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# Chamber catalogues of optical and fluorescent signatures distinguish bioaerosol classes

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**Abstract.** Rapid bioaerosol characterization has immediate applications in the military, environmental and public health sectors. Recent technological advances have facilitated single-particle detection of fluorescent aerosol in near real time; this leverages controlled ultraviolet exposures with single or multiple wavelengths, followed by the characterization of associated fluorescence. This type of ultraviolet induced fluorescence has been used to detect airborne microorganisms and their fragments in laboratory studies, and it has been extended to field studies that implicate bioaerosol to compose a substantial fraction of supermicron atmospheric particles. To enhance the information yield that new-generation fluorescence instruments can provide, we report the compilation of a referential aerobiological catalogue including more than 50 pure cultures of common airborne bacteria, fungi and pollens, recovered at water activity equilibrium in a mesoscale chamber (1 m<sup>3</sup>). This catalogue juxtaposes intrinsic optical properties and select bandwidths of fluorescence emissions, which manifest to clearly distinguish between major classes of airborne microbes and pollens.

ously reporting from multiple bandwidths, ultraviolet induced fluorescence (UV-LIF) has successfully detected airborne microbes in bench-scale chamber studies (Healy et al., 2012; Kaye et al., 2005; Toprak and Schnaiter, 2013) and been applied to support large-scale aerosol monitoring campaigns where fluorescent particle cohorts indicated significant bioaerosol contributions to airborne organic carbon pools (Gabey et al., 2010; Pöschl et al., 2010; Gabey et al., 2013; Perring et al., 2015).

A variety of biogenic fluorophores have evolved in microorganisms and pollen grains, many of which are relevant to UV-LIF characterization of aerosols (Lakowicz, 2006; Poehlker et al., 2012). These fluorophores include metabolic mediators (e.g., nicotinamide adenine dinucleotide hydroxide (NADH) and its phosphorylated derivatives, NADPH) that are widely conserved throughout the microbial and plant worlds. Other fluorescently active biopolymers include a wide variety of structural proteins and pigments; however, these microbial compounds are tremendously variable in conformation and intracellular quantity (Kepner and Pratt, 1994; Madigan et al., 2008; Hill et al., 2014). As such, the distribution of fluorescence yields from some (whole-cell) microbial bioaerosols has been reported to have sensitivities to age (Kanaani et al., 2007) as well as cultivation history and environmental conditions (Saari et al., 2014; Uk Lee et al., 2010).

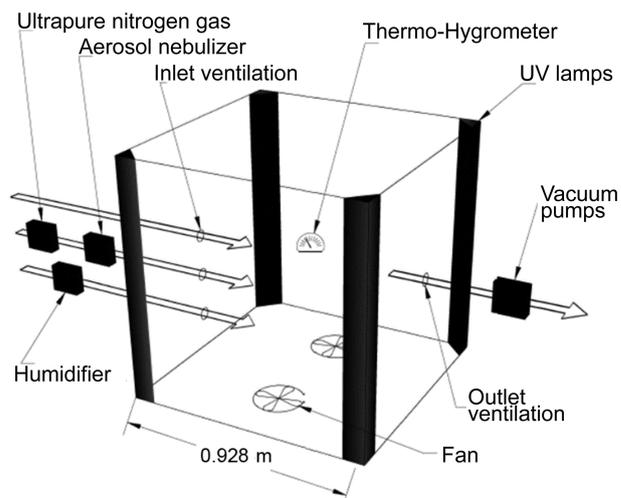
Sensitivity analyses of bioaerosol spectra from different UV-LIF instrument configurations are emerging in the litera-

## 1 Introduction

Fluorescence aerosol interrogation is gaining increased attention for its ability to characterize particulate matter suspended in both indoor and outdoor environments (Huffman et al., 2010; Sivaprakasam et al., 2011). When simultane-

ture (Agranovski et al., 2003; Huffman et al., 2010; Poehlker et al., 2012), as are generalized performance indices designated as apparatus-specific detection efficiencies (Saari et al., 2014). These UV-LIF metrics include, but are not limited to, photomultiplier thresholds, fluorescent particle fraction (FPF) recoveries and specific quantum yield comparisons. The calibration of modern UV-LIF instruments has been predominantly executed with thermoplastic spheres that are impregnated with colloidal metals or artificial fluorophores. While conventional microsphere calibrants are well characterized from a chemical perspective, they are poor biophysical analogues to the bioaerosols they are meant to reference, and they have sensitive stability windows limited to the order of weeks in cold, dark storage. In this context, no pure culture(s) of microbial bioaerosol, under any formally defined suite of cultivation and aerosol challenge conditions, has yet to be broadly recognized as a UV-LIF calibrant.

There is increasing interest in the utility of UV-LIF for environmental and autecological studies (Despres et al., 2012), yet there remains no unified approach for calibrating and cataloguing optical properties of fluorescence emissions associated with primary biological airborne particles (PBAPs). While many PBAPs fluoresce, interpreting UV-LIF measurements for bioaerosol characterization presents significant interpretive challenges, some of which have been successfully addressed using cluster analyses (Pinnick et al., 2013; Gabey et al., 2013; Crawford et al., 2015; Pan et al., 2015). In addition to classic Bayesian clustering approaches, referential methods can also be applied to analyze fluorescence distributions with selected optical properties, which may in turn be leveraged for environmental comparison(s). To this end, limited laboratory characterizations of some common bioaerosols have been introduced to different UV-LIF instruments, including single- and multiple-wavelength whole-cell microbial challenges in a variety of equipment configurations (Sivaprakasam et al., 2011; Healy et al., 2012; Saari et al., 2014). Some recent UV-LIF studies have demonstrated a library-based approach and specifically cite a need for expanding the database of intrinsic fluorescence properties associated with different PBAPs (Healy et al., 2012). In response, a unified approach for cataloguing intrinsic fluorescence from pure cultures of airborne bacteria (14), fungal spores (29) and pollen grain isolates (12) using a Wideband Integrated Bioaerosol Sensor (WIBS; Droplet Measurement Technologies, Boulder, CO, USA) is presented here. Using a mesoscale ( $1\text{ m}^3$ ) bioaerosol chamber, a methodical compilation of optical and fluorescence properties is reported at a defined humidity range, residence time and cell cycle. This serves as a referential model that can be reproduced, expanded and/or modified for economically cataloguing intrinsic bioaerosol fluorescence observations from UV-LIF instruments, under carefully controlled environmental conditions.



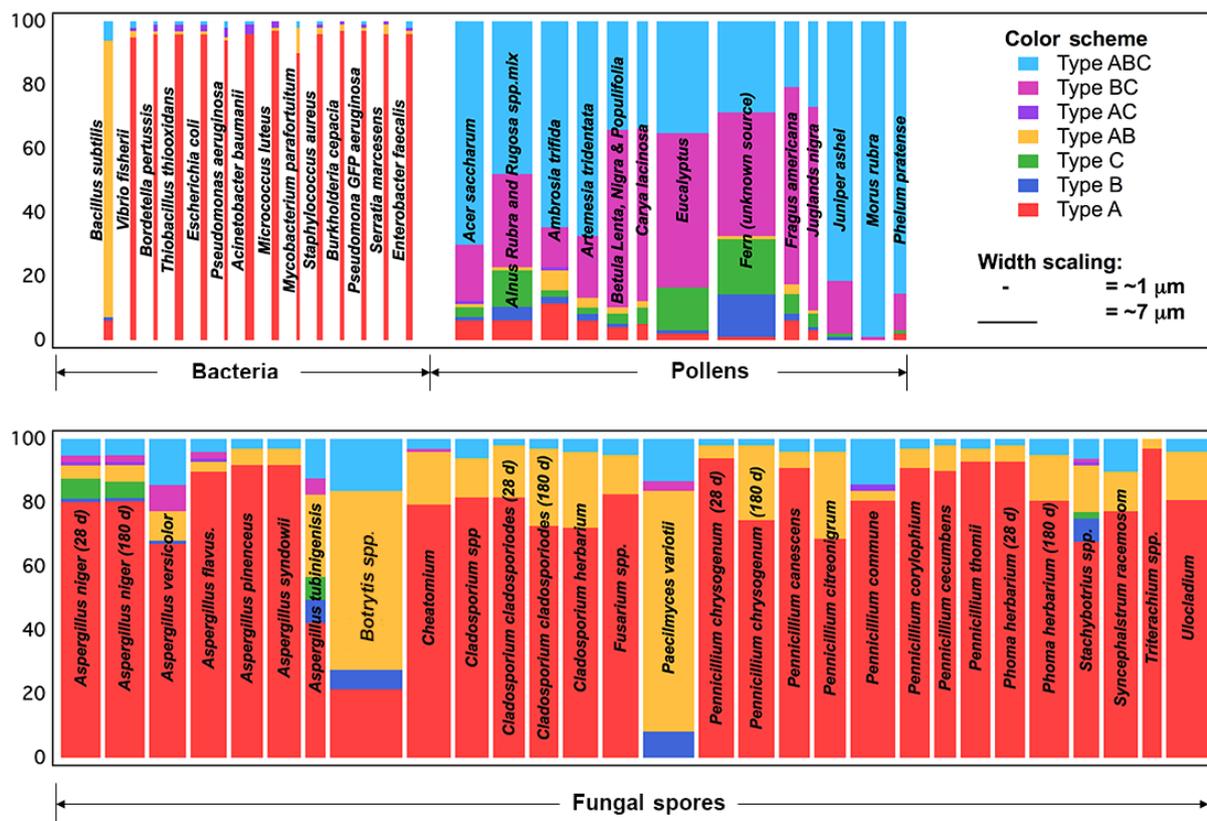
**Figure 1.** Schematic of bioaerosol test chamber. In independent trials, dry ultrapure nitrogen aerosolized and diluted different pure cultures of bacteria, fungi and pollens into a temperature ( $20\text{ }^{\circ}\text{C}$ )- and RH ( $25\text{--}40\%$ )-controlled chamber ( $890\text{ L}$ ). Bioaerosol delivery vessels were fitted with steel ports, and all gas streams were contained in  $0.4\text{ cm}$  conductive neoprene tubing, carrying  $151\text{ kPa}$  at a flow rate of  $12\text{ L min}^{-1}$  into the chamber.

## 2 Materials & methods

A cohort of light-scattering and fluorescent bioaerosol spectra were obtained from consecutively aerosolizing pure cultures of bacteria, fungi and pollens in an environmentally controlled chamber. The chamber into which all bioaerosol cultures were introduced is cubic (ca.  $1\text{ m}^3$ ) and constructed of grounded,  $1.25\text{ cm}$  (thick) Lucite; chamber air was continuously mixed by two  $1.5\text{ W}$  fans (Fig. 1). Comprehensive details of chamber construction and its operation for pure bioaerosol culture characterization have been previously described (Peccia et al., 2000). The WIBS used in this study was placed immediately adjacent to the outside chamber and continuously drew bioaerosol from the chamber interior at  $1.2\text{ L min}^{-1}$  through  $0.25\text{ m}$  of conductive neoprene tubing for these tests. The chamber was vented to a hood at ambient pressure (ca.  $88\text{ kPa}$ ); relative humidity (RH) was held between  $25$  and  $40\%$ ; temperature was held between  $20$  and  $22\text{ }^{\circ}\text{C}$ ; and bioaerosol was allowed to equilibrate to these conditions for more than  $5\text{ min}$  prior to UV-LIF sampling.

### 2.1 Bacteria

Fifteen pure bacterial cultures were grown into early stationary phase using standard protocols described by the American Type Culture Collection (ATCC) and German Collection of Microorganisms and Cell Cultures (DSMZ). These cultures, listed with their accompanying names (Table A1 in Appendix A) and fluorescence distributions (Fig. 2), have been used as models for the environmental behavior of airborne bacteria (Hurst et al., 2007). As judged by tracking the



**Figure 2.** Juxtaposition of optical and fluorescence properties of aerosolized pure cultures of bacteria and pollens (top), and fungal spores (bottom). Fluorescence type distribution is defined by excitation and emission from any of three possible channels alone (A, B or C) or in any combination. Bar width is proportional to equivalent optical diameter (scaling shown below color scheme legend in top panel).

optical density of the cultures (Madigan et al., 2008) upon entering stationary phase, bacteria were washed by sequential centrifugations in cold, sterile deionized water and immediately aerosolized into a chamber using a six-jet Collison nebulizer. Direct phase contrast and fluorescence microscopy (Hernandez et al., 1999) showed these bacteria remained intact after inspecting nebulizer reflux immediately following their aerosolization. Based on previous studies with many of the same bacterial models used here, some microbes, particularly Gram negative bacteria harvested in their log growth phase, can experience significant viability losses after being refluxed in Collison nebulizers even after 10 min (Stone and Johnson, 2002; Thomas et al., 2011). To minimize aerosolization stress and viability variance, bacterial cultures were harvested in early stationary phase, and nebulization times were limited to 2 min.

## 2.2 Fungi

Twenty-nine pure cultures of commonly occurring indoor fungi (Vesper et al., 2007) were cultured until they presented an obvious spore-bearing physiology, after being inoculated on malt extract agar (MEA) and held at 20 °C and 25 % RH. As judged by direct phase contrast microscopy, copious fun-

gal spores were obviously presented by these cultures between 18 and 28 days following their inoculation, depending on the species. These fungal spores, also presented in Fig. 2, were aerosolized directly off their host agar into the chamber using dry ultrapure nitrogen guided through sterile glass containments specifically designed for this purpose (modified fungal spore source strength tester, FSSST; Sivasubramani et al., 2004). Direct microscopy showed these fungal spores were intact and dominated the airborne biomass, with less than 1 % of supermicron bioaerosol particle volume as associated hyphae.

## 2.3 Pollen

Thirteen pure stocks of common temperate tree and grass pollens, collected under conditions similar to those reported by O'Connor et al. (2013), were obtained from a collection maintained at the Denver Botanic Garden. These pollen stocks, also listed in Fig. 2, were aerosolized into the chamber with the same FSSST configuration used for fungi, with notable fractionation of some grains as judged by direct phase contrast microscopy using accepted methods (Taylor et al., 2004).

To refine and expand upon previous UV-LIF PBAP cataloging efforts, we chose cohorts of pure cultures of bacteria, fungi and pollens that have been (repeatedly) recovered from aerosols in both indoor and outdoor environments as judged by culturing and microscopy. Of the 14 bacterial cultures used here, all but two are medically relevant to public health and/or bioterrorism (CDC, 2013), and eight cultures have (commonly) been used as bioaerosol models to their persistence under different atmospheric/disinfection conditions (e.g., UV; Peccia and Hernandez, 2006). The three bacterial cultures used here that have not been previously recovered from the atmospheric environment or otherwise used in prior bioaerosol studies (*Thiobacillus* sp., *Vibrio* sp. and *Enterobacter* sp.), were chosen to broaden the range of bacterial phenotypes used for fluorescence challenges such that all major bacterial physiologies were represented: Gram positive, Gram negative, bacterial spores, cocci, bacilli and vibrio (filamentous bacteria were purposely not included). With respect to the fungi and pollens chosen, the same logic applied in choosing which cultures to use for this demonstrative library (challenge): all members of the 26 fungal cultures used and all 13 pollen grain stocks have been commonly recovered from different atmospheric environments (Hurst et al., 2007; Vesper et al, 2007; O'Connor et al., 2013). While certainly not an exhaustive environmental inventory, these cultures cover a broad range of microbial physiologies commonly recovered from the atmospheric environment.

## 2.4 Optical spectra acquisition

The WIBS (v 4.0), previously described by Healy et al. (2012) uses dual-wavelength excitation and fluorescence detection, while simultaneously measuring characteristic dimensions from scattered light. The WIBS variant used here (version VI) detects fluorescence of airborne particles by sequential exposure to UV irradiation from a flashlamp filtered at 280 and 370 nm. Fluorescence emitted due to 280 nm excitation was detected in two wavebands, 310–400 and 420–650 nm, using dedicated photomultipliers. Fluorescence emitted due to 370 nm excitation was detected between 420 and 650 nm. Optical diameter was determined by light scattered from exposure to a 635 nm laser; it is reported here as an equivalent optical diameter (EOD), defined as the diameter of a spherical particle, with a fixed refractive index (relative to that calibrated with 2.0 and 2.8  $\mu\text{m}$  latex beads in air between 25 and 45 % RH), scattering the same light intensity as the measured bioaerosol. These optical properties were recovered from a minimum of between  $10^3$  and  $10^5$  cells, fungal spores or pollen grains. Each pure culture aerosolized was held in the chamber long enough to be at equilibrium with the contained humidity condition (>5 min; Peccia et al., 2001), and bioaerosol was collected over at least a 5 min period, when operating in the following gain modes: fungi and pollens, low gain; bacteria, higher gain. Between each aerosolization challenge, the chamber was evacuated using a

high-volume HEPA filter such that total particle counts were below  $10^2 \text{ m}^{-3}$ , and the chamber was subsequently purged with ethanol vapor and/or ozone while illuminated with UV light.

Following the annotation introduced by Perring et al. (2015), particle fluorescence was categorized as one of seven types, which considers each of three fluorescence bandwidths individually, as well as in all possible combinations. Here the subscripts denote excitation wavelengths, and parentheses indicate the emission bandwidths observed (nanometers):

TYPE A<sub>280</sub> = (310 – 400); TYPE B<sub>280</sub> = (420 – 650);

TYPE C<sub>370</sub> = (420 – 650).

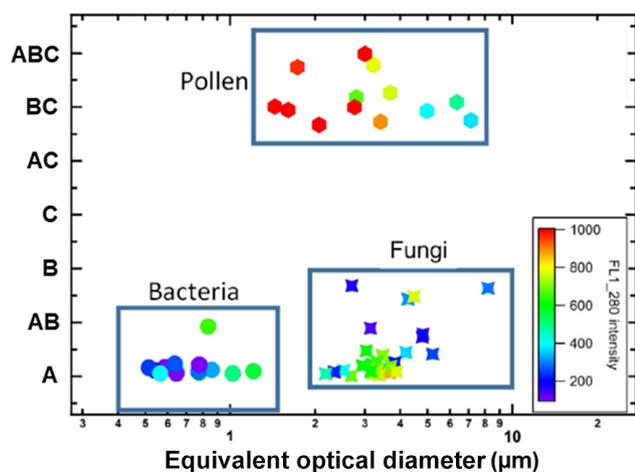
While this notation departs from the fluorescence emission notations used in studies previously describing WIBS performance, it provides a convenient symbolization system for visualizing and discussing fluorescence as described by distributions of emissions. Leveraging these measured quantities, the following metrics were analyzed and compiled for each bioaerosol: (i) the frequency of particles that could be segregated by fluorescence signal (bandwidth) into any of the seven types; (ii) the average fluorescence intensity within each bandwidth (if any); and (iii) the average EOD of each particle type. As is customary with interpreting UV-LIF data (Healy et al., 2012), we report results where a mean forced trigger value plus 3 standard deviations was used as the minimum data recovery threshold.

To demonstrate detection of the potential effects aging can have on intrinsic fluorescence emissions under these conditions, airborne spores of *Phoma* spp., *Penicillium chrysogenum*, *Cladosporium cladosporioides*, and *Aspergillus versicolor* that were less than 28 days old were compared to the spectra of spores from these same cultures, which were allowed to age 180 days (raised and maintained under identical environmental conditions).

## 3 Results & discussion

### 3.1 Composite optical recognition patterns of bioaerosol classes

The seven fluorescence categories, the fluorescence intensity and the relative size expressed by the EOD provide a multi-dimensional matrix from which composite signatures can be constructed for the three bioaerosol classes tested here. Figure 2 shows the EOD and the distribution of fluorescence type for each of the cultures studied, while Fig. 3 shows the normalized fluorescence intensity emitted and composite weighting of their distributions considering all fluorescence types. As judged by EOD and fluorescence type distributions, the pure bioaerosol cultures used to challenge this WIBS distinctly assembled into their major respective physiologic groups. All of the bacterial cultures aerosolized as



**Figure 3.** Assemblages of equivalent optical diameter and fluorescence type from aerosolized pure cultures of bacteria (●), fungi spores (X) and pollen grains (◆). The fluorescence type reported is a weighted composite of fluorescent type occurrence observed from each culture or pollen stock aerosolized; intensity is a relative scale.

vegetative cells assembled with common fluorescence patterns below an EOD of  $1.5\ \mu\text{m}$  and presented the weakest fluorescence intensities of the bioaerosols observed. With the exception of spore-forming *Bacillus subtilis*, bacterial bioaerosol was dominated by a single fluorescence type (A). The fungal spores aerosolized were in a markedly higher range with respect to their equivalent optical diameters ( $2\text{--}9\ \mu\text{m}$ ) but presented several fluorescence types: A, AB, BC and ABC, with a prevalence of A and AB. While the pollens aerosolized were overlapping in equivalent optical diameter ranges with some of the fungal spores observed, they presented significantly different fluorescence type distributions, which were dominated by combinations of fluorescent types BC and ABC. Note that many of the pollen EODs presented here are considerably smaller than the true diameter of their nascent intact-grain size, which is attributed here to pollen fragmentation during collection, storage or aerosolization. Because of this phenomenon, the EOD distributions for pollen grains are much wider than their fungal or bacterial counterparts, and the means of these distributions are less than unadulterated pollen grains, some of which have true optical diameters larger than  $10\ \mu\text{m}$ . These results are consistent with laboratory and environmental studies which indicate that pollen grain fragmentation routinely occurs in the atmosphere (D'Amato et al., 2007; Miguel et al., 2006; Taylor et al., 2002, 2007). The pollens assembled together based on the ascendancy of type C fluorescence and their markedly higher fluorescence intensities relative to the other bioaerosol classes observed (Fig. 3).

As judged by WIBS, the multi-channel fluorescence response observed from the seven *Penicillium* species aerosolized here are in general agreement with those of

the *Penicillium notatum* by Toprak and Schnaiter (2013); they reported fluorescence emissions from *P. notatum* commensurate with those observed here (dominance of type A, with ascendancy of type C), where these aerosolized microbes were mixed with ammonium sulfate aerosol in small-chamber studies. Pan et al. (2010) also reported fluorescence (type B) associated with aerosolized *B. subtilis* spores similar to that reported here. The spectral distributions of *B. subtilis* spores reported here, and by Pan et al. (2010), suggest that these aerosolized spores had markedly different fluorescence than that observed from vegetative bacterial cells with nearly the same EOD. Although they did not aerosolize into bioaerosol chambers which facilitate water activity equilibrium, Pan et al. (2011) also reported a limited catalogue of dual-wavelength fluorescence spectrometer (DWFS) observations by directly introducing 12 dry pollen stocks and four different types of fungal spores to the optical chamber of a DWFS. Of those pollens that overlapped with the genera of the pollen grains used here (*Betula* spp., *Ambrosia* spp.) and of those fungal cultures which were the same as those reported here (*Aspergillus flavus*, *Cladosporium herbarum* and *Penicillium chrysogenum*), there were general similarities in fluorescence distributions such that these fungal spore expressed more dominance of type A fluorescence, while the pollens expressed ascendancy of type C fluorescence.

In a cataloguing approach similar to that reported here, Healy et al. (2012) challenged a WIBS 4 with seven different types of fungal spores and eight different pollen grain stocks in a bench-scale chamber under mid-range humidity conditions (ca. 54%). As was the case here, the pollens aerosolized by Healy et al. (2012) presented a much broader diversity of fluorescence emissions recovered across the three fluorescent WIBS channels when compared with the two *Cladosporium* spp. they aerosolized; however, the same fluorescence emission pattern was not true when juxtaposing fluorescence observations with that of the other airborne fungi they tested. When comparing fluorescence response of the seven *Aspergillus* and eight *Penicillium* species tested here to the response of the single *Aspergillus* (*A. fumigatus*) and *Penicillium* (*P. notatum*) species reported by Healy et al. (2012), there were significant differences in fluorescence, particularly with respect to the prevalence of type C fluorescence. Because chamber volume and bioaerosol residence time were undisclosed, it was unclear if the bioaerosol challenges executed by Healy and coworkers were at water activity equilibrium with the chamber humidity. This physical condition, the aerosolization scheme, chamber operations and the limited number of fungi cultures observed may account for these differences. Indeed, Healy and coworkers cited the need for “further studies to generate and add to the current library/database of WIBS-4 particle signatures”; in part, this study responds to that prescribed need and refines culturing, aerosolization and (larger) environmental chamber conditions, and, in particular, equilibrium water activity with respect to the bioaerosol observed.

### 3.2 Fluorescence response to spore aging

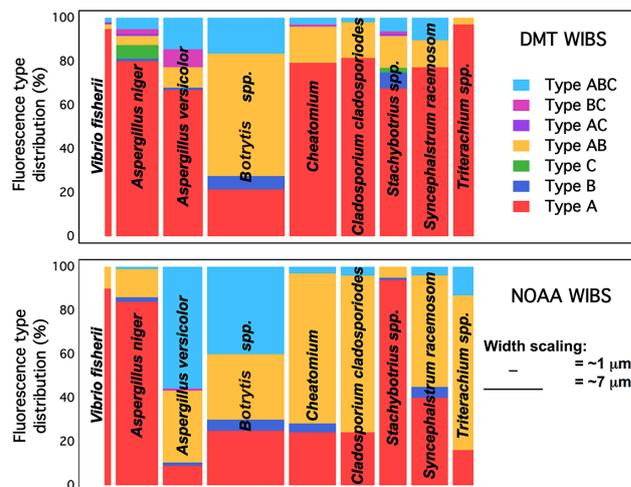
Fluorescence spectra from four pairs of young and aged fungal spores were compared through these analyses (aged 28 vs. 180 days). Of these, two presented no discernable change in the EOD and fluorescence distributions used as cataloguing metrics here: *Cladosporium cladosporioides* and *Aspergillus niger*. There was, however, a significant shift in the fluorescence distribution, from younger to older spores, of the species *Penicillium chrysogenum* and *Phoma herbarium* (Fig. 2 bottom). More than 90% of the spores of these species presented their fluorescence as type A when aerosolized at an age of 28 days, but this decreased to near 70% after 180 days of culture aging under identical conditions, with the balance becoming type AB.

### 3.3 Instrumental variability and gain considerations

We present here the response of a single WIBS instrument to pure cultures of bioaerosols grown, collected and aerosolized under carefully controlled conditions. The instrument manufacturer (Droplet Measurement Technologies) typically sets detector gains based on photomultiplier fluorescence detection from monodisperse challenges with commercially available fluorescent microspheres (polystyrene latex, PSL), the intensity of which is known to vary between manufacturing batches. The fluorescence of PSLs also significantly degrades on the timescale of weeks to months (even with proper cold storage). While this strategy provides general consistency across different instruments, the lack of an absolute calibration means that there can be appreciable variability in fluorescence signal recovery of a particular bioaerosol between one UV-LIF instrument and another. In response, we aerosolized a subset of test bioaerosol cultures to challenge two WIBS in parallel: one owned by the manufacturer, which was used for the larger set of measurements presented here, and another owned by the Chemical Sciences Division of the National Oceanic and Atmospheric Sciences Administration (NOAA). We find differences in spectral classification (Fig. 4), likely attributable to differing gain settings, with the NOAA WIBS recording a larger fraction of a given particle population as having signal above threshold in the B and C channels (both of which are detected using the same photomultiplier) than does the manufacturer's market issue WIBS. This operational example highlights a condition that different WIBS instrumentation, while configured the same and challenged under the same conditions, can report somewhat different fluorescence distributions.

### 3.4 Implications for real-time detection of atmospheric bioaerosol

Given the potential variability introduced with culture age and between instruments, genera- or species-type classifications for different bioaerosols should not be treated as ab-



**Figure 4.** Comparison of fluorescence type distributions for two different WIBS instruments from a subset of the pure bioaerosol cultures and pollen grains used in this analysis. The top panel shows distributions from the DMT WIBS, which was used for all of the bioaerosol chamber challenges presented here. The bottom panel shows the fluorescence type distributions from the NOAA WIBS, used in parallel for a subset of the bioaerosol chamber challenges using pure cultures of bacterial cells and fungal spores.

solute “signatures” but, rather, as likely assemblages that, when considered in conjunction with EOD, allow for discrimination between broad bioaerosol categories. There is also particle-to-particle variability in the type manifestation of a given sample (i.e., not all particles of a given species present the same spectral type) that makes reliable identification on a single-particle basis unlikely and requires, instead, a statistical treatment. For example, bacterial populations tend to present as mostly type A at small (< 1 μm) EOD. Fungi present a mixture of type A, AB and ABC fluorescence at larger EOD sizes (2–9 μm), although the distribution between the dominant types may differ substantially from that presented here, depending on instrumental parameters, culture circumstances and environmental conditions (e.g. RH, *T*). Finally, pollen tends to present a mixture of C, BC and ABC and is the only class of bioaerosol for which types C or BC are significant contributors. This facilitates discrimination between fungi and pollen despite the fact that the likelihood of pollen fragmentation reduces confidence in the utility of EOD as a discriminating parameter. Another distinguishing feature of pollens is their relatively high fluorescence intensity, although we note that very little is known about the temporal evolution of fluorescent intensity over atmospherically relevant timescales of photochemical processing.

Type A was dominant in all bacterial and most fungal populations aerosolized in this study (at least for certain gain settings, like those in the DMT WIBS), and thus EOD is an important consideration in distinguishing bacteria from fungi.

Under laboratory conditions constrained by this work, we typically sampled single, intact, bacterial cells. Atmospheric observations of bacteria and DNA in aerosol, however, indicate that bacteria are frequently found associated with particle sizes larger than that of a single cell. (Burrows et al., 2009; Shaffer and Lighthart, 1997; Tong and Lighthart, 2000; Wang et al., 2008). It is hypothesized that this is due to the presence of either multicell bacterial agglomerates or particle mixtures including bacteria and dust or other vegetative detritus (Bovallius et al., 1978; Lighthart, 1997), and we acknowledge that such agglomerates could be easily confused with intact fungal spores given their spectral similarities as detected by the WIBS. Another notable observation of the ensemble of experiments presented here is the lack of an appreciable fraction of type B particles associated with these pure bioaerosol culture challenges. Type B particles are detected as a minor fraction (up to 15 %) in only a few fungal and pollen species; therefore, any atmospheric sampling in which type B is a dominant fluorescent type should be examined carefully for potential (non-biological) interferents. Type AC was also strikingly absent from the measurements presented here.

#### 4 Conclusions

Recent advances in UV-LIF instrumentation have led to a new generation of atmospheric particle sensors with promising utility for rapid characterization of fluorescent bioaerosols, both indoors and out. Here we have demonstrated the ability of this technique to characterize whole, airborne microbial cells and pollen grains in a relational capacity by refining a chamber-based library approach with environmental conditions controlled at a mesoscale.

Where conventional light-scattering analyses are coupled with the spectral analyses of fluorescence emission(s), a powerful combination of phenotypic information becomes available on a high-throughput, single-particle basis: the broader the excitation and emission spectra observed, the higher the potential for the fundamental characterization of (bio)aerosol in situ. As has been previously described, the WIBS technology employed here demonstrates the potential of multi-channel fluorescence analyses in a laboratory setting; however, this work has refined the approach previously reported by Healy et al. (2012) for expanding existing fluorescence emission catalogues through a simple, unified reference method created under defined chamber conditions. Such a library can be reproduced, expanded and shared by users for interlaboratory comparisons and/or associations with field observations.

In the course of this analysis, the composite evaluation of optical diameter and fluorescent spectra associated with airborne fungal spores, pollens and bacteria revealed that these primary physiologies can be unambiguously differentiated from each other when airborne under defined conditions (culture condition, RH and temperature). While deeper cluster analyses were beyond the scope of this demonstrative report, within the fungi and pollens there emerged optically defined phenotypes, which also formed distinct sub-assemblages – an exciting cataloguing outcome with implications for future algorithmic developments and the potential for confident identification of “interferences” which may complicate bioaerosol quantitation in the field (Despres et al., 2012; O’Connor et al., 2013; Poehlker et al., 2012).

The identity of fluorescent aerosol and the fraction thereof composed of primary biological materials remain unknown in many different environments (Despres et al., 2012; Heald and Spracklen, 2009). To this end, these findings suggest that a composite analysis of wideband UV-LIF, as referenced to robust bioaerosol libraries, offers a promising approach to help better characterize airborne particulate matter in the laboratory as well as during wide-area environmental surveillance.

#### 5 Data availability

All data (fluorescence counts) recorded, used and analyzed in this investigation are presented in Table A1. Raw WIBS data files can be obtained by request from Droplet Measurements Technologies (darrel.baumgardner@gmail.com).

## Appendix A

A table containing discrete EOD and intrinsic fluorescence values observed for each culture aerosolized (used to construct Figs. 2 and 3) is presented here.

**Table A1.** Summary of properties observed for airborne bacteria, fungi and pollens.

Genus species ATCC or DSMZ no.	n	EOD	$\sigma$	Intensity	Fluorescence type frequency						
					A	B	C	AB	AC	BC	ABC
Bacteria											
<i>Bacillus subtilis</i> 6051	19786	0.7	0.3	656	11	2	0	87	0	0	0
<i>Vibrio fischeri</i> 7744	15073	0.7	0.4	581	97	0	0	1	2	0	0
<i>Bordetella pertussis</i> 9797	12834	0.5	0.4	84	93	0	1	1	3	0	1
<i>Thiobacillus thiooxidans</i> 19377	10100	0.9	0.4	178	94	0	0	1	3	0	2
<i>Escherichia coli</i> K-12	11900	0.9	0.4	265	95	0	0	1	3	0	1
<i>Pseudomonas aeruginosa</i> 10145	28190	0.4	0.1	248	94	0	0	1	3	0	2
<i>Acinetobacter baumannii</i> 19606	30006	1.1	0.5	556	96	0	0	0	3	0	1
<i>Micrococcus luteus</i> 4698	28197	0.9	0.4	502	97	0	0	1	2	0	0
<i>Mycobacterium parafortuitum</i> 19686	18197	0.4	0.2	265	90	0	0	8	0	0	2
<i>Staphylococcus aureus</i> 12600	30015	0.7	0.2	283	96	0	0	2	1	0	1
<i>Burkholderia cepacia</i> 25416	30003	0.5	0.2	399	97	0	0	2	1	0	0
<i>Pseudomonas aeruginosa</i> GFP 15692	16015	0.6	0.3	112	97	0	0	1	1	0	1
<i>Serratia marcescens</i> (DSMZ 3021)	30027	0.6	0.2	393	96	0	0	3	1	0	0
<i>Enterococcus faecalis</i> 19433	17401	0.8	0.3	327	96	0	0	1	1	0	2
Fungi											
<i>Aspergillus niger</i> (28 days) 16888	30772	4.4	2	157	79	2	7	4	2	2	5
<i>Aspergillus niger</i> (180 days) 16888	23086	4.4	2	181	78	2	5	5	2	2	5
<i>Aspergillus versicolor</i> 9577	32620	4	2.1	304	67	2	0	9	0	8	14
<i>Aspergillus flavus</i> 16883	15847	4	1.3	364	87	0	0	3	1	2	4
<i>Aspergillus pinenceus</i> (sp. proposed)	33085	3.6	1.1	206	90	0	0	5	0	0	3
<i>Aspergillus sydowii</i> 16844	5098	3.5	1.4	515	90	0	0	5	0	0	3
<i>Aspergillus tubingensis</i> 10550	28801	2.5	1.5	172	41	7	7	25	0	5	12
<i>Botrytis</i> spp.	7024	8	3.7	294	21	6	0	55	0	0	16
<i>Cheatomium</i> spp.	19264	4.9	2.1	258	77	0	0	16	0	1	3
<i>Cladosporium</i> spp.	39294	3.3	1	710	80	0	0	16	0	0	2
<i>Cladosporium cladosporioides</i> (28 days) 16022	26474	3	0.9	570	72	0	0	24	0	0	3
<i>Cladosporium cladosporioides</i> (180 days) 16022	34085	3.3	1.2	702	70	0	0	23	0	0	4
<i>Cladosporium herbarum</i> 58927	2962	3.2	1.4	651	80	0	0	12	0	0	6
<i>Fusarium</i> spp.	1188	3.3	1.5	638	81	0	0	12	0	0	5
<i>Paecilomyces variotii</i> 18502	11188	4.2	1.5	772	0	8	0	74	0	3	13
<i>Penicillium chrysogenum</i> (28 days) 10106	14577	3.5	1.1	867	92	0	0	4	0	0	2
<i>Penicillium chrysogenum</i> (180 days) 10106	33452	3.2	1	733	73	0	0	23	0	0	2
<i>Penicillium canescens</i> 10419	7840	3.1	1.3	659	90	0	0	5	0	0	4
<i>Penicillium citreonigrum</i> 22350	8470	2.8	1	596	68	0	0	27	0	0	4
<i>Penicillium commune</i> 10428	9255	3.3	1.3	762	79	0	0	3	2	0	14
<i>Penicillium corylophilum</i> 9784	12654	2.6	0.9	687	90	0	0	6	0	0	3
<i>Penicillium cecumbens</i> (sp. proposed)	13676	2.4	0.5	420	89	0	0	8	0	0	2
(50) <i>Penicillium thomii</i> 18313	5739	3.1	0.9	721	91	0	0	4	0	0	3
<i>Phoma herbarum</i> (28 days) 44739	5518	2.9	1.2	607	91	0	0	5	0	0	2
<i>Phoma herbarum</i> (180 days) 44739	2648	3.5	1.9	688	79	0	0	14	0	0	5
<i>Stachybotrys</i> spp.	12697	2.8	1.2	137	65	7	2	14	1	1	6
<i>Syncephalastrum racemosum</i> 14832	6753	2.8	2.3	618	75	0	0	12	0	0	10
<i>Tritirachium</i> spp.	5595	2.2	0.9	227	95	0	0	3	0	0	0
<i>Ulocladium</i> sp.	4513	3.8	1.7	767	80	0	0	15	0	0	4
Pollens											
<i>Acer saccharum</i>	800	3.3	4.9	753	6	1	3	1	1	17	68
<i>Abus rubra</i> and <i>rugosa</i> mix	389	4.7	3.1	387	6	4	11	1	0	28	46
<i>Ambrosia trifida</i>	1219	3.2	4.7	876	11	2	2	6	1	12	62
<i>Artemisia tridentata</i>	1703	2.5	4.1	666	6	2	2	3	0	19	66
<i>Betula lenta</i> , <i>nigra</i> & <i>populifolia</i>	2208	2.5	4.2	990	4	1	3	2	0	54	33
<i>Carya laciniata</i>	1942	1.3	2.1	1043	5	0	5	2	0	53	33
<i>Eucalyptus</i>	1095	6.1	5	481	2	1	13	0	0	47	34
<i>Fern</i> (unknown source)	1571	6.8	5.7	373	1	13	17	1	0	38	28
<i>Fagus americana</i>	1991	1.8	2.4	1155	6	2	6	3	0	60	20
<i>Juglans nigra</i>	1968	1.2	2.2	1197	3	1	4	1	0	62	26
<i>Juniperus ashei</i>	2536	3	5.2	766	0	1	1	0	0	16	79
<i>Morus rubra</i>	4232	2.8	5.5	1229	0	0	0	0	0	1	96
<i>Phleum pratense</i>	3390	1.5	3.4	936	2	0	1	0	0	11	82

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