



# Technical Note: Bioaerosol identification by wide particle

# 2 size range single particle mass spectrometry

3

- 4 Xuan Li<sup>1,2</sup>, Lei Li<sup>1,2</sup>, Zeming Zhuo<sup>1,2</sup>, Guohua Zhang<sup>3</sup>, Xubing Du<sup>1,2</sup>, Xue Li<sup>1,2</sup>,
- 5 Zhengxu Huang<sup>1,2</sup>, Zhen Zhou<sup>1,2</sup>, and Zhi Cheng<sup>4</sup>
- <sup>1</sup>Institute of Mass Spectrometry and Atmospheric Environment, Jinan University,
- 7 Guangzhou, 510632, China
- <sup>2</sup>Guangdong Provincial Engineering Research Center for On-Line Source
- 9 Apportionment System of Air Pollution, Guangzhou, 510632, China
- <sup>3</sup>Guangzhou Institute of Geochemistry, Chinese Academy of Sciences, Guangzhou,
- 11 510640, China
- <sup>4</sup>Institute of Systems Engineering, Academy of Military Sciences, Tianjin, 300161,
- 13 China

14 15

Correspondence: Zhi Cheng(chengzh@npec.org.cn)

16 17

18

19

20

21

22

23

24

2526

27

28

29

30 31

32

Abstract The sources of bioaerosols are complex and diverse, which have a direct impact on the environment, climate, and human health. The effective identification of bioaerosols in the atmosphere is greatly significant for accurately obtaining the atmospheric chemical characteristics of bioaerosols and making biological early warnings and predictions. To improve the identification ability of bioaerosols, this study detected a variety of bioaerosols and abiotic aerosols based on a single particle aerosol mass spectrometry (SPAMS). Furthermore, the bioaerosol particle identification and classification algorithm based on the ratio of phosphate to organic nitrogen was optimized to distinguish bioaerosols from abiotic aerosols. The results show that 15 kinds of pure fungal aerosols were detected by SPAMS based on a wide range sampling system and that fungal aerosols with a particle size up to 10 μm could be detected. Through the mass spectra peak ratio method of PO<sub>3</sub>-/PO<sub>2</sub>- and CNO-/CN-, when discriminating abiotic aerosols, such as disruptive biomass combustion particles, automobile exhaust, and dust, from pure bacterial aerosols, the discrimination degree was up to 97.7%. The optimized ratio detection method of phosphate to organic nitrogen has strong specificity, which can serve as the discriminant basis for identifying bioaerosols in SPAMS source analysis or other analytical processes.

33 **Keywords** Bioaerosol; Single particle aerosol mass spectrometer (SPAMS); Online identification

36 37

38

39

40

41 42

43

44

45 46

47

48

49

50

51

52

53 54

55

56

57

58

59

60 61

62

63

64 65

66

67

68





### 1 Introduction

As a crucial component of atmospheric organic aerosols, bioaerosols participate in the weather and climate process as cloud condensation nuclei and ice nuclei (Fröhlich-Nowoisky et al., 2016). Moreover, some aerosols are human allergens, which pose a great threat to human health. At present, the importance of bioaerosols (Burrows et al., 2009) has been fully recognized; however, the sources of bioaerosols are sometimes difficult to identify, given their wide and scattered sources (Li et al., 2021), in addition to the obvious influence of meteorological conditions. This makes the identification of bioaerosols in the environment difficult (Rosch et al., 2006).

At present, laser-excited fluorescence spectroscopy is widely employed to detect bioaerosols (Li et al., 2018) due to its strong fluorescence signal, relative ease of operation, long-distance identification of bioaerosols and abiotic aerosols, and determination of single-molecule particle spectra. The fluorescent groups contained in bioaerosol particles are used for their detection in the fluorescence spectrometry method. However, since some inorganic minerals also fluoresce under ultraviolet light excitation, it is difficult to exclude the interference of abiotic fluorescent particles in the identification process. For instance, polycyclic aromatic compounds or humic acids can have similar fluorescence properties (Gabey et al., 2010) and cigarette smoke has similar fluorescence properties to bacteria (Hill et al., 1999). In recent years, the single particle mass spectrometry detection technology of bioaerosols has been developed rapidly, which can obtain the particle size information and chemical composition of single particles in real-time online. However, single particle aerosol mass spectrometry (SPAMS) also has its shortcomings in identifying environmental bioaerosols (Kleefsman et al., 2007). As phosphorus and nitrogen are components of nucleic acids and cell membranes, there is a large number of phosphate ions (PO, PO<sub>2</sub>-, and PO<sub>3</sub>-) and organic nitrogen ions (CN- and CNO-). Therefore, particles containing phosphate and organic nitrogen in the ambient air (such as biomass combustion products (Wei et al., 2019), fly ash, road dust (Yu et al., 2017), vehicle exhaust (Sodeman et al., 2005), and soil dust (Silva et al., 2000) are often confused with bioaerosols in the detection process. To improve the identification of bioaerosols, Zawadowicz et al. (2017) proposed a classification algorithm of spectral peak ratio based on PO<sub>3</sub>/PO<sub>2</sub> and CN-/CNO. When using particle analysis by laser mass spectrometry to discriminate dust and combustion by-products from pollen and bacterial aerosols, the degree of confidence is up to 98%. However, the research on the detection and discrimination of bioaerosols from fungi and other bioaerosols remains insufficient.

In addition, the particle size distribution of bioaerosols is generally  $0.3-100 \mu m$ , while that of viruses is less than  $0.3 \mu m$  (Smets et al., 2016). The typical particle size of bacteria is  $0.25-8 \mu m$ , while that of fungi is  $1-30 \mu m$ , among which the particle size that can cause harm to the human





body is generally between 0.4– $10~\mu m$ . For most of the existing SPAMS, the particle size analysis ability is about 0.1– $3~\mu m$ , thus the ability of SPAMS to detect fungi, spores, and other large particles is limited. Williams et al. (2013) designed a 7-stage aerodynamic lens (A-lens) to improve the ability of aerodyne aerosol mass spectrometry to detect biological particles. By optimizing the buffer cavity and increasing the sampling pressure of the lens, the transport efficiency of aerosol particles in the size range of 200–5000 nm can reach 100% but that of 10  $\mu m$  particles is only 22%. At the same time, Cahill et al. (2014) constructed a 7-stage A-lens for the transmission of a single particle in the range of 4–10  $\mu m$  and found that the transmission efficiency of 10  $\mu m$  particles was less than 20%, while that of 3  $\mu m$  particles was close to 0. ATOFMS is attempted to extend to the study of single-cell metabolomics.

To improve the ability of SPAMS to detect bioaerosols, a new sampling system (wide particle size range is 0.1–10 µm) was verified in the preliminary design. The particle transport efficiency can reach 100% in the particle size range of 0.15–10 µm. In this study, the advantages of high performance-SPAMS (HP-SPAMS) in the detection of bioaerosols were explored and 10 µm fungal particles were successfully detected. Furthermore, the discrimination method based on the ion ratio of bioaerosol characteristic peak was further optimized and verified, which could successfully discriminate bioaerosols from common disruptors, such as road dust, vehicle exhaust, and biomass combustion products, with a discrimination degree of 97.7%. In addition, this discrimination method had good statistical significance for bacterial aerosols. The analysis of single particle mass spectra is a hard ionization process and laser energy has little effect on the discrimination of this classification method. This method can be used as a discriminant basis for identifying bioaerosols in SPAMS source analysis or other analyses.

## 2 Experiment

#### 2.1 SPAMS

The constitution and working principle of SPAMS have been described in detail by Li et al. (2011). In short, SPAMS uses an aerodynamic lens to introduce aerosols from the atmosphere into a vacuum system and focus them into a collimating particle beam. Two successive laser beams are then used to measure the flight speed of particles and calculate their aerodynamic diameter in turn. The high-energy pulsed laser ionizes particles into positive and negative ions at the center of the ion source, which is detected by time-of-flight mass spectrometry.

In this study, the performance of HP-SPAMS was improved on the original basis and the resolution of the time-of-flight mass spectrometer was improved using delayed extraction technology (Li et al., 2018). The multi-channel superimposed signal acquisition system improves the sensitivity and dynamic range of instrument detection (Shen et al., 2018). In particular, for the





improvement of the sampling system (Zhuo et al., 2021), the whole sampling system consists of five modules: pre-focusing sampling interface, virtual impact concentration device, buffer cavity, 7-stage aerodynamic lens, and acceleration nozzle. The pore size was expanded from 0.11  $\mu$ m to 0.22  $\mu$ m. The numerical results show that the theoretical transport efficiency of particles could reach 100% in the particle size range of 0.15–10  $\mu$ m, which improved the sampling capacity of SPAMS for coarse particles and provided conditions for the detection of bioaerosols.

#### 2.2 Sample Determination

The 10 strains of bacteria and five strains of fungi determined in this study were standard strains provided by Hexin Kangyuan Medical Technology Co., Ltd. (Guangzhou). The specific names are shown in Table 1. Biological aerosol disruptors often found in the real environment, such as road dust, vehicle exhaust, and biomass combustion products (wheat stalk, corn stalk, and oblate leaf stalk), were selected as the research objects.

The preparation steps of the strain sample solution are as follows: 1) Sampling: first, the strains refrigerated at -80°C were taken out, thawed at room temperature for 1–2 hours, and vortexed using an oscillator to shake the centrifugation tube of the strain sample evenly. 2) Inoculation: on a clean laboratory table, the strain solution adhered to the disposable sterilized inoculating loop, and streak inoculation on the blood agar plate medium was performed. 3) Culture: the streaked culture medium was placed horizontally in a 37°C constant temperature box for about 24 hours. 4) Sampling: the growth of the samples of 15 strains after the culture is shown in Table 1. The colonies on the surface of the blood agar were slightly scraped with a disposable sterilized inoculating loop, dissolved with 1 mL deionized water in the centrifuge tube, and shaken well. 5) Dilution with water-soluble salt: the bacterial sample aqueous solution was centrifuged for 3 min at the rotation speed of 3,000–5,000 rpm. After centrifugation, the sample was precipitated at the bottom of the centrifuge tube and the aqueous solution was absorbed. Then, 1 mL of deionized water was added to dissolve the precipitate, followed by thorough shaking. Step 5 was repeated thrice.

The main components of blood agar plate medium used in this experiment were peptone, beef powder, sodium chloride, defiber sheep's blood, agar and deionized water. All media were autoclaved prior to use. Scrape only the upper layer of the culture medium surface to avoid small contaminants of the culture medium itself. Repeated rinsing with deionized water removes excess salt. It should be stressed that no additional fixatives or epoxies were added to the cells before analysis, reducing complications in the interpretation of the mass spectra.

The prepared pure bacterial sample solution was mixed with 20 mL of deionized water and atomized using a single nozzle aerosol generator (TSI Inc., Model 9302) to obtain the aerosol particles of the samples. A sheath gas of 80 kPa of clean air was used. The atomized sample





aerosol was connected to a silica gel drying tube, whose outlet was connected to the SPAMS inlet and an exhaust port with a high-efficiency particulate air filter. When sampling, 1,000 effectively ionized particle size, and spectrum data were stored in each sample. The experimental flow of HP-SPAMS is shown in Fig. 1.

143144

142

139

140 141

Table 1 Sample numbers and names of the 15 strains

Table 1 Sample numbers and names of the 15 strains						
number of	name	state of bacteria				
samples		or fungal				
#01	Klebsiella pneumoniae					
#02	Salmonella pneumoniae					
#03	Shiga virulent Escherichia coli					
#04	Bordetella bronchitis					
#05	Escherichia coli					
#06	Staphylococcus aureus					
#07	Listeria monocytogenes					
#08	Enterococcus faecium					
#09	Enterobacter cloacae					
#10	Staphylococcus epidermidis					
#11	Candida albicans					
#12	Candida tropicalis					
#13	Candida glabrata					
#14	Aspergillus brasiliensis					
#15	Saccharomyces cerevisiae					

145

147

148

149

150 151

152

153

154

155

156 157

158

159

160 161

162

163

164

165

166

167

168

169

170

171

172

173





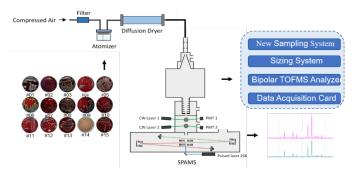


Fig. 1 Experimental flow of HP-SPAMS

#### 3 Results and Conclusions

#### 3.1 Distribution of bioaerosol particle size

The vacuum aerodynamic particle size distribution of the bacterial and fungal particles is shown in Fig. 2. Preliminary experimental results showed that the overall particle size of bacteria was relatively smaller, except for #08 Enterococcus faecium. The overall particle size distribution of bacteria was mainly within the range of 0.3-1 µm, thus showing an approximately normal distribution. Jung and Lee (2013) used scanning electron microscopy to observe Escherichia coli and Bacillus subtilis cells at room temperature, both bacteria with a diameter in the range of 0.5-0.7 µm and a length in the range of 1.1-1.6 µm. Fungi samples (#11, #12, #13, #14, and #15) obtained from pure strain cultures had a much larger proportion of particle size distribution above 1 μm than bacteria, and the particle size distribution of three fungi, Candida albicans, Candida glabrata, and Saccharomyces cerevisiae were around 0.5-2 µm. Sample #13 C. glabrata was concentrated in the range of 1–2.5 µm. Compared with other samples, the samples #11 C. albicans and #12 Candida tropicalis were similar to the samples #14 Aspergillus brasiliensis and #15 S. cerevisiae in terms of particle size distributions, while the particle size distribution of samples #11 and #15 was mainly in the range of 0.25-1.5 µm and the peak was around 0.4 µm. It is worth noting that the particle size of C. tropicalis and A. brasiliensis were evenly distributed between 0.1 μm and 10 μm. Li et al. (2020) used transmission electron microscopy and scanning electron microscopy to investigate primary biological aerosol particles (PBAPs) collected from boreal coniferous forests in the Xiao Hinggan Mountains of China in summer and speculated that the size of rod PBAPs was distributed at 1.4 µm and 3.5 µm and that the two typical peaks were bacterial and fungal particles, respectively.

Due to the low transmission efficiency of the aerodynamic lens for large particles and the tendency of large particles to produce inertial impinging wall loss in the process of air transport, especially large particles above 1 µm, the detected particle size distribution was smaller than the real one. Therefore, for the first time, HP-SPAMS was used to measure 10 µm coarse particulate

6 / 16

177

178

179180

181

182

183

184

185

186

187

188

189

190 191

192

193

194 195

196

197

198





matter under the improvement of the sampling system.

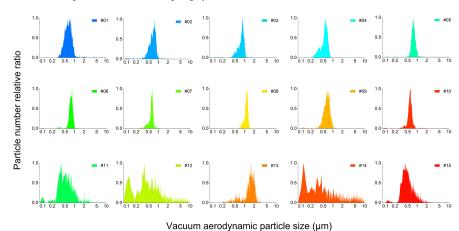


Fig. 2 Vacuum aerodynamic size distribution of 15 biological samples detected by SPAMS

## 3.2 Characteristic spectrum of bioaerosols

Zeng et al. (2019) used SPAMS to detect 13 strains of bacteria in 2019 to obtain similar bioaerosol characteristic ions; however, there were fewer characteristic peaks in the negative mass spectrum, and no negative ions with a mass charge ratio greater than 200 were detected, and the overall ionic peak signal was 50-10,000. HP-SPAMS detected the mass spectra of 15 strains of pure bacteria as shown in Fig. 3. Similar to the SPAMS detection results of the same type in the world, they all could effectively measure the phosphate and organic nitrogen ionic peaks of active bacterial aerosols, as well as some amino acid decarboxylic ionic peaks in the positive mass spectrum. Czerwieniec et al. (2005) found similar peaks when detecting vegetative cells of Bacillus atrophaeus and speculated that +30, +70, +72, +74, +86, +110, and +120 were decarboxylic ionic peaks of amino acids. The positive ion peaks were mainly 30[Glycine-COOH]+, <sup>70</sup> [Proline-COOH]<sup>+</sup>, <sup>72</sup> [Valine-COOH]<sup>+</sup>, <sup>74</sup> [Threonine-COOH]<sup>+</sup>, <sup>86</sup> [Leucine-COOH]<sup>+</sup>, <sup>110</sup> [Histidine-COOH]<sup>+</sup>, and <sup>120</sup> [Phenylalanine-COOH]<sup>+</sup>. Srivastava et al. (2005) speculated that +59, +81, +84, and +88 ionic peaks were organic fragments containing nitrogen, among which  $^{84}$  [C<sub>5</sub>NH<sub>10</sub>]<sup>+</sup> with a strong signal was also found in the detection of this study. The negative ionic peaks were mainly organic nitrogen <sup>26</sup>CN, <sup>42</sup>CNO, phosphate <sup>63</sup>PO<sub>2</sub>, <sup>79</sup>PO<sub>3</sub>, <sup>97</sup>H<sub>2</sub>PO<sub>4</sub>, <sup>159</sup>H(PO<sub>3</sub>)<sub>2</sub>, <sup>199</sup> NaH<sub>2</sub>P<sub>2</sub>O<sub>7</sub>, and other common biological ionic peaks.

More abundant ion characteristics were obtained by HP-SPAMS. On the original basis, the decarboxylic ionic peaks of serine and alanine <sup>44</sup> [Alanine-COOH]<sup>+</sup> and <sup>60</sup> [Serine-COOH]<sup>+</sup> were supplemented and the signal intensity in the HP-SPAMS detection results was relatively strong. The negative ion peaks <sup>261</sup>NaH(PO<sub>3</sub>)<sub>3</sub> and <sup>277</sup>NaH(PO<sub>3</sub>)<sub>2</sub>(PO<sub>4</sub>)<sup>-</sup> with a mass charge ratio greater than 250 were speculated and added. Exponential pulse delayed extraction technology (Chen et al.,





2020) not only solves the hit rate and resolution problems of SPMS but also improves the ion signal intensity by multiple times, thereby providing conditions for obtaining the complete mass spectrum characteristics of bioaerosols. More characteristic peaks can make it easier to distinguish whether or not a single particle is a bioaerosol.

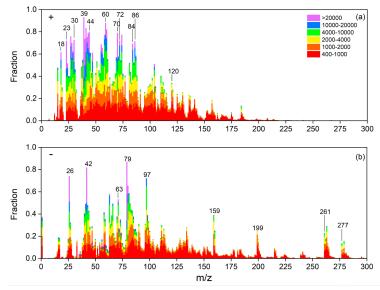


Fig. 3 Stacking diagram of the area of all bioaerosol ion peaks; (a) Positive mass spectrum; (b) Negative mass spectrum

## 3.3 Bioaerosol identification based on characteristic peak ratio

HP-SPAMS can accurately screen out bioaerosols according to their characteristic ions. However, due to the different intensity of ion signals in the single-particle spectrum, when the signal of the characteristic peak is weak, the spectrum will be ignored and the bioaerosols cannot be completely extracted. Zawadowicz proposed that <sup>26</sup>CN<sup>-</sup>, <sup>42</sup>CNO<sup>-</sup>, <sup>63</sup>PO<sub>2</sub><sup>-</sup>, and <sup>79</sup>PO<sub>3</sub><sup>-</sup> could be used as characteristic peaks of bioaerosol discrimination to distinguish bioaerosols from abiotic aerosols in a larger proportion. In the actual environment, many inorganic particles contain bioaerosol characteristic ion phosphate and organic nitrogen peaks. Organic nitrogen and phosphate ionic peaks with strong signals also appear in biomass combustion products, vehicle exhaust, and road dust measured by HP-SPAMS, as shown in Fig. 4. Therefore, when using phosphate and organic nitrogen ionic peaks alone to identify bioaerosols in the environment, at least 89% of vehicle exhaust, 49.5% of dust, and 58.3% of biomass combustion products have interference, which cannot be directly used as a sufficient condition to distinguish bioaerosols.

221

222

223

224

225

226

227

228

229

230

231

232

233

234

235

236237

238





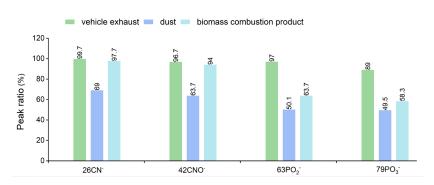


Fig. 4 Comparison of the frequency of four ion peaks in abiotic aerosols

Data classification based on HP-SPAMS, which is the peak height ratio of the mass spectra peak of PO<sub>3</sub>-/PO<sub>2</sub>- and CNO-/CN-, was used to distinguish bioaerosols from their disruptors. The scatter distribution of PO<sub>3</sub>/PO<sub>2</sub> and CNO CN was obtained by capturing the corresponding ionic peak height. As shown in Fig. 5, the distribution of bioaerosols was significantly different from that of biomass combustion products, vehicle exhaust, and dust. In particular, the scatter distribution positions of PO<sub>3</sub>-/PO<sub>2</sub>- and CNO-/CN<sup>-</sup> were significantly different from that of vehicle exhaust. Moreover, it was concentrated in a certain range of values. Bioaerosols were classified as one class, while aerosols produced by vehicle exhaust, dust, and biomass combustion products were classified as another class. The proportion interval of the bioaerosols PO<sub>3</sub>-/PO<sub>2</sub>- and CNO CN measured by SPAMS were concentrated at (3,200) and (0.7, 7), respectively, while those of abiotic aerosols were at (0.2, 3) and (0.02, 2), respectively. Furthermore, using the support vector machine (SVM), a supervised machine learning algorithm, the discrimination degree between bioaerosols and abiotic aerosols was up to 97.7%. This indicates that HP-SPAMS had a strong detection specificity on the phosphate and organic nitrogen ratio between bioaerosols and abiotic aerosols. It is possible to use it as the discriminant basis for identifying bioaerosols in the SPAMS source analysis or other analyses. Compared with the traditional single discrimination method via life characteristic elements, such as nitrogen and phosphorus, this method has a higher discrimination degree.





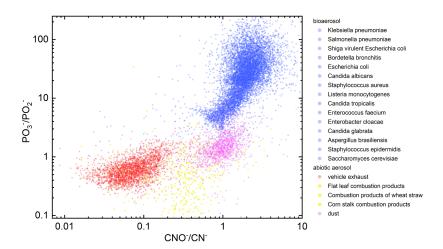


Fig. 5 Scatterplot of the bioaerosol and abiotic aerosol CNO-/CN- and PO<sub>3</sub>-/PO<sub>2</sub>-

The premise of this classification method is that there are four characteristic peaks. However, the actual research showed that when the threshold of the effective peak was set high, part of the weak signal peaks would be filtered out; however, when the threshold was set low, there was noise interference in the collected signals. Through a series of equivalent gradient threshold settings, the effective peak threshold of 10 mV was determined. The average frequency of the characteristic peaks in the bacterial aerosols was generally higher than that of the fungal aerosols, as shown in Table 2. At least, 82.9% of bacterial aerosols and 52.8% of fungal aerosols could be effectively discriminated against. The discrimination method based on the characteristic peak ratio had a high identification rate for bacterial aerosols. Fungi and bacteria have great differences in terms of morphology and structure. Bacteria are mainly coccus, bacillus, and spiral, while fungi are mainly subcellular and multicellular. In addition, fungi have nuclei. In the detection process, bacterial aerosols are more easily ionized to produce effective mass spectra peaks.

Table 2 The average frequency of the characteristic ionic peak of bioaerosol

Species	CN-	CNO-	PO <sub>2</sub> -	PO <sub>3</sub> -
Bacteria	92.9%	96.5%	82.9%	97.6%
Fungus	63.8%	70.4%	52.8%	75.3%

#### 3.4 Influence analysis of laser energy

SPAMS single-step laser desorption is a difficult ionization process in which organic compounds produce ion fragments of different degrees. Noble et al. (1996) proposed that in single particle mass spectrometry analysis, it is difficult to determine the morphology of organic compounds due to the extensive fragmentation caused by the ionization process. Cornwell et al.

10 / 16





(2022) proposed that the ion signals of dust and biological particles were very sensitive to ionization conditions and that total positive ion intensity was used to characterize the mass spectral relationship between different dust and biological particles. Through this method, environmental particles with both dust and characteristic biological spectra fingerprints were successfully excluded from the classification of biological particles. Liu et al. (2021) used *Bacillus thuringiensis* to explore the influence of different laser pulse energies on SPAMS and found that particles did not ionize when the laser energy was lower than 0.2 mJ and that the ionic peak increased significantly when the laser energy was higher than 1.5 mJ; they also found that the ionic peak integrity was the best when the laser energy was about 0.5 mJ. Too high or too low energy was not conducive to the discovery of the characteristic mass spectrum. To verify the influence of ionized laser energy on this analysis method, different laser energies of 0.5, 0.75, 1.0, 1.25, and 1.5 mJ were selected and *Staphylococcus aureus* was taken as an example to explore the influence of ionized laser energy on the ionic peak ratio. As shown in Fig. 6, as the energy increased, the ionization degree increased with more fragmented ions, the ratio of PO<sub>3</sub>-/PO<sub>2</sub>- and CNO-/CN- decreased, and more PO<sub>3</sub>-and CNO-were ionized as PO<sub>2</sub>- and CN-, respectively.

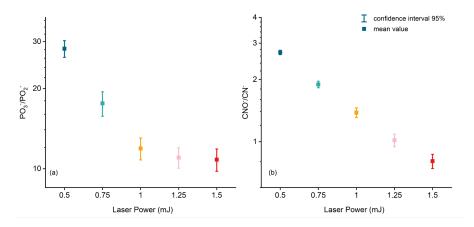


Fig. 6 Distribution of PO<sub>3</sub>-/PO<sub>2</sub>- (a) and CNO-/CN-(b)

Furthermore, the discrimination degree of ionized laser energy on bioaerosols under different ratios was compared. The biological aerosol (*S. aureus*) and abiotic aerosol (dust) were selected under the laser energy of 0.5 mJ. Using SPAMS, it was found that the ratio interval of PO<sub>3</sub>-/PO<sub>2</sub>- and CNO-/CN- was concentrated in (5, 70) and (0.8, 3), while that of dust aerosol was (0.7, 3) and (0.5, 2), respectively. When the laser energy was 1.5 mJ, the ratio interval of bioaerosols was (2, 30) and (0.7, 3), while that of dust aerosols was (0.6, 3) and (0.6, 2). As shown in Fig. 7, The interval of biological aerosols was gradually changing and the trends of horizontal and vertical coordinates were both decreasing, while the scatter interval of abiotic aerosols was almost unchanged. According to the SVM algorithm, the discrimination degree of bacterial aerosols and





dust under 0.5, 0.75, 1.0, 1.25, and 1.5 mJ energies were 96.6%, 97.4%, 97.1%, 96.5%, and 97.8%, respectively, indicating that the ionized laser had little effects on discriminating biological aerosols and dust disruptors.

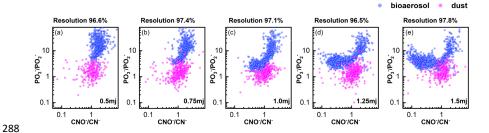


Fig. 7 Scatter distribution of the bioaerosol and dust aerosol CNO<sup>-</sup>/CN<sup>-</sup> and PO<sub>3</sub><sup>-</sup>/PO<sub>2</sub><sup>-</sup>

Under the condition of a constant effective peak threshold, the frequency of phosphate and organic nitrogen ionic peaks of bacterial aerosol and dust changed with the change of the ionized laser energy, as shown in Fig. 8. When the laser energy was 0.5 mJ, the peak output rate of both bioaerosol and dust was the lowest and the influence on abiotic aerosols was larger, with a peak output rate of 34%. When using this classification method for discrimination, it is only necessary to discriminate against 28.7% of dust particles. When the laser energy was 1.5 mJ, the peak output of four *S. aureus* ionic peaks was the highest, while that of the dust was the lowest. At this time, the highest proportion of bacterial aerosols (94.6%) and the lowest proportion of dust particles (31.1%) could be statistically discriminated against. Under the same laser energy, the overall peak output of abiotic aerosols was about 40% lower than that of biological aerosols. Different types of particles had different laser energy requirements. In addition, the variation trend of CNO<sup>-</sup> and PO<sub>3</sub><sup>-</sup> was the same as that of CN<sup>-</sup> and PO<sub>2</sub><sup>-</sup>, respectively; however, the phosphate ionic peaks (PO<sub>3</sub><sup>-</sup> and PO<sub>2</sub><sup>-</sup>) were more affected by the laser energy. In conclusion, when the ionized laser energy was 1.5 mJ, the classification method of the ionic peak ratio was more effective in discriminating bioaerosols.





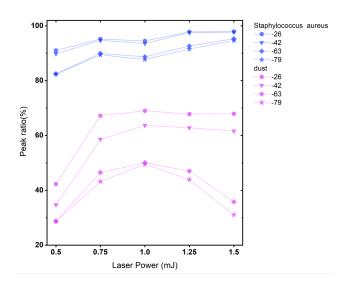


Fig. 8 Peak rate diagram of the characteristic peak output under different laser energies

### **4 Conclusions**

The performance of SPAMS and the improvement of the sampling system have improved the ability to identify bioaerosols. HP-SPAMS was used for the first time to detect fungal particles with a particle size of 10 μm, which provided a good technical basis for the detection of biological aerosols with large particle sizes in the environment. With the improvement of the instrument performance, the single-particle spectrum of bioaerosol showed decarboxylic ionic peaks of serine and alanine <sup>44</sup>[Alanine-COOH]<sup>+</sup> and <sup>60</sup>[Serine-COOH]<sup>+</sup> and phosphate ionic peaks <sup>261</sup>NaH(PO<sub>3</sub>)<sub>3</sub><sup>-</sup> and <sup>277</sup>NaH(PO<sub>3</sub>)<sub>2</sub>(PO<sub>4</sub>)<sup>-</sup>. A more unique fingerprint spectrum than the original study was obtained. The bioaerosol identification method based on the characteristic peak ratios PO<sub>3</sub>-/PO<sub>2</sub><sup>-</sup> and CNO<sup>-</sup>/CN<sup>-</sup> can effectively discriminate bioaerosol from three kinds of commonly seen abiotic disruptors, with the discrimination degree up to 97.7%. In addition, due to the influence of laser ionization efficiency, the effective mass spectra peak ratio of bacterial aerosol generation is higher, thus it is more suitable for this method. The ionized laser energy has a certain influence on the integrity of the ionic peak but hardly affects the identification accuracy of bioaerosols. This study showed that the SPAMS detection technology of bioaerosols has the potential to be a new method for real-time online identification of bioaerosols.

Data availability. These data can be publicly accessible in free.

Author contributions. LL and ZC designed the study; XL and ZMZ performed the experiments; GHZ, XBD, XL, ZXH and ZZ participated in data analysis and result discussion; XL and LL 13 / 16





328 wrote the paper with the input from all authors. 329 330 Competing interests. The authors declare that they have no conflict of interest. 331 332 Acknowledgements. We would like to thank engineer Huang Fugui of the Guangzhou Hexin Mass 333 Spectrometer Co., Ltd., for his technical support . 334 335 Financial support. This research has been supported by the National Natural Science Foundation 336 of China (grant no. 41905106). 337 338 339 References 340 341 Burrows, S. M., Elbert, W., Lawrence, M. G., and Poschl, U.: Bacteria in the global atmosphere - Part 342 1: Review and synthesis of literature data for different ecosystems, Atmos. Chem. Phys., 343 9,10777-10827, https://doi.org/10.5194/acp-9-9263-2009, 2009. 344 Cahill, J. F., Darlington, T. K., Wang, X., Mayer, J., Spencer, M. T., Holecek, J. C., Reed, B. E. and 345 Prather, K. A.: Development of a High-Pressure Aerodynamic Lens for Focusing Large Particles (4-346 10 µm) into the Aerosol Time-of-Flight Mass Spectrometer, Aerosol Sci. Technol., 48, 948-956, 347 https://doi.org/10.1080/02786826.2014.947400, 2014. 348 Chen, Y., Kozlovskiy, V., Du, X., Lv, J., Nikiforov, S., Yu, J., Kolosov, A., Gao, W., Zhou, Z., Huang, Z. 349 and Li, L., Increase of the particle hit rate in a laser single-particle mass spectrometer by pulse 350 extraction 13. 941-949. delayed technology, Atmos. Meas. Tech.. https://doi.org/10.5194/amt-13-941-2020, 2020. 351 352 Cornwell, G. C., Sultana, C. M., Petters, M. D., Al-Mashat, H., Rothfuss, N. E., Mohler, O., Demott, P. 353 J., Martin, A. C., and Prather, K. A.: Discrimination between individual dust and bioparticles using 354 aerosol time-of-flight spectrometry, Aerosol Sci. Technol., mass 355 https://doi.org/10.1080/02786826.2022.2055994, 2022. 356 Czerwieniec, G. A., Russell, S. C., Lebrilla, C. B., Coffee, K. R., Riot, V., Steele, P. T., Frank, M., and 357 Gard, E. E.: Improved sensitivity and mass range in time-of-flight bioaerosol mass spectrometry 358 using an electrostatic ion guide, J. Am. Soc. Mass Spectrom., 16, 1866-1875, https://doi.org/10.1016/j.jasms.2005.06.013, 2005. 359 Fröhlich-Nowoisky, J., Kampf, C. J., Weber, B., Huffman, J. A., Pöhlker, C., Andreae, M. O., 360 361 Lang-Yona, N., Burrows, S. M., Gunthe, S. S., Elbert, W., Su, H., Hoor, P., Thines, E., Hoffmann, T., Després, V. R., and Pöschl, U.: Bioaerosols in the Earth system: Climate, health, and ecosystem 362 363 interactions, Atmos. Res., 182, 346-376, https://doi.org/10.1016/j.atmosres.2016.07.018, 2016. 364 Gabey, A. M., Gallagher, M. W., Whitehead, J., Dorsey, J. R., Kaye P. H., and Stanley W. R.: 365 Measurements and comparison of primary biological aerosol above and below a tropical forest 366 canopy using a dual channel fluorescence spectrometer, Atmos. Chem. Phys., 10, 4453-4466, 367 https://doi.org/10.5194/acp-10-4453-2010, 2010.

Hill, S. C., Pinnick, R. G., Niles, S., Pan, Y. L., Holler, S., Chang, R. K., Bottiger, J., Chen, B. T., Orr C.

368





- S., and Feather, G.: Real-time measurement of fluorescence spectra from single airborne biological
- particles, Field Anal. Chem. Technol., 3,
- 371 https://doi.org/10.1002/(SICI)1520-6521(1999)3:4/53.0.CO;2-7, 1999.
- 372 Jung, J. H. and Lee, J. E.: In situ real-time measurement of physical characteristics of airborne bacterial
- particles, Atmos. Enviro., 81, 609-615, https://doi.org/10.1016/j.atmosenv.2013.09.018, 2013.
- 374 Kleefsman, I., Stowers, M. A., Verheijen, P. J. T., Wuijckhuijse, A. L. van, Kientz, C. E., and
- 375 Marijnissen, J. C. M.: Bioaerosol analysis by single particle mass spectrometry, Part. Part. Syst.
- 376 Charact., 24 ,85-90, https://doi.org/10.1002/ppsc.200601049, 2007.
- 377 Li, L., Liu, L., Xu, L., Li, M., Li, X., Gao, W., Huang, Z., and Cheng, P.: Improvement in the Mass
- Resolution of Single Particle Mass Spectrometry Using Delayed Ion Extraction, Int. J. Mass
- 379 Spectrom., 29 ,2105-2109, https://doi.org/10.1007/s13361-018-2037-4, 2018.
- 380 Li, L., Huang, Z., Dong, J., Li, M., Gao, W., Nian, H., Fu, Z., Zhang, G., Bi, X., Cheng, P., and Zhou,
- 381 Z.: Real time bipolar time-of-flight mass spectrometer for analyzing single aerosol particles, Int. J.
- 382 Mass Spectrom., 303 , 118-124, https://doi.org/10.1016/j.ijms.2011.01.017, 2011.
- 383 Li, X., Ran, B., Wu, W., Wang, Q., Tong, Z., Zhang, X., and Li, Y.: Progress in the application of
- fluorescence spectroscopy in biological aerosol monitoring, Mil. Med., 42 , 464-470,
- 385 https://doi.org/10.7644/j.issn.1674-9960.2018.06.015, 2018.
- 386 Li, W., Liu, L., Xu, L., Zhang, J., Yuan, Q., Ding, X., Hu, W., Fu, P., and Zhang, D.: Overview of
- primary biological aerosol particles from a Chinese boreal forest: Insight into morphology, size, and
- mixing state at microscopic scale, Sci. Total Environ., 719 , 137520-137533,
- 389 https://doi.org/10.1016/j.scitotenv.2020.137520, 2020.
- 390 Li, Y., Ma, T., Du, S., Xie, W., Zhang, H., and Xie, Z.: Review on source emission and source
- identification of bioaerosols in the atmosphere, J. Earth Sci. Environ., 43, 315-331,
- 392 https://doi.org/10.19814/j.jese.2020.11018, 2021.
- 393 Liu, C., Li, B., Liu, C., Li, M., and Zhou, Z.: Analysis of single-cell microbial mass spectra profiles
- from single-particle aerosol mass spectrometry, Rapid Commun. Mass Spectrom., 35, 9069-9078,
- 395 https://doi.org/10.1002/rcm.9069, 2021.
- 396 Noble, C. A., and Prather, K. A.: Real-Time Measurement of Correlated Size and Composition Profiles
- 397 of Individual Atmospheric Aerosol Particles, Environ. Sci. Technol., 30 , 2667-2680,
- 398 https://doi.org/10.1021/es950669j, 1996.
- 399 Rosch, P., Harz, M., Peschke, K. D., Ronneberger, O., Burkhardt, H., Schule, A., Schmauz, G., Lankers,
- 400 M., Hofer, S., Thiele, H., Motzkus, H. W., and Popp, J.: On-line monitoring and identification of
- 401 bioaerosols, Anal. Chem., 78, 2163-2170, https://doi.org/10.1021/ac0514974, 2006.
- 402 Shen, W., Dai, X., Huang, Z., Hou, Z., Cai, W., Du, X., Zhou, Z., Li, M., and Li, L.: Improvement of
- 403 the Dynamic Range of Data Acquisition System in Single particle mass spectrometry, J. Chinese
- 404 Cinemas., 39 ,331-336, https://doi.org/10.7538/zpxb.2017.0119, 2018.
- 405 Silva, P. J., Carlin, R. A., and Prather, K. A.: Single particle analysis of suspended soil dust from
- 406 Southern California, 34, 1811-1820. https://doi.org/10.1016/S1352-2310(99)00338-6, 2000.
- 407 Smets, W., Moretti, S., Denys, S., and Lebeer, S.: Airborne bacteria in the atmosphere: Presence,
- 408 purpose, and potential, Atmos. Environ., 139,214-221,
- 409 https://doi.org/10.1016/j.atmosenv.2016.05.038, 2016.
- 410 Sodeman, D. A., Toner, S. M., and Prather, K. A.: Determination of single particle mass spectral
- 411 signatures from light-duty vehicle emissions, Environ. Sci. Tech., 39 , 4569-4580,
- 412 https://doi.org/10.1021/es0489947, 2005.

https://doi.org/10.5194/egusphere-2022-598 Preprint. Discussion started: 26 July 2022 © Author(s) 2022. CC BY 4.0 License.





- 413 Srivastava, A., Pitesky, M. E., Steele, P. T., Tobias, H. J., Fergenson, D. P., Horn, J. M., Russell, S. C.,
- 414 Czerwieniec, G. A., Lebrilla, C. B., Gard, E. E., and Frank, M.: Comprehensive Assignment of Mass
- 415 Spectral Signatures from Individual Bacillus atrophaeus Spores in Matrix-Free Laser
- 416 Desorption/Ionization Bioaerosol Mass Spectrometry, Anal. Chem., 77 ,3315-3323,
- 417 https://doi.org/10.1021/ac048298p, 2005.
- 418 Wei, M., Xu, C., Xu, X., Zhu, C., Li, J., and Lv, G.: Size distribution of bioaerosols from biomass
- 419 burning emissions: Characteristics of bacterial and fungal communities in submicron (PM1.0) and
- 420 fine (PM2.5) particles, Ecotoxicol. Environ. Saf., 171 , 37-46,
- 421 https://doi.org/10.1016/j.ecoenv.2018.12.026, 2019.
- Williams, L. R., Gonzalez, L. A., Peck, J., Trimborn, D., McInnis, J., Farrar, M. R., Moore, K. D.,
- Jayne, J. T., Robinson, W. A., Lewis, D. K., Onasch, T. B., Canagaratna, M. R., Trimborn, A., Timko,
- 424 M. T., Magoon, G., Deng, R., Tang, D., Blanco, E. d. l. R., Prévôt, A. S. H., Smith, K. A., and
- Worsnop, D. R.: Characterization of an aerodynamic lens for transmitting particles greater than 1
- 426 micrometer in diameter into the Aerodyne aerosol mass spectrometer, Atmos. Meas. Tech., 6,
- 427 5033-5063, https://doi.org/10.5194/amtd-6-5033-2013, 2013.
- 428 Yu, N., Huang, B., Li, M., Cheng, P., Li, L., Huang, Z., Gao, W., and Zhou, Z.: Single particle mass
- spectrometry characteristics of atmospheric fine particulate dust sources, China Environ. Sci.,
- 430 37,1262-1268, https://doi.org/10.3969/j.issn.1000-6923.2017.04.008, 2017.
- 431 Zawadowicz, M. A., Froyd, K. D., Murphy, D. M., and Cziczo, D. J.: Improved identification of
- primary biological aerosol particles using single-particle mass spectrometry, Atmos. Chem. Phys.,
- 433 17,7193-7212, https://doi.org/10.5194/acp-17-7193-2017, 2017.
- 434 Zeng, Z., Yu, J., Liu, P., Huang, F., Chen, Y., Huang, Z., Gao, W., Li, M., Zhou, Z., and Li, L.: Analysis
- of bacterial aerosol particles by single particle aerosol mass spectrometer, Anal. Chem., 47,
- 436 1344-1351, https://doi.org/10.19756/j.issn.0253-3820.191202, 2019.
- 437 Zhuo, Z., Su, B., Xie, Q., Li, L., Huang, Z., Zhou, Z., Mai, Z., and Tan G.: Simulation design and
- 438 experimental study of aerodynamics particle concentrator for single particle mass spectrometry, J.
- 439 Vac. Sci. Technol., 41, 441-447, https://doi.org/10.13922/j.cnki.cjvst.202008026, 2021.

440 441