizing particles, including soot, will  
157 bear heavily on the fluorescent properties observed. In particular, different aerosolization  
158 methods are likely to produce very different size distributions, which then will dictate the overall  
159 fluorescence properties. For this reason, we included the following statements in the Savage et al.,  
160 2017 paper:  
161

162 From Page 4292, Section 4.3: “It is important to note, however, that the method chosen  
163 for particle generation in the laboratory strongly impacts the size distribution of  
164 aerosolized particles. For example, higher concentrations of an aqueous suspension of  
165 particle material generally produce larger particles, and the mechanical force used to  
166 agitate powders or aerosolize bacteria can have strong influences on particle viability and  
167 physical agglomeration or fragmentation of the aerosol (Mainelis et al., 2005). So, while  
168 the absolute size of particles shown here is not a key message, the relative fluorescence at  
169 a given size can be informative.”  
170

171 The referee points out that the work by Toprak and Schnaiter (2013) presented small soot  
172 particles that also exhibited relatively weaker fluorescence in FL1. This is consistent with the  
173 expectation that fluorescence intensity will scale strongly with particle size. Differences in  
174 particle size could also impact clustering separation properties somewhat, and so further  
175 investigation of clustering using multiple narrow size ranges of different types of particles could  
176 further explore this process. This exhaustive process was beyond the scope of this work, however.  
177

178 To make sure these points are clear in the revised manuscript we have added the following text at  
179 L327:

180 *“It is also important to note here that the method of aerosolization for each particle type plays an*  
181 *important role in the observed size distribution and so results involving laboratory particles*  
182 *should be interpreted with this in mind. Observed fluorescence properties, in contrast, are*  
183 *expected to be conserved at a given particle size and intrinsically related to particle*  
184 *composition.”*  
185

186 [R1.7] L384: Would we expect to be able to differentiate between 2 different particles of the same type  
187 with such coarse spectral resolution?  
188

189 [A1.7] The referee’s implied point is correct. No, we would not expect to be able to separate  
190 between very similar types of particles using such coarse resolution as is available in the WBS.

191 Frankly, the fact that HAC paired with WIBS data was able to separate as well as it did was  
192 somewhat remarkable and surprising. To make the point clearer, we added text at the end of that  
193 paragraph as follows at L390:

194 “...separating more finely to quantify differences between types of individual biological particles  
195 is ~~likely to be~~ significantly more challenging *and not likely to be possible in most situations.*”  
196

197 [R1.8] L415: Again I wonder if the use of too coarsely separated bins may compromise the 9-sigma  
198 thresholding and cause misclassification?  
199

200 [A1.8] This question also loops back to [R1.3] and stems from a miscommunication. Values of  
201 the five WIBS data parameters were not separately binned (either during the logging process or  
202 when used as recorded), but are input into the cluster algorithm in the same spacing provided in  
203 the raw output of the instrument. The bin resolution is therefore limited by the WIBS optics and  
204 PMT settings.  
205

206 Further, fluorescence intensity is relayed by a integer units between 0 and 2047, and resolution is  
207 not a limiting factor. For example, see Figure 5 of the Savage et al. 2017 paper. Biological  
208 particles typically exhibit median fluorescence intensity much higher than non-biological  
209 particles, thus using different threshold strategies can help separate particle classes from one  
210 another by this strategy.  
211

212 [R1.9] L514: Can the authors comment on the applicability of their findings to new high resolution UV-  
213 LIF instruments that are beginning to become commercially available. Some of these new instruments  
214 have significantly more channels/greater fluorescent resolution than the WIBS.  
215

216 [A1.9] This is a helpful suggestion. To extend the applicability of results, the text was amended  
217 as follows:

218 “Results here are ~~only~~ generally extendable to other UV-LIF instruments, *however, whether they*  
219 *offer single or many channels of emission spectral resolution, in that the methods of particle pre-*  
220 *preparation and the impact of particle number ratio are likely to relay similar effects on*  
221 *clustering strategy.*”  
222

223 [R1.10] Technical corrections

224 L63: instruments, not instrument.

225 L370: grains, not gains.

226 L112: Suggest “Experimental and Computational Methods”

227 L131: “each of the three”

228 L181: “was the best”  
229

230 [A1.10] All typos corrected.