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**Origin, distributions, and environmental significance of ubiquitous humic-like fluorophores in**

**Antarctic lakes and streams**

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**Morimaru Kida<sup>1†‡\*</sup>, Taichi Kojima<sup>1</sup>, Yukiko Tanabe<sup>2, 3</sup>, Kentaro Hayashi<sup>4</sup>, Sakae Kudoh<sup>2, 3</sup>,**

**Nagamitsu Maie<sup>5</sup>, Nobuhide Fujitake<sup>1</sup>**

<sup>1</sup> Graduate School of Agricultural Science, Kobe University, 1-1 Rokkodai, Nada, Kobe, Hyogo  
657-8501, Japan

<sup>2</sup> National Institute of Polar Research, Research Organization of Information and Systems, 10-3  
Midori-cho, Tachikawa, Tokyo 190-8518, Japan

<sup>3</sup> Department of Polar Science, SOKENDAI (The Graduate University for Advanced Studies), 10-3  
Midori-cho, Tachikawa, Tokyo 190-8518, Japan

<sup>4</sup> Institute for Agro-Environmental Sciences, NARO, 3-1-3 Kannondai, Tsukuba, Ibaraki 305-8604,  
Japan

<sup>5</sup> School of Veterinary Medicine, Kitasato University, Towada, Aomori 034-8628, Japan

† Research Fellow of Japan Society for the Promotion of Science

19 ‡ Present address: Research Group for Marine Geochemistry (ICBM-MPI Bridging Group), Carl  
20 von Ossietzky University of Oldenburg, Institute for Chemistry and Biology of the Marine  
21 Environment (ICBM), Carl-von-Ossietzky-Str. 9-11, 26129 Oldenburg, Germany

22

23 \* corresponding author

24 morimaru.kida@people.kobe-u.ac.jp

25 ORCID ID: 0000-0002-9908-2012 (M.K.), 0000-0002-2936-9544 (K.H.)

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## 27    **Abstract**

28    This study characterized dissolved organic matter (DOM) obtained from 47 lakes and 2 streams on  
29    ice-free areas at Lützow-Holm Bay and Amundsen Bay in East Antarctica ( $n = 74$ ), where few  
30    biogeochemical studies have been historically conducted. Samples were analyzed for basic water  
31    chemistry and by resin fractionation, UV-vis spectroscopy, and excitation emission matrix  
32    spectroscopy combined with parallel factor analysis (EEM-PARAFAC). Salinity of the samples  
33    ranged very broadly from fresh to hypersaline as a result of evaporative concentration. There was a  
34    clear positive correlation between log-salinity and the spectral slopes of DOM ( $S_{275-295}$ ), an indicator  
35    of photodegradation. Thus, we interpreted the correlation as a progression of photodegradation by  
36    prolonged water retention time. Of the identified seven PARAFAC components, three ubiquitous  
37    humic-like components decreased as photodegradation progressed, while a photorefractory UVC  
38    humic-like component increased its relative abundance. A non-humic component, traditionally  
39    defined as Peak N, did not show a trend depending on photodegradation, and its level was high in  
40    nutrient-rich lakes, presumably due to high in-situ production. We found robust correlations between  
41    the relative abundance of the ubiquitous humic-like components and that of the Peak N component  
42    in the bulk DOM irrespective of water types or ice-free areas. We proposed there were common  
43    processes that generated the ubiquitous humic-like components from the Peak N component in the  
44    Lützow-Holm Bay and Amundsen Bay lakes and streams, such as bacterial processing of primary

45 production-derived DOM and photochemical transformation of microbial DOM.

46

47 **Keywords**

48 autochthonous; DOM; fluorescence; microbial process; PARAFAC; photodegradation

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## 1. Introduction

Dissolved organic matter (DOM) is a highly heterogeneous collection of hundreds of thousands of organic compounds that collectively play an essential role in key ecosystem processes (Zark et al., 2017). DOM ubiquitously exists in water and comprises a similar mass of carbon as atmospheric CO<sub>2</sub> or the global biomass (Hedges et al., 1997). DOM is involved in various biogeochemical processes, such as light attenuation, metal complexation, pH buffering, and influencing microbial metabolism (Findlay and Sinsabaugh, 2003), and is one of the most important components on Earth that influences the health of ecosystems.

The chemical composition and functionality of DOM is strongly influenced by the original source and ecosystem activities (Roth et al., 2014). DOM in streams and peatland rivers often contains a high proportion of terrestrial lignin-derived materials, while that in large lakes and oceans usually contains a high proportion of microbial-derived matter. Aside from the ecosystem-specificity of DOM, molecularly and photochemically indistinguishable ubiquitous components of DOM have been recently identified in samples from diverse environments (Murphy et al., 2018; Zark and Dittmar, 2018). These findings imply that there are some common mechanisms that produce these ubiquitous components in DOM, such as either biotic or abiotic pathways that lead to the production of ubiquitous, recalcitrant components of DOM that persist in the environment. Unraveling the underlying mechanisms of the generation of these components is one of the key questions in

understanding transformation and long-term turnover of DOM (Murphy et al., 2018; Zark and Dittmar, 2018).

In ice-free areas of continental Antarctica, where no vascular plants are present, DOM is originated dominantly from in-situ biological activity (both benthic and planktonic phototrophs) in water and phototrophs (mosses, algae, lichens, cyanobacteria) in the catchments (Matsumoto, 1989; McKnight et al., 1994, 1991). Furthermore, because lakes are isolated from each other with independent water catchment areas by impermeable bedrocks, they function as closed hydrologic systems. The simplicity of the organic matter sources and the lack of the connection with other hydrologic systems limit the number of environmental factors that would otherwise make it difficult to elucidate processes responsible for DOM transformations. Due to the simplicity of the sources, the chemical characteristics of Antarctic lake DOM have been considered to vary little among Antarctic lakes (Aiken et al., 1996; McKnight et al., 1991). However, past studies were exclusively conducted in the McMurdo Dry Valleys of East Antarctica (Matsumoto, 1989; McKnight et al., 1994, 1991), and very few biogeochemical studies on DOM chemistry have been conducted in other regions in Antarctica (Farzadnia et al., 2017).

Here, we report an extensive study conducted in 47 lakes and 2 streams in 6 ice-free areas at Lützow-Holm Bay (East Dronning Maud Land) and Amundsen Bay (Enderby Land) in East Antarctica. This was by far the largest and most diverse dataset that has been historically obtained

from Antarctic lakes. The overall goal of this study was to use spectroscopic techniques to evaluate the potential sources and sinks of DOM in these lakes and streams. Spectroscopic techniques that were used in this study, such as UV-vis spectroscopy and excitation-emission matrix spectroscopy (EEM), have been successful in studying concentrations, compositions, and dynamics of chromophoric (CDOM) and fluorescent DOM (FDOM) in aquatic systems including Antarctic lakes (Cawley et al., 2016, 2013; McKnight et al., 1994). We hypothesized that DOM in the Antarctic lakes and streams has much larger compositional diversity than has been considered and such diversity is mostly introduced by external factors such as photodegradation because the simplicity of the sources would reduce the original DOM heterogeneity. Studying the geochemistry of diverse microbially-dominated lake systems should lead to a better understanding of lower latitude aquatic systems as well, where several end-members form a continuum of DOM characteristics.

## **2. Materials and Methods**

### **2.1. Study area**

The main sampling campaign was conducted during the austral summer during December 2016 and February 2017 in ice-free areas along Sôya Coast (Lützow-Holm Bay, East Dronning Maud Land), located in the southern area of Syowa Station (Japan's base) (Fig. S1). In Sôya Coast, there are several ice-free areas, including Skarvsnes, Langhovde, Skallen, Breidvågnipa, and Rundvågshetta,



where hundreds of lakes of various sizes and water chemistry exist (Imura et al., 2003; Kimura et al., 2010; Kudoh and Tanabe, 2014). These ice-free areas were created during the last 7,000 years under the glacier retreat and iso-static uplifts that occurred after the Last Glacial Maximum (Kudoh and Tanabe, 2014). Almost no lakes have any in- or out flows, and they are fed by melt water from adjacent glaciers and/or snow and ice in their catchments. The lakes have been isolated from each other with independent water catchment areas, and they function as closed hydrologic systems. A 1–2 m thick ice cover usually develops on all lakes during March–December (or January) (Imura et al., 2003). Additional water sampling was conducted in an ice-free area at the foot of Mt. Riiser-Larsen, Amundsen Bay, West Enderby Land. A large number of Adélie penguins (*Pygoscelis adeliae*) (> 1800 breeding pairs) and emperor penguins (*Aptenodytes forsteri*) have been reported in this region (Takahashi et al., 2000), which would influence the lake water chemistry and biology.

Water chemistry of the lakes in Sôya Coast is very diverse despite sharing the same climate, suggesting that it would induce the diversity in lake water DOM as well. For example, salinity ranges from completely fresh to saline to hypersaline, i.e., more than seven times the salinity of seawater (Matsumoto, 1989). Although most of the saline lakes are considered to have been generated via the evaporative concentration of lake water under dry conditions, others are marine relict lakes, which were in the shallow coastal sea and formed after the iso-static uplift of the present lake shores. The evapo-concentration of Sôya Coast lakes occurs at variable rates reflecting the

balance between evaporation and freshwater inputs. When freshwater inputs dominate, even marine relict lakes can have zero salinity after a long period (see Table 1). In contrast to salinity, concentrations of essential nutrients for phototroph growth, such as nitrate and phosphate, are quite low in most of the lakes, rendering them oligotrophic. Waters in proglacial lakes (lakes located in front of or close to a glacier) exhibit ultra-oligotrophic freshwater conditions. They typically contain inorganic particles derived from glacial deposits; thus, the water transparency is low despite the ultra-oligotrophic state. Proglacial lakes directly attached to the glacier terminal tend to have permanent ice covers, which can develop up to 4-m thick in winter. Geological, limnological, and ecological characteristics of many of the sampled lakes have been described and reviewed elsewhere (Imura et al., 2003; Kimura et al., 2010; Kudoh and Tanabe, 2014)

## **2.2. Water sampling and handling**

Water samples were collected at the deepest point ( $n = 19$ ) or shore ( $n = 28$ ) of 47 lakes in the ice-free areas at Lützow-Holm Bay and Amundsen Bay (Fig. S1). Two streams (Yukidori and Yatsude Valleys in Langhovde,  $n = 16$ ) were additionally sampled for comparison to lakes. The sampling locations, limnological characteristics, and basic optical and chemical properties of DOM for each lake are presented in Table 1, while those for streams are presented in Table S1. Stream samples were collected at approximately regular intervals from the beginning (glacial melt) to the

end (before the outlet to the sea) of each stream. At the streams and lake shores, samples were directly collected into 550-mL volume polyethylene terephthalate bottles after rinsing more than three times with the collected water. The pH, salinity (*S*), and water temperature were recorded in situ using portable water quality meters (LAQUA series, Horiba, Kyoto, Japan). At the deepest point of the lakes (checked using a portable sonar depth meter, PS-7, Hondex, Aichi, Japan), after the confirmation of uniformity in water properties by a multi-water quality logger (YSI-6600V2, YSI Inc., OH, USA) (Fig. S2), lake water was collected at mid-depth of the water column using a Teflon cylindrical water sampler into the bottles. When water stratification was observed, water was collected at several depths to capture changes in water chemistry (Figs. S2 i–l). At five lakes (Figs. S2 b–d, p, and q), porewater was sampled from sediment cores that were collected using a gravity corer with a transparent acrylic pipe (diameter 5 cm, length 50 cm) (Fig. S1e). After gently aspirating water above the sediment cores (bottom water) using a glass pipet, porewater was collected and stored in the dark.

The collected waters were filtered in field laboratories as soon as possible, usually within a few hours after sampling. Fourteen samples collected in Skallen, Breidvågnipa, and Rundvågshetta were filtered in the field laboratory in Skarvsnes 1–3 days after sampling because of a logistical limitation. Samples collected in Amundsen Bay were filtered in a laboratory on the ice breaking ship “Shirase” on the day of sampling. Samples for dissolved organic carbon (DOC), total dissolved

nitrogen (TDN), and optical analysis were filtered through pre-combusted (450°C, > 3 h) glass fiber filters (nominal pore size 0.3 µm, GF-75, Advantec, Tokyo, Japan) into pre-combusted glass bottles. A 50-mL aliquot of GF-75 filtered samples was stored in a plastic bottle and kept frozen (−20°C) for inorganic nutrient analysis. The samples for DOC and TDN analysis were spiked with sodium azide (final concentration of 0.02%) and hydrochloric acid (final concentration of 1%), respectively (Wako Pure Chemical, Osaka, Japan), and stored for further analysis in Japan. Optical analysis was performed immediately after filtration in the field laboratory in Skarvsnes or on Shirase.

### **2.3. Water analysis**

UV-vis absorption spectra (240–600 nm) were measured using a UV-visible spectrophotometer (GENESYS 10S UV-Vis, Thermo Fisher Scientific, Tokyo, Japan) with a 5-cm path length, acid-cleaned, quartz cuvette at ambient temperature. Some samples with a high optical density were measured with a 1-cm path length cuvette. Measurements were baseline-corrected using ultrapure water (Wako Pure Chemical, Osaka, Japan). Overestimation of absorbance by scatter was evident in some samples, which was most likely due to clay-sized inorganic particles that passed through the GF-75 filter. Such samples were mostly from proglacial lakes, where lake water was turbid from silt and clay-sized particles fed by glaciers (Table 1). Absorbance data in this paper are expressed as Napierian absorption coefficients ( $a_\lambda$ , m<sup>−1</sup>). Absorption coefficients of DOM were calculated from

177  $a_{\lambda} = 2.303A_{\lambda}/l$

178 where  $A_{\lambda}$  is the absorbance at wavelength  $\lambda$  and  $l$  is the cell path length in meters (Helms et al.,  
179 2008).

180 Fluorescence EEMs were collected with a spectrofluorometer (FluoroMax4, Horiba,  
181 Kyoto, Japan) equipped with a 150 W xenon arc lamp at room temperature using 5 nm excitation  
182 and emission slit widths, integration time of 0.25 s, excitation (Ex) range of 240–450 nm at 5 nm  
183 increments, and an emission (Em) range of 300–550 nm at 2 nm increments. A sharp peak of the  
184 xenon lamp emission (467 nm) and Raman peak (397 nm) were checked daily and corrected if they  
185 deviated  $> 0.5$  nm from the desired values. The EEMs were corrected for all necessary steps to  
186 remove biases and normalized in Raman units (RU) (Murphy et al., 2018). Blank EEM (ultrapure  
187 water) used for blank subtraction and Raman normalization was measured daily.

188 In the laboratory in Japan, we measured DOC and TDN using a total organic carbon  
189 analyzer combined with a total nitrogen measuring unit (TOC-L<sub>CPH</sub>, Shimadzu, Kyoto, Japan) as in  
190 Kida et al. (2019). Also, DOM in the samples was fractionated into hydrophobic (HPO) and  
191 hydrophilic DOM based on adsorption onto Supelite™ DAX-8 resin in quadruplicate according to  
192 Kida et al. (2019). DOM fractions retained on the DAX-8 resin (HPO) included hydrophobic acids  
193 (i.e., humic substances) and neutrals, which are too hydrophobic to be eluted by dilute alkali solution,  
194 such as less oxygenated humic substances, hydrocarbons, carbonyl compounds, and some

anthropogenic matter. The proportion of HPO in DOM (%HPO) was calculated by  $[\text{HPO}]/[\text{DOM}] \times 100$  on a carbon basis. The coefficient of variation of replicate %HPO determination measurements was  $< 17\%$  (average  $\pm$  one standard deviation SD,  $4.9\% \pm 4.4\%$ ; Fig. S3b).

Dissolved inorganic nutrients ( $\text{PO}_4^{3-}$ ,  $\text{NH}_4^+$ ,  $\text{NO}_2^-$ ,  $\text{NO}_3^-$ , and  $\text{SiO}_3^{2-}$ ) were determined colorimetrically using an auto analyzer as previously reported (Tanabe et al., 2017). The limits of detection and quantitation were determined as the average plus 3 or 10 SD of the  $10 \times$  blank measurements, respectively.  $\text{NO}_3^-$  concentrations were determined as the difference between ( $\text{NO}_2^- + \text{NO}_3^-$ ) concentrations minus  $\text{NO}_2^-$  concentrations with or without a  $\text{NO}_3^-$  reduction step. A dissolved organic nitrogen (DON) concentration was calculated as the difference between TDN and the sum of dissolved inorganic nitrogen concentrations. If any of the dissolved inorganic nitrogen concentrations were below the detection limit, a DON concentration was calculated as the difference between TDN and the sum of the rest of the detected dissolved inorganic nitrogen concentrations. The proportion of DON in TDN (%DON) was calculated by  $[\text{DON}]/[\text{TDN}] \times 100$  on a nitrogen basis.

#### 2.4. Optical index and parallel factor analysis

We calculated optical indices, such as the spectral slope ( $S_{275-295}$ ), DOC-specific ultraviolet absorbance ( $\text{SUVA}_{254}$ ), and commonly used fluorescence-based indices (Gabor et al., 2014; Helms et

al., 2008; Weishaar et al., 2003), from the absorbance and EEM data to estimate the structural properties and origin of DOM (See Supporting Information, SI 1, Figs S3–S5). Other spectral slope indices that use longer wavelengths were not used because many samples were optically too dilute for the accurate calculation. In addition, we analyzed the EEMs by parallel factor analysis (PARAFAC) to mathematically deconvolute them into distinct fluorescent components with independent spectral characteristics (SI 2) (Murphy et al., 2018). The identified seven fluorescent components were compared with the published data on an open-access spectral database (OpenFluor; Murphy et al., 2014), with the Tucker congruence coefficient (TCC) as similarity criteria. The components were named according to their rounded fluorescence emission maximum:  $C_{325}$ ,  $C_{415}$ ,  $C_{360}$ ,  $C_{500}$ ,  $C_{440}$ ,  $C_{460}$ , and  $C_{300}$ , with variation explained by each component decreasing sequentially from  $C_{325}$  to  $C_{300}$ .  $C_{300}$  was named so because it had a fluorescence maximum at  $Ex/Em = 270/302$  nm when  $Em < 318$  nm was not removed before modeling (data not shown). The fluorescence intensity at the maximum for each component was represented by  $F_{max}$  (in RU) (Murphy et al., 2014). The  $F_{max}$  of individual components divided by its DOC ( $F_{max}/DOC$ ) was calculated to track changes in its overall importance in the bulk DOM.

## 2.5. Statistical Analysis

The statistical difference between multiple parameters was tested by Tukey-Kramer's multiple

comparisons of means. Linear regressions were fitted using the ordinary least squares method with log<sub>10</sub> transformation of variables when needed. Correlations among F<sub>max</sub>/DOC values of the identified PARAFAC components were evaluated using Spearman's rank correlation coefficient ( $\rho$ ). A significance level of 0.05 was selected for multiple comparisons to test the null hypothesis. Linear correlations were considered statistically significant when  $P < 0.01$ . All analyses were conducted with Origin 2016 (OriginLab Corporation, MA, USA).

### 3. Results

#### 3.1. Water chemistry and DOM composition

Salinity of the samples ranged very broadly from fresh to hypersaline ( $S = 179$ , 5 times that of seawater; Table 1). The pH was neutral to slightly basic, and dissolved silica ( $\text{SiO}_3^{2-}$ ) concentrations were high (Table S2). Dissolved inorganic phosphorus and nitrogen concentrations were generally very low and undetectable or below the quantification limit (Table S2). As a result, the %DON in lakes was as high as  $93.3\% \pm 13.6\%$ . Among the measured nutrients, only  $\text{SiO}_3^{2-}$  concentrations increased along the salinity gradient of  $S = 0\text{--}0.5$ , but the trend in  $\text{SiO}_3^{2-}$  concentrations became unclear in higher salinity lakes. Other nutrient concentrations did not show any trend along salinity.

The distribution patterns of C, N, and the absorption-based optical indices along the salinity gradient are shown in Fig. 1. The DOC concentration ranged  $0.25\text{--}146 \text{ mg C L}^{-1}$  and



increased as a function of salinity. A similar trend was observed for DON and  $a_{254}$  (quantitative measures of CDOM). Salinity alone explained 70%, 61%, and 53% of the variability of DOC, DON, and  $a_{254}$  values, respectively (linear regression of log-log plots). Consequently, a strong log-log linear relationship was observed between DOC and  $a_{254}$  ( $R = 0.94$ ,  $P < 0.01$ ) (Table 1). The low DOC/DON ratios indicate the microbial origin of DOM. The %HPO ranged from 18.3% to 60.5% ( $40.4\% \pm 8.5\%$ ). The %HPO tended to be lower in proglacial lakes and higher in the porewater of lakes.

The UV-absorption indices ( $SUVA_{254}$  and  $S_{275-295}$ ) showed typical non-terrestrial properties (Fig. 1, Table 2). The  $SUVA_{254}$  values were low ( $0.22-1.66 \text{ L mg C}^{-1} \text{ m}^{-1}$ ) and did not show a clear trend along salinity (Fig. 1e). Amundsen Lake 3, which receives large N inputs from the catchment, had the highest  $SUVA_{254}$  value among others (Fig. 1e). This lake had a very high abundance of phytoplankton, which was obvious from its green color (Fig. S1f). We note that the influence of  $\text{NO}_3^-$  to the  $SUVA_{254}$  value of this lake was negligible due to the low absorbance of  $\text{NO}_3^-$  ( $A_{254} < 0.01$  at  $100 \text{ mg/L NO}_3^-$ , Weishaar et al., 2003). In contrast, the values of  $S_{275-295}$  showed a clear increasing tendency with the increase in salinity (log scale) ( $R = 0.80$ ,  $P < 0.01$ ; Fig. 1f). Porewater and stream samples had lower  $S_{275-295}$  values than others at a given salinity (Fig. 1f, Table 2). The fluorescent indices also showed typical non-terrestrial properties (Table 2), but values varied along salinity in spite of the anticipated predominantly microbial origin of DOM (Fig. S6).

### 3.2. Assignments of PARAFAC components

Seven PARAFAC components were identified with our dataset (Fig. 2). For all of the components except for C<sub>300</sub>, multiple strong matches ( $TCC_{ex \times em} > 0.95$ ) were observed (18, 23, 10, 40, 3, and 10 matches for C<sub>325</sub>, C<sub>415</sub>, C<sub>360</sub>, C<sub>500</sub>, C<sub>440</sub>, and C<sub>460</sub>, respectively) on OpenFluor. C<sub>300</sub> had seven matching components with  $TCC_{ex \times em} > 0.9$ . Assignments of the components are summarized in Table S3 with descriptions of each component provided in the past literature. Other than C<sub>300</sub>, C<sub>325</sub>, and C<sub>500</sub>, the identified components have been reported as either having allochthonous or autochthonous origins (Table S3).

We assigned C<sub>300</sub> and C<sub>325</sub> as protein-like components (Table S3) based on their fluorescence positions ( $Em < 350$  nm) and the scarcity of polyphenols in Antarctic lakes. C<sub>360</sub> had fluorescence peaks at  $Em$  362 nm with the primary excitation maximum at 290 nm. The position of C<sub>360</sub> was clearly different from a microbial humic-like component (peak M) that occurs at  $Em$  370–430 and  $Ex$  290–325, which is often found in aquatic systems (Coble et al., 1998). Therefore, we assigned this component as a non-humic peak N ( $Em$  360–370 nm,  $Ex$  280 nm), which has been associated with new DOM production in water (Coble et al., 1998).

C<sub>460</sub> was assigned as a UVC humic-like component (Table S3), which was reported as photorefractory because of the rapid attenuation of UVB light in water (Stedmon and Markager,

2005). The rest of the humic-like components ( $C_{415}$ ,  $C_{500}$ , and  $C_{440}$ ) have been reported as ubiquitous components, which are found in almost all environments and derived from both allochthonous and autochthonous sources (Table S3) (Murphy et al., 2018; Stedmon and Markager, 2005). We further compared our model to the “ubiquity” models of Murphy et al. (2018), which were modeled by the one-sample PARAFAC of a set of photo-decomposed DOM isolates (Fig. S7). Quite good matches were found between  $C_{415}$ ,  $C_{500}$ , and  $C_{440}$  with several ubiquitous components in their models, supporting our assertion that these components were ubiquitous humic-like components.

### 3.3. Distribution of PARAFAC components

Fluorescence intensity of the PARAFAC components ( $F_{max}$ ) less clearly increased along the salinity gradient than the other quantitative measures (Fig. S8a), and salinity explained less than 42% of the variability of  $F_{max}$  (linear regression of log-log plots). The explained variability was especially small for the ubiquitous humic-like components (5.7%–8.2%). When normalized by DOC,  $C_{325}$  had the highest intensities followed by  $C_{415}$ , while the other components had lower intensities and they were not significantly different from each other (Fig. S9). It was evident that  $F_{max}/DOC$  values of the PARAFAC components differed among water types (Fig. S8b). Porewater samples typically had a 5–20 times higher average  $F_{max}/DOC$  values than lake surface water samples, except for  $C_{460}$  and  $C_{300}$  (Table 2). Stream samples also had 2–9 times higher average  $F_{max}/DOC$  values than lake

samples for the ubiquitous humic-like components ( $C_{415}$ ,  $C_{500}$ , and  $C_{440}$ ) (Table 2). Saline lake samples had the lowest average  $F_{max}/DOC$  values for all the components (Table 2).

Figure 3 was informative regarding the distribution of the PARAFAC components in response to photodegradation, which was assessed by the spectral slope  $S_{275-295}$  (Fichot and Benner, 2012; Helms et al., 2008): 1)  $C_{360}$ , which is derived from recent in-situ production (Coble et al., 1998), did not show changes in its relative abundance depending on the degree of photodegradation (Fig. 3b); 2) photodegradation decreased the ubiquitous humic-like components ( $C_{415}$ ,  $C_{500}$ , and  $C_{440}$ ), and these components dominated more in the porewater and streams and less at the epilimnion in saline lakes (Fig. 3c); 3) UVC humic-like component  $C_{460}$ , which is photorefractory (Stedmon and Markager, 2005), increased its relative importance as photodegradation of bulk DOM progressed (Fig. 3d); 4) Saline lakes exhibited unique PARAFAC compositions with more contribution from protein-like components ( $C_{325}$  and  $C_{300}$ ) and much less contribution from  $C_{460}$ , which deviated from the relationship between  $\%C_{460}$  and  $S_{275-295}$  (Figs. 3a and 3d); and 5) Amundsen Lake 3 (“penguin” lake), which received substantial nutrient inputs from nearby penguin colonies (Table S2), had the lowest protein-like components and highest ubiquitous humic-like components as well as high  $\%C_{360}$ .

#### 4. Discussion

#### 4.1. Factors controlling DOM and nutrient concentrations

It was obvious that the bulk DOM concentration was significantly influenced by evapo-concentration in the Lützow-Holm Bay and Amundsen Bay lakes because of the dry climate (Figs. 1a and 1b). This was as expected because evapo-concentration or dilution by melt water inputs results in either the increase or decrease of any solute in water in closed hydrologic systems. Similar concentration effects have been observed in lower latitude arid regions. Among such studies, Osburn et al. (2011) compared DOC-conductivity (in  $\text{mS cm}^{-1}$ ) relationships among saline lakes reported in past studies and found that the slope of regressions between log DOC and log conductivity varied from 0.322 to 0.796 ( $0.577 \pm 0.156$ ,  $n = 6$ ). Our Antarctic lake samples had an equation of  $\log \text{DOC} = 0.345 + 0.394 \times \log \text{conductivity}$  ( $R^2 = 0.68$ ,  $P < 0.001$ ). The shallower slope of the regression compared to those reported in lower latitude arid regions suggested that the net DOM accumulation in the studied Antarctic lakes was lower, possibly due to the lack of nutrients necessary for primary production in lake water (Table S2) and the harsh Antarctic environment.

In contrast, nutrients other than dissolved silica were not concentrated by evapo-concentration (Table S2). Consequently, there were no correlations between the  $F_{\text{max}}$  values of the PARAFAC components and nutrient concentrations, as opposed to those observed in arid lakes (Osburn et al., 2011). This implies that the Antarctic lakes are substantially N- and P-deficient compared to the arid lakes, rendering them oligotrophic (Kudoh and Tanabe, 2014). As a

result, %DON was more influenced by nutrient inputs from the surrounding areas than by evapo-concentration. For example, two lakes in Amundsen Bay (Amundsen Lake 1 and 3) had low %DON and had high nitrate ( $\text{NO}_3^-$ ) concentrations because of N inputs from surrounding penguin colonies and south polar skua nests (Table S2). Similarly, five lakes (lakes other than Nurume) and the streams located in Yukidori and Yatsude Valleys in Langhovde are affected by inputs from abundant snow petrels that nest in the valleys and had low %DON (Table S2). These results suggested that the animal-related point source is important in influencing the nutrient distribution in Antarctic lakes.

#### **4.2. Factors controlling the DOM composition**

It appeared that both in-situ production and photodegradation substantially influenced the composition of DOM in the studied Antarctic lakes. The observed increase in  $S_{275-295}$  values with increasing salinity (Fig. 1f) suggests the progression of photodegradation of DOM as a result of prolonged water retention time (Fichot and Benner, 2012; Helms et al., 2008). Most of  $S_{275-295}$  values were higher than those previously reported in coastal waters where terrestrial inputs are low (for example,  $0.024 \text{ nm}^{-1}$  in nearshore water of Georgia Bright; Helms et al., 2008), which supports the predominantly microbial (autochthonous) origin and/or the highly photodegraded state of the studied DOM. We note that the effects of salinity and pH on the optical properties were investigated

and found to be negligible (Table S4); thus, the compositional change was the primary driver for variability in CDOM and FDOM properties. The negligible effect of water chemistry on  $S_{275-295}$  was particularly important because we used the index as the only indicator for DOM photodegradation. Fichot and Benner (2012) has also shown that neither biological processes nor changes in pH, ionic strength,  $\text{NO}_3^-$ , or  $\text{NO}_2^-$  interferes with  $S_{275-295}$  determination.

The distribution patterns of the PARAFAC components along  $S_{275-295}$  (Fig. 3) could largely be interpreted by the following three biogeochemical processes: **i)** in-situ production, **ii)** photodegradation, and **iii)** precipitation due to high salinity. The four samples that had the top four highest % $C_{360}$  (Fig. 3b) were those having the highest nutrient (especially  $\text{NO}_3^-$ ,  $\text{NH}_4^+$ , and  $\text{PO}_4^{3-}$ ) concentrations among others (Table S2), and water color of these lakes were yellow-green to green (Fig. S1f), strongly supporting that  $C_{360}$  was derived from recent in-situ production. It is worth noting that there were stronger log-log correlations between DON concentrations with the levels of  $C_{360}$  ( $R = 0.73$ ) and the protein-like components ( $R = 0.79\text{--}0.81$ ) than with those of the ubiquitous humic-like components ( $R = 0.45\text{--}0.51$ ). Thus, we propose that the level of  $C_{360}$  in the studied Antarctic lakes was primarily driven by in-situ production, and not by photodegradation. It is currently not established whether  $C_{360}$  (Peak N) is directly produced by phytoplankton or requires bacterial processing of organic matter originated from primary production. However, the occurrence of Peak N in axenic diatom cultures does indicate that some phytoplankton can produce Peak N

directly (Castillo et al., 2010).

Photodegradation substantially influenced the relative contribution of the ubiquitous humic-like components and  $C_{460}$  (Figs. 3c and 3d). This was in accordance with the photolabile and photorefractory nature of the ubiquitous humic-like components and  $C_{460}$ , respectively (Murphy et al., 2018; Stedmon and Markager, 2005). A decrease in %C(ubiquitous humic) (and their  $F_{max}/DOC$ ) at the epilimnion of saline lakes further supported the photodegradation of the ubiquitous humic-like components (Fig. 3c). In shallow freshwater Sôya Coast lakes ( $\leq 3$  m), Tanabe et al. (2010) observed that 45%–60% of photosynthetically active radiation (PAR, 400–700 nm) and 20%–60% of ultraviolet radiation (300–400 nm) reached the lake beds, implying that photodegradation of the ubiquitous humic-like components could occur throughout the water column. It was as expected that porewater samples had higher contribution from the photolabile ubiquitous humic-like components because of the lack of sunlight penetration into the lake sediment (less than 1 cm) (Fig. 5 of Tanabe et al. 2010). Interestingly, stream samples had similar optical characteristics as porewater samples, namely, high  $SUVA_{254}$  and %C(ubiquitous humic) and low  $S_{275-295}$  and % $C_{460}$  (Figs. 1 and 3, Table 2). We presume that this was because the water residence time of the streams (at most a few days) was much shorter than the most of lakes (except for lakes directly fed by a glacier) and, thus, stream DOM was not photodegraded substantially. The continuous DOM inputs from decomposed, partially humified mosses within the stream and in stream banks (Kanda et al.,



2004) could also be a source of the ubiquitous humic-like components. The low %C<sub>460</sub> in saline lakes (Fig. 3d) and concomitant increases in %C(protein) (Fig. 3a) could be caused by precipitation of C<sub>460</sub> due to high salinity (Sholkovitz, 1976). Another possibility may be the greater primary production of protein-like DOM in saline lakes (Osburn et al., 2011). However, it appeared that this was not true, because Lake Funazoko, which had the much higher nutrient concentrations than other saline lakes (Table S2), had lower C(protein)/DOC and thus lower %C(protein). Also, the phototroph abundance is typically small in the studied saline lakes (Tanabe, unpublished data).

The lack of correlations between the protein-like component levels and nutrient concentrations may be due to rapid microbial processing of the protein-like components originated from primary production (Stedmon and Markager, 2005). It could also be partly due to the timing of sampling. In Lakes Oyako and Hotoke in Sôya Coast, Tanabe et al. (2008) observed the maximum chlorophyll a fluorescence under dim-light conditions during austral spring and autumn and the minimum chlorophyll a fluorescence in ice-free summer, and they concluded that the too strong summer light acts as an inhibitor of Antarctic phytoplankton growth. Since our samples were collected during austral summer, they may have already undergone extensive algae breakdown and photolytic/microbial modification of primary production-derived DOM. Insight as to intra-annual variation of DOM concentrations and compositions should be gained when samples collected during the winter time are analyzed. Finally, the fluorescence-based indices failed to adequately capture the

changes in FDOM compositions described above (Fig. S10).

### **4.3. Generation of ubiquitous humic-like components and its ecological implications**

The evapo-concentration of the studied lakes occurs at variable rates reflecting the balance between evaporation and freshwater inputs. As such, the correlation analysis of the fluorescence intensities of the PARAFAC components would result in false-positive if not corrected for concentration effects. Therefore, we compared the levels of the PARAFAC components after dividing by the DOC concentrations ( $F_{\max}/\text{DOC}$ ), as the DOC concentrations in this study were largely (70%) explained by salinity and dividing by DOC could effectively reduce concentration effects. In addition,  $F_{\max}/\text{DOC}$  represents the relative abundance of the PARAFAC components in the bulk DOM. Thus, if  $F_{\max}/\text{DOC}$  values of two components vary and are strongly correlated among samples, it means that the components behave in parallel independent of the behavior of the other constituents of DOM. Figure 4 shows the correlations between  $C_{360}/\text{DOC}$  and the ubiquitous humic-like components  $C_{415}/\text{DOC}$ ,  $C_{440}/\text{DOC}$ , and  $C_{500}/\text{DOC}$ . All the ubiquitous humic-like components had a significant strong correlation with  $C_{360}$ , while the other protein-like components and  $C_{460}$  had a much weaker correlation. The y-intercepts for  $C_{415}$  and  $C_{500}$  were not statistically different from zero ( $P > 0.01$ ). These universal correlations were observed despite the fact that the samples were collected from closed hydrologic systems of distinct water types and different ice-free areas. This strongly

suggested that there were common processes that generated the ubiquitous components (C<sub>415</sub>, C<sub>440</sub>, and C<sub>500</sub>) from C<sub>360</sub> in the Lützow-Holm Bay and Amundsen Bay lakes and streams, with a certain proportion of C<sub>360</sub> being transformed into the ubiquitous components. The inconsistent assignments of these components in literature (allochthonous or autochthonous, Table S3) also imply their multiple origins or common environmental processing that ultimately leads to these fluorescence. Because the lakes have been isolated from each other (~7000 years) with independent water catchment areas and they function as closed hydrologic systems, such a process should work on areas larger than the catchment in size or it should work commonly among lakes.

In water, humic-like fluorophores can be generated in a culture of bacterial communities amended with phytoplankton-derived DOM or simple substrates, such as glucose, in a relatively short period (< a few weeks) (Kramer and Herndl, 2004; Rochelle-Newall and Fisher, 2002; Shimotori et al., 2012). The bacterial processing of C<sub>360</sub> could be a source of the ubiquitous humic-like fluorophores in the studied lakes. Although C<sub>500</sub> has not been linked to autochthonous sources (Table S3), its relatively high contribution to our Antarctic dataset suggested that it was of autochthonous origin in this study, re-emphasizing that one cannot assign humic-like fluorophores as terrestrial just because their spectral features resemble that of terrestrially-derived natural organic matter (Murphy et al., 2018). The fact that the same ubiquitous humic-like fluorophores were found in Lake Fryxell and the deep Sargasso Sea (4534 m) (Fig. S7), where sunlight penetration is

negligible (discussed below), supports the biotic generation of these fluorophores. Although recent studies demonstrated that humic-like fluorophores can also be generated directly from phytoplankton in axenic cultures (Castillo et al., 2010; Chari et al., 2013; Fukuzaki et al., 2014), its contribution to the generation of the ubiquitous humic-like fluorophores was likely low in our study considering the lack of relationships between nutrient concentrations and ubiquitous humic-like fluorophore levels.

Furthermore, abiotic photochemical humification (photohumification) of DOM in the presence of dissolved nitrogen has also been proposed (Kieber et al., 1997). Photooxidation has been shown to cross-link lipid precursors, incorporating N (free amine groups or ammonia) into the newly formed marine humic substance and creating humic-like fluorescence (Kieber et al., 1997). Although McKnight et al. (1991) noted that photohumification would not occur in McMurdo Dry Valleys Antarctic lakes (such as Lake Fryxell), which are permanently ice-covered and thus have low light intensities (photon flux density is 3–10  $\mu\text{mol photons m}^{-2} \text{ s}^{-1}$  below the ice) (McKnight et al., 1991), it could occur in Lützow-Holm Bay and Amundsen Bay lakes because most lakes are completely ice-free during the 2-month austral summer and they can have two orders brighter light ( $> 600 \mu\text{mol photons m}^{-2} \text{ s}^{-1}$  PAR up to 4 m) (Tanabe et al., 2008) than the lakes in McMurdo Dry Valleys. Considering the richness of DON (Fig. 1) and the strong light intensity, photohumification in the presence of DON (and DIN in some nutrient-rich lakes) may play an important role in abiotic transformation of microbial DOM into distinct humic-like fluorophores in the studied Antarctic lakes.

The %HPO in this study ( $40.4\% \pm 8.5\%$ ) was generally higher than previously reported for lakes in the McMurdo Dry Valleys (Lake Vida, 12%–17%; Lakes Fryxell and Hoare; 17%–24% and 16%–18% as fulvic acid, respectively) and on Ross Island of East Antarctica (Pony Lake, 23%–34%) (Aiken et al., 1996; Cawley et al., 2016, 2013; McKnight et al., 1991), and similar to that of temperate clear water lakes (Table 1 of Sato et al., 2019). The higher %HPO of our samples than of lakes in the McMurdo Dry Valleys suggested the higher abundance of humic substances in the Lützow-Holm Bay and Amundsen Bay lakes and streams despite prevailing photodegradation. Levels of the ubiquitous humic-like components in the studied Antarctic lakes may reflect the balance between the abiotic/biotic production and photodegradation.

The ecological importance of CDOM in Antarctic lakes has been discussed mainly on its light-shading effect for primary producers (McKnight et al., 1994). The formation of humic-like fluorophores could be very important in regulating the depth of the euphotic zone to which visible wavelength radiation can penetrate. This light-shading role that CDOM plays in the studied lakes will be assessed in the future when we will finish analyzing the light penetrating properties of each lake. Another important attribute of Antarctic DOM could be its influence on the position of the thermocline. An example where DOM substantially influenced the thermocline depth could be seen from the depth trend in the water temperature of a saline lake Funazoko (Fig. S2k). The water temperature of Lake Funazoko showed a drastic change; it was approximately 6.8°C at the surface,

increased to 20°C at 1.7 m, then linearly decreased to −5°C at 6.8 m. Because the depth trend in conductivity did not correspond with that of temperature, the stratification was not a halocline but a thermocline, in contrast to other saline lakes (Figs. S2 i, j, l). The very high DOC concentration (92.8 mg C L<sup>-1</sup>) and absorption coefficient ( $a_{254} = 96.1 \text{ m}^{-1}$ ) of the surface water of this lake (Table 1) likely absorbed sunlight energy and increased the temperature. The depth trend in the temperature was likely the balance between heating by the sunlight absorption and cooling by the atmosphere and deeper water.

## 5. Concluding remarks

We found large compositional diversity in our Antarctic samples in terms of DOC/DON, %HPO, UV-absorbing characteristics (SUVA<sub>254</sub> and S<sub>275–295</sub>), and FDOM compositions, and such diversity was largely due to a different degree of photodegradation. Therefore, the hypothesis that DOM in the Antarctic lakes and streams has large compositional diversity and such diversity is mostly introduced by external factors such as photodegradation was confirmed in our study. In addition, we proposed that there were common processes that generated the ubiquitous humic-like components from C<sub>360</sub> (traditionally defined as Peak N) in the Lützow-Holm Bay and Amundsen Bay lakes and streams. Although we consider biotic and photochemical processes primarily responsible for the generation of the ubiquitous humic-like components, more studies are needed to elucidate the underlying

mechanisms.

We note that the CDOM and FDOM mainly studied in this work only captured a small portion of the bulk DOM. Future work should broaden the analytical window and incorporate chemical structural techniques, such as high-field nuclear magnetic resonance spectroscopy and Fourier-transformed ion cyclotron resonance mass spectrometry, to unravel the unique chemistry of Antarctic DOM, which should lead to a better understanding of the other pristine lakes in the world and lower latitude aquatic systems as well.

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626

**Table 1.** The sampling locations, limnological characteristics, and basic optical and chemical properties of DOM for each lake of Lützow-Holm Bay and Amundsen Bay ice-free area.

Area name	Lake name	Type	Water type	Altitude (m)	Length (km)	Area ( $\times 10^4$ m <sup>2</sup> )	Catchment area ( $\times 10^4$ m <sup>2</sup> )	Maximum depth (m)	Sampling depth (m)	Sampling date	Salinity	pH	Temp. (°C)	DOC (mg C L <sup>-1</sup> )	%HPO (%)	<i>a</i> <sub>254</sub> (m <sup>-1</sup> )
A	Amundsen Lake1	ND	others	ND	0.34	1.3	ND	0.5	shore	19 Feb. 2017	0.035	6.97	3.9	0.76	59.2	scatter
A	Amundsen Lake2	ND	others	ND	0.13	0.6	ND	ND	shore	23 Feb. 2017	0.030	7.87	2.7	2.00	45.2	4.42
A	Amundsen Lake3	ND	others	ND	0.15	0.2	ND	ND	shore	23 Feb. 2017	0.193	10.5	3.5	11.1	43.6	42.4
A	Richardson	ND	proglacial	ND	4.76	544	ND	ND	shore	19 Feb. 2017	0.023	6.59	3.5	1.16	30.1	scatter
B	Hiroe	3	proglacial	<b>215</b>	0.96	35.2	ND	<b>9.5</b>	4.0	19 Jan. 2017	0.016	7.84	2.9	0.36	46.5	0.60
B	Breivagnipa Lake1	3	proglacial	285	0.15	0.6	ND	ND	shore	19 Jan. 2017	0.198	7.86	6.4	1.36	36.0	2.04
L	Yukidori	<b>3</b>	others	<b>125</b>	<b>0.28</b>	<b>4.1</b>	ND	<b>8.6</b>	3.5	10 Jan. 2017	0.034	8.22	6.6	0.89	48.0	0.71
L	Yukidori	<b>3</b>	pore	125	<b>0.28</b>	<b>4.1</b>	ND	<b>8.6</b>	pore	10 Jan. 2017	0.311	ND	ND	3.73	57.3	10.2
L	Mitsuike Ue	ND	others	40	0.05	0.2	20.6	1.0	shore	13 Jan. 2017	0.606	8.36	7.8	1.67	32.1	2.63
L	Mitsuike Naka	ND	others	40	0.07	0.2	9.7	1.0	shore	13 Jan. 2017	0.794	8.24	8.2	1.22	29.2	1.68
L	Mitsuike Shita	ND	others	30	0.06	0.2	24.9	1.5	shore	13 Jan. 2017	0.879	8.39	8.2	2.22	31.0	3.55
L	Nurume	<b>1</b>	saline	<b>0</b>	<b>0.31</b>	<b>3.5</b>	17.4	<b>16.2</b>	3.0	12 Jan. 2017	29.1	8.13	6.6	2.24	40.7	3.56
L	Nurume	<b>1</b>	saline	<b>0</b>	<b>0.31</b>	<b>3.5</b>	17.4	<b>16.2</b>	8.5	12 Jan. 2017	32.2	8.13	3.2	6.21	40.7	10.4
L	Nurume	<b>1</b>	saline	<b>0</b>	<b>0.31</b>	<b>3.5</b>	17.4	<b>16.2</b>	14	12 Jan. 2017	54.7	7.17	2.1	11.2	48.4	scatter
L	Heito	<b>3</b>	proglacial	<b>165</b>	0.51	4.9	ND	<b>4.5</b>	shore	9 Jan. 2017	0.024	7.32	1.8	0.64	31.0	1.19
R	Maruwan-Oike	<b>2</b>	proglacial	<b>10</b>	0.63	<b>25.2</b>	ND	<b>37</b>	9.0	17 Jan. 2017	0.015	7.96	3.6	0.56	20.4	scatter
R	Maruwan Kita	2	proglacial	15	<b>0.20</b>	<b>4.6</b>	ND	ND	shore	17 Jan. 2017	0.016	7.44	2.3	1.02	18.3	1.04
R	Maruwan Minami	2	proglacial	15	<b>0.55</b>	<b>2.0</b>	ND	<b>11.5</b>	shore	17 Jan. 2017	0.022	7.15	5.2	0.72	27.6	0.47
Sl	Skallen-Oike	2	others	5	1.10	22.5	115.0	<b>9.2</b>	3.5	4 Jan. 2017	0.132	8.06	4.1	1.43	48.4	2.57
Sl	Skallen-Oike	2	pore	5	1.10	22.5	115.0	<b>9.2</b>	pore	4 Jan. 2017	0.138	ND	ND	7.86	58.7	19.7
Sl	Skallen Lake1	ND	others	30	0.23	1.3	ND	ND	shore	4 Jan. 2017	0.072	8.14	6.5	0.71	40.5	1.34
Sl	Skallen Lake2	ND	others	30	0.11	0.3	ND	ND	shore	4 Jan. 2017	0.050	8.40	8.6	0.78	30.1	1.92
Sl	Skallen Lake3	ND	others	55	0.14	0.9	ND	ND	shore	5 Jan. 2017	0.078	8.50	6.0	1.25	35.9	1.87
Sl	Skallen Lake4	ND	others	75	0.17	1.2	ND	ND	shore	5 Jan. 2017	0.040	8.48	9.5	0.71	32.0	0.67
Sl	Skallen Lake5	ND	others	45	0.11	0.6	ND	ND	shore	5 Jan. 2017	0.034	8.22	10.1	1.07	37.6	2.57
Sl	Glacial Lake	ND	proglacial	10	0.34	4.6	ND	ND	shore	4 Jan. 2017	0.074	7.35	7.7	0.25	41.1	scatter
Sl	Glacial Water	-	proglacial	15	-	-	-	-	melt	4 Jan. 2017	0.017	6.52	1.5	0.33	30.4	scatter
Sr	Oyako	2	others	5	<b>0.43</b>	<b>6.5</b>	280.0	<b>8.0</b>	4.0	27 Dec. 2016	0.364	8.27	5.7	1.19	54.2	2.37
Sr	Naga	3	others	70	<b>0.38</b>	<b>4.8</b>	23.9	<b>10.8</b>	5.0	28 Dec. 2016	0.879	8.56	4.9	3.51	36.1	2.33
Sr	Naga	3	pore	70	<b>0.38</b>	<b>4.8</b>	23.9	<b>10.8</b>	pore	28 Dec. 2016	0.880	ND	ND	ND	ND	27.1
Sr	Nyorai	3	others	130	0.18	0.6	7.5	<b>3.0</b>	1.5	29 Dec. 2016	0.330	8.30	7.1	1.54	44.6	1.94
Sr	Nyorai	3	pore	130	0.18	0.6	7.5	<b>3.0</b>	pore	29 Dec. 2016	0.331	ND	ND	8.47	ND	18.3
Sr	Hotoke	3	others	120	0.12	0.4	6.7	<b>3.0</b>	1.0	30 Dec. 2016	0.421	8.31	4.8	1.08	40.5	1.66
Sr	Hotoke	3	pore	120	0.12	0.4	6.7	<b>3.0</b>	pore	30 Dec. 2016	1.018	ND	ND	ND	ND	19.5
Sr	Kuwai	3	others	160	0.08	0.3	4.0	<b>4.3</b>	2.0	30 Dec. 2016	1.608	7.57	10.3	0.94	38.3	1.19
Sr	Bosatsu	3	others	130	0.19	0.8	10.4	<b>4.8</b>	2.0	25 Jan. 2017	0.181	8.05	8.4	1.38	38.4	1.50

Sr	Jizo	<b>3</b>	others	<b>120</b>	0.10	0.3	2.1	<b>3.1</b>	1.5	25 Jan. 2017	1.227	8.54	8.2	1.46	41.4	2.47
Sr	Himago	1	others	−5	0.08	0.4	9.6	ND	shore	27 Jan. 2017	3.897	8.42	7.8	5.23	43.2	10.3
Sr	Mago	<b>1</b>	others	−5	0.18	1.1	35.3	<b>1.7</b>	shore	27 Jan. 2017	2.226	8.56	7.3	4.57	46.3	8.03
Sr	Sara	<b>1</b>	others	<b>0</b>	0.18	1.1	24.3	<b>2.5</b>	1.5	27 Jan. 2017	0.365	8.75	7.1	2.09	45.0	4.59
Sr	Tokkuri	<b>3</b>	others	45	0.24	1.9	65.5	6.0	shore	28 Jan. 2017	0.873	8.04	6.5	2.77	48.6	3.17
Sr	Kumogata	<b>3</b>	others	65	0.32	3.2	78.5	0.8	shore	28 Jan. 2017	0.648	7.42	6.3	3.90	37.6	4.25
Sr	Hyotan	<b>3</b>	others	<b>75</b>	<b>0.37</b>	<b>5.3</b>	40.0	<b>12.4</b>	6.0	31 Jan. 2017	1.070	8.80	5.9	2.69	42.6	2.44
Sr	Hamagiku	ND	others	95	0.28	1.9	10.5	<b>4.0</b>	2.0	31 Jan. 2017	0.064	8.05	4.2	0.90	38.5	1.33
Sr	Ebi	ND	others	50	0.31	2.2	31.8	<b>6.3</b>	3.0	31 Jan. 2017	0.842	8.81	5.3	2.94	40.3	2.89
Sr	Kikuno	ND	others	55	0.29	1.3	14.3	2.5	shore	31 Jan. 2017	1.232	8.53	4.9	1.75	36.9	2.44
Sr	Hashioki	1	others	10	0.26	1.2	34.1	2.0	shore	27 Jan. 2017	0.267	8.42	7.3	1.73	39.8	3.18
Sr	Skarvsnes Lake2	ND	others	55	0.09	0.4	0.6	ND	shore	31 Jan. 2017	0.964	8.56	5.8	2.57	41.6	3.15
Sr	A-5	ND	others	90	0.07	0.2	7.8	2.0	shore	31 Jan. 2017	0.392	8.59	5.1	3.36	38.6	5.38
Sr	Hechima	<b>3</b>	others	<b>85</b>	0.22	1.4	16.0	<b>6.3</b>	shore	6 Feb. 2017	1.669	8.73	5.2	1.43	44.8	ND
Sr	Magobachi	<b>3</b>	others	<b>60</b>	0.27	1.5	17.0	<b>5.0</b>	shore	6 Feb. 2017	1.017	8.83	5	1.27	44.9	2.12
Sr	Misumi	ND	others	<b>90</b>	0.13	0.7	6.1	6.3	shore	6 Feb. 2017	0.093	8.17	5	1.79	54.3	1.44
Sr	Kobachi	<b>1</b>	saline	<b>25</b>	0.34	4.4	75	<b>12</b>	1.5	27 Jan. 2017	7.37	8.86	7.2	4.39	42.9	5.62
Sr	Kobachi	<b>1</b>	saline	<b>25</b>	0.34	4.4	75	<b>12</b>	6.0	27 Jan. 2017	13.6	8.75	9.7	6.62	43.9	8.53
Sr	Suribachi	<b>1</b>	saline	−33	1.2	58.8	387	<b>32</b>	2.0	27 Jan. 2017	69.8	7.66	11.8	12.2	39.6	6.11
Sr	Suribachi	<b>1</b>	saline	−33	1.2	58.8	387	<b>32</b>	6.0	27 Jan. 2017	121	7.88	20.7	35.1	39.7	17.7
Sr	Funazoko	<b>1</b>	saline	−23	0.66	13.1	118	<b>9.2</b>	2.0	28 Jan. 2017	155	7.40	20.3	92.8	29.9	96.1
Sr	Funazoko	<b>1</b>	saline	−23	0.66	13.1	118	<b>9.2</b>	6.0	28 Jan. 2017	179	7.34	−3.5	146	29.0	158

GPS data (Interactive Map file .kmz) are provided online.

ND = not determined, scatter = samples that contained clay-sized particles after filtration (0.3 µm) and thus absorbance data not presented, while that of Nurume 14 m was due to flocculation upon oxidation.

Length = longitudinal length, melt = glacier melt that drains into Glacial Lake, pore = porewater

Area name; A = Amundsen Bay, B = Breivågnipa, L = Langhovde, R = Rundvågshetta, Sl = Skallen, Sr = Skarvsnes.

Type = geographical type; 1 = Marine relict and lentic lakes, 2 = marine relict and lotic lakes, 3 = non-marine relict lakes.

Geographical and morphologic data in bold italics were compiled from (Kimura et al., 2010), (Kudoh and Tanabe, 2014), and (Imura et al., 2003).

Table 2. Average (standard error) values of the optical parameters in each water type.

	$a_{254}$	SUVA <sub>254</sub>	S <sub>275–295</sub>	FI	HIX	BIX	C <sub>325</sub> /DOC	C <sub>415</sub> /DOC	C <sub>360</sub> /DOC	C <sub>500</sub> /DOC	C <sub>440</sub> /DOC	C <sub>460</sub> /DOC	C <sub>300</sub> /DOC
Water type	(m <sup>-1</sup> )	(L mg C <sup>-1</sup> m <sup>-1</sup> )	(×10 <sup>-3</sup> nm <sup>-1</sup> )				(×10 <sup>-3</sup> RU L mg C <sup>-1</sup> )	(×10 <sup>-3</sup> RU L mg C <sup>-1</sup> )	(×10 <sup>-3</sup> RU L mg C <sup>-1</sup> )	(×10 <sup>-3</sup> RU L mg C <sup>-1</sup> )	(×10 <sup>-3</sup> RU L mg C <sup>-1</sup> )	(×10 <sup>-3</sup> RU L mg C <sup>-1</sup> )	(×10 <sup>-3</sup> RU L mg C <sup>-1</sup> )
Proglacial lake	1.07 <sup>a</sup>	0.579 <sup>a</sup>	25.6 <sup>ab</sup>	1.452 <sup>a</sup>	0.335 <sup>a</sup>	0.935 <sup>a</sup>	38.9 <sup>c</sup>	8.75 <sup>c</sup>	7.69 <sup>c</sup>	5.18 <sup>c</sup>	2.12 <sup>c</sup>	3.86 <sup>bc</sup>	15.5 <sup>a</sup>
N = 5–9	(0.62)	(0.211)	(5.35)	(0.098)	(0.070)	(0.100)	(21.7)	(4.43)	(3.80)	(3.62)	(1.04)	(1.79)	(8.80)
Saline lake	38.2 <sup>bc</sup>	0.486 <sup>a</sup>	49.5 <sup>c</sup>	1.776 <sup>c</sup>	0.384 <sup>ab</sup>	1.218 <sup>b</sup>	9.29 <sup>c</sup>	4.18 <sup>c</sup>	4.00 <sup>c</sup>	1.45 <sup>c</sup>	0.98 <sup>c</sup>	2.35 <sup>c</sup>	6.43 <sup>b</sup>
N = 8–9	(57.4)	(0.190)	(12.9)	(0.171)	(0.117)	(0.181)	(5.76)	(3.76)	(3.06)	(1.33)	(0.88)	(1.67)	(3.74)
Other lake <sup>1</sup>	2.88 <sup>a</sup>	0.652 <sup>a</sup>	32.2 <sup>b</sup>	1.581 <sup>b</sup>	0.488 <sup>c</sup>	0.900 <sup>a</sup>	22.7 <sup>c</sup>	9.87 <sup>c</sup>	6.85 <sup>c</sup>	4.49 <sup>c</sup>	2.65 <sup>c</sup>	7.39 <sup>ab</sup>	9.80 <sup>ab</sup>
N = 32–33	(2.00)	(0.206)	(6.76)	(0.086)	(0.074)	(0.054)	(12.9)	(4.81)	(3.82)	(2.44)	(1.48)	(3.02)	(5.21)
Porewater	19.0 <sup>ab</sup>	1.071 <sup>c</sup>	18.1 <sup>a</sup>	1.781 <sup>c</sup>	0.481 <sup>bc</sup>	0.936 <sup>a</sup>	111 <sup>ab</sup>	91.4 <sup>a</sup>	37.7 <sup>a</sup>	23.9 <sup>a</sup>	27.0 <sup>a</sup>	6.56 <sup>abc</sup>	13.4 <sup>ab</sup>
N = 3–5	(6.02)	(0.124)	(0.48)	(0.062)	(0.105)	(0.060)	(4.05)	(23.5)	(12.0)	(1.90)	(4.43)	(1.83)	(1.29)
Stream	1.09 <sup>a</sup>	0.862 <sup>bc</sup>	17.2 <sup>a</sup>	1.569 <sup>ab</sup>	0.573 <sup>d</sup>	0.836 <sup>a</sup>	37.9 <sup>b</sup>	32.1 <sup>b</sup>	11.3 <sup>b</sup>	13.3 <sup>b</sup>	8.58 <sup>b</sup>	7.79 <sup>a</sup>	12.1 <sup>ab</sup>
N = 16	(0.33)	(0.200)	(3.05)	(0.088)	(0.099)	(0.060)	(11.6)	(14.9)	(3.76)	(6.01)	(4.61)	(3.11)	(6.37)

<sup>1</sup> Excluding Amundsen Lake 3.

Different letters in each parameter indicate significant differences among water types at  $P < 0.05$  based on Tukey-Kramer's multiple comparisons of means.

$a_{254}$  = absorption coefficient of dissolved organic matter at 254 nm, SUVA<sub>254</sub> = specific UV absorbance at 254 nm, S<sub>275–295</sub> = spectral slope determined between 275 nm and 295 nm, FI = fluorescence index, HIX = humification index, BIX = biological index, and C<sub>x</sub>/DOC = fluorescence intensity of each PARAFAC component normalized by its sample dissolved organic carbon concentration



Fig. 1. Distribution of dissolved C or N properties (a–d) and UV absorption properties (e, f) of Lützow-Holm Bay and Amundsen Bay lake and stream samples along the salinity gradient (log scaled). Samples were categorized into six water types and represented by color. Open black circle (“penguin”) represents a sample collected at Amundsen Lake 3 with abundant phytoplankton because of penguin guano inputs. Error bar for %HPO indicates one standard deviation of quadruplicate experiments. Correlation coefficients between log DOC, log DON, DOC/DON, SUVA<sub>254</sub>, and S<sub>275–295</sub> with log salinity were 0.83, 0.78, 0.51, –0.38, and 0.80, respectively (all  $P < 0.01$ ).

Fig. 2. Counter plots of three-dimensional spectra of the identified PARAFAC components from Lützow-Holm Bay and Amundsen Bay lake and stream samples. The components were named according to their rounded fluorescence emission maximum.

Fig. 3. Relationships between relative contribution of PARAFAC components and the spectral slope (S<sub>275–295</sub>). %C(protein) = relative contribution of the sum of the protein-like components (C<sub>325</sub> and C<sub>300</sub>), while %C(ubiquitous humic) = relative contribution of the sum of the ubiquitous humic-like components (C<sub>415</sub>, C<sub>500</sub>, and C<sub>440</sub>). C<sub>360</sub> = analogous to traditional Peak N (Coble et al., 1998), while C<sub>460</sub> = photorefractory UVC humic-like component. Samples were categorized into six water types as in Fig. 1. Tentative interpretations on the pattern and directions of changes due to environmental processing (salinity-induced precipitation and photodegradation) are shown (see main text). In (b) and (c), saline lake samples are labeled as follows: S2 & 6 = Suribachi 2 & 6 m, F2 & 6 = Funazoko 2 & 6 m, K1.5 & 6 = Kobachi 1.5 & 6 m, N3 & 8.5 = Nurume 3 & 8.5 m, where numbers after the lake names represent the depths at which samples were collected (See Table 1).

Fig. 4. Correlation between C<sub>360</sub>/DOC versus C<sub>415</sub>/DOC, C<sub>440</sub>/DOC, and C<sub>500</sub>/DOC (ubiquitous humic-like fluorophore). The inset shows correlation matrix among the levels of the identified PARAFAC components (Fmax/DOC) with Spearman’s correlation coefficient ( $\rho$ ) indicated by color.

Figure 1

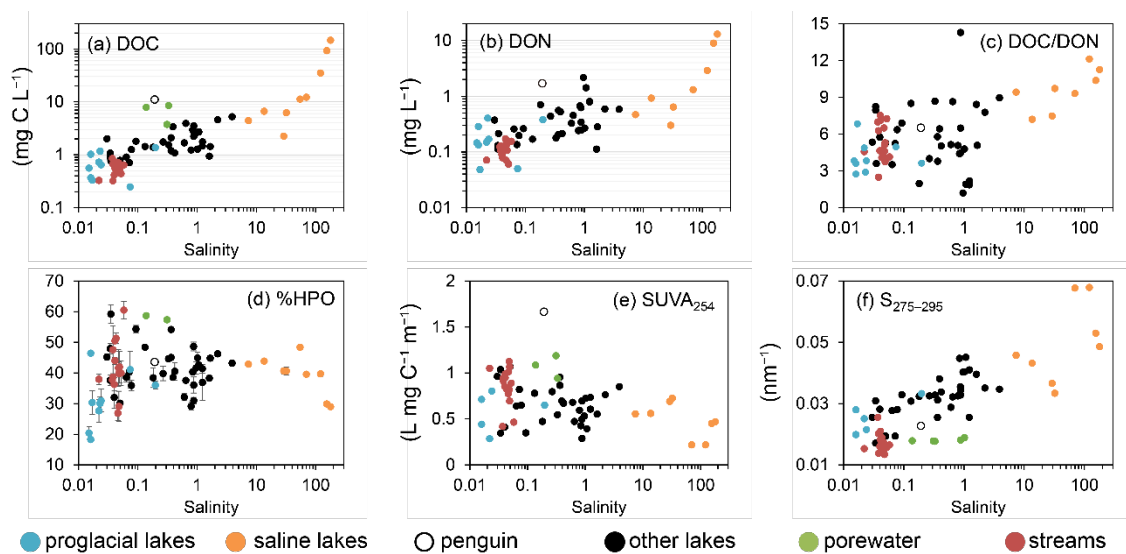


Figure 2

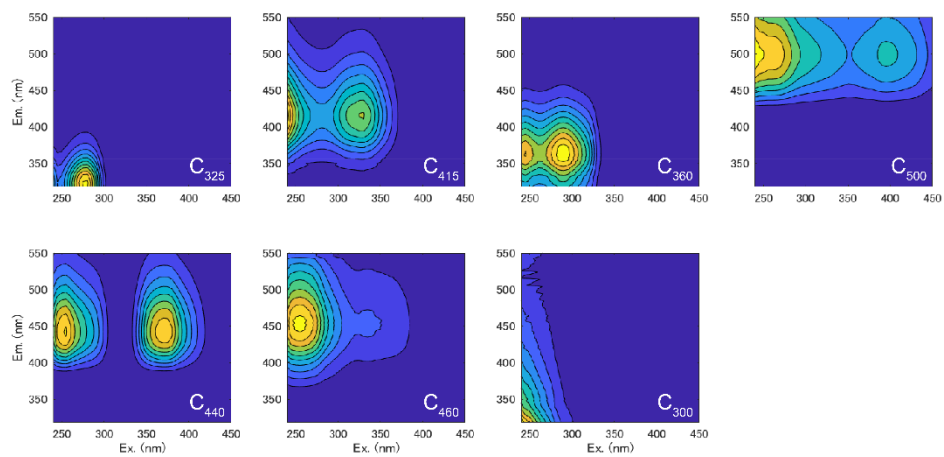


Figure 3

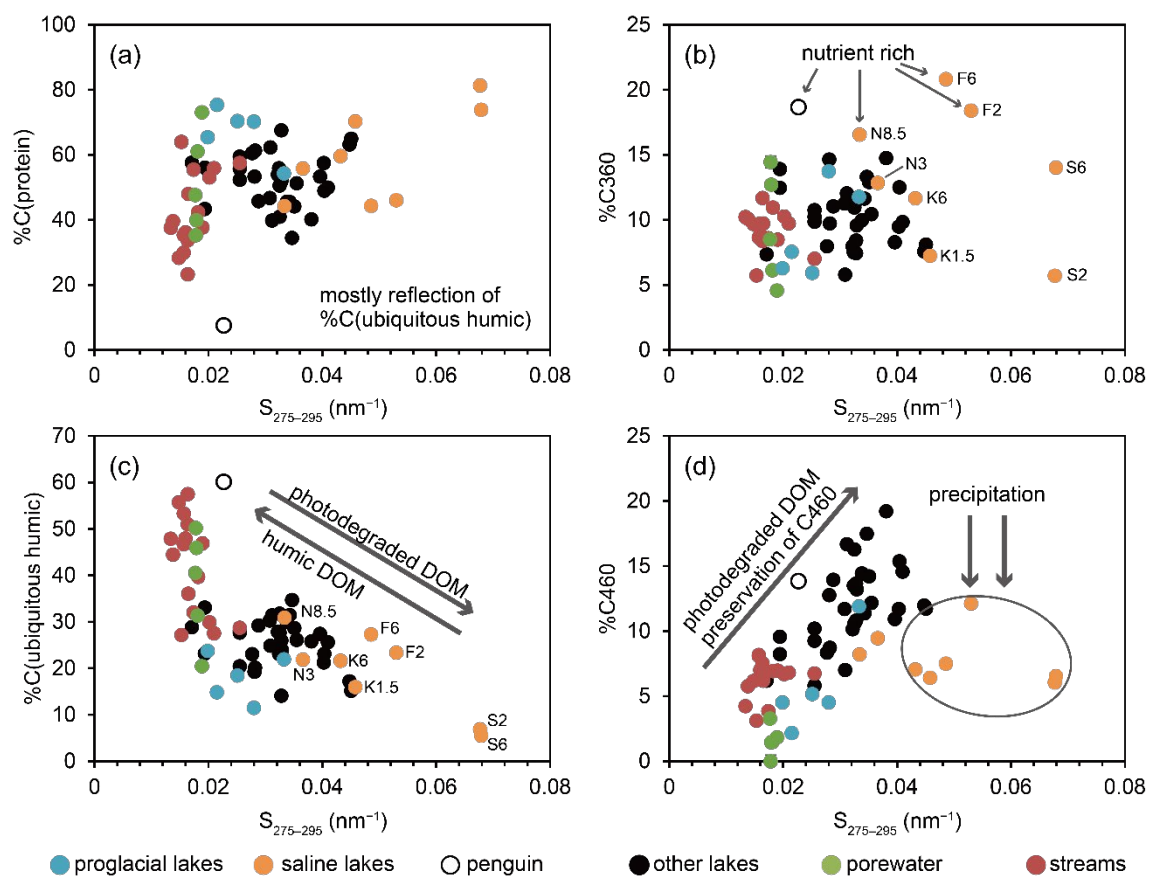
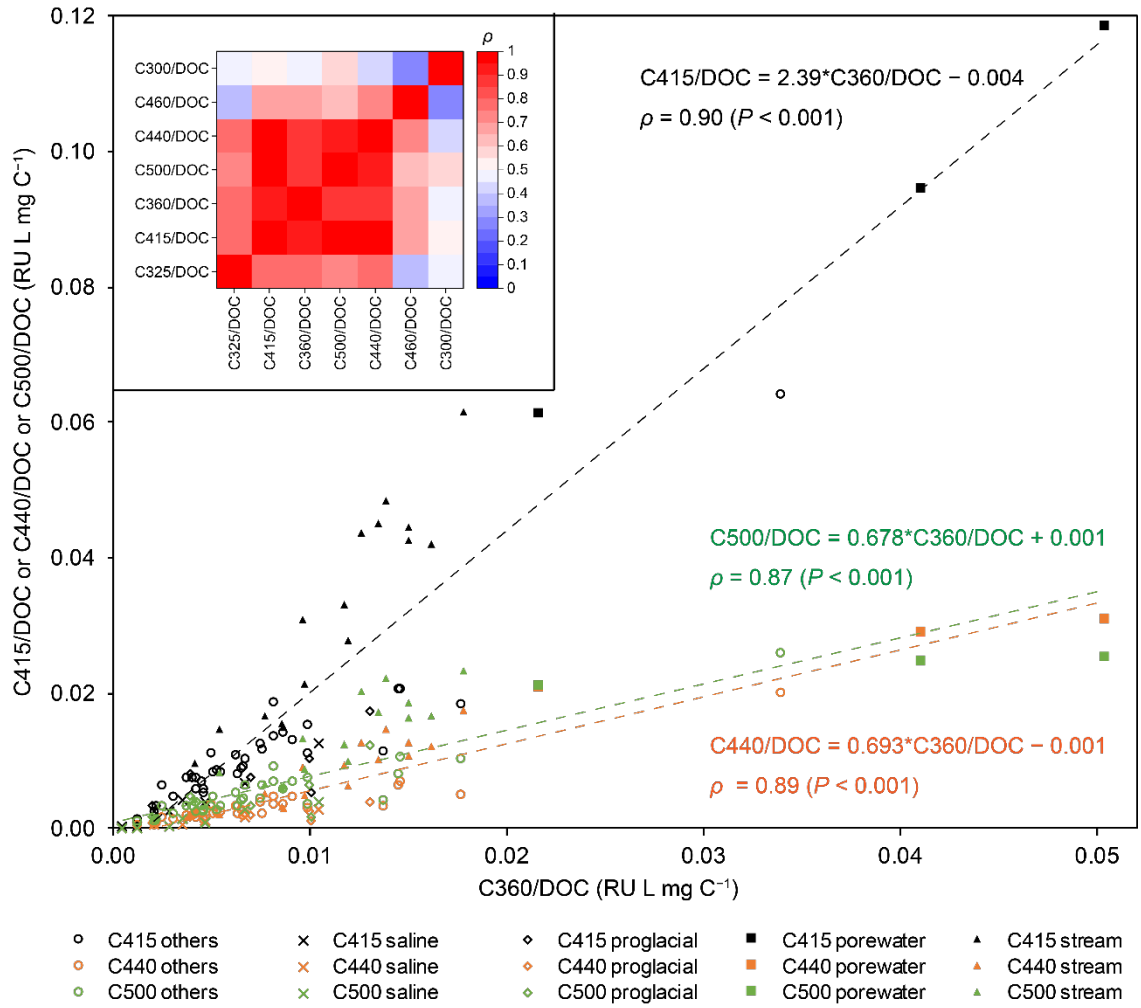


Figure 4



Journal: Water Research

Supporting information for:

# **Origin, distributions, and environmental significance of ubiquitous humic-like fluorophores in Antarctic lakes and streams**

**Morimaru Kida<sup>1†‡\*</sup>, Taichi Kojima<sup>1</sup>, Yukiko Tanabe<sup>2, 3</sup>, Kentaro Hayashi<sup>4</sup>, Sakae Kudoh<sup>2, 3</sup>, Nagamitsu Maie<sup>5</sup>, Nobuhide Fujitake<sup>1</sup>**

<sup>1</sup> Faculty of Agriculture, Graduate School of Agricultural Science, Kobe University, 1-1 Rokkodai, Nada, Kobe, Hyogo 657-8501, Japan

<sup>2</sup> National Institute of Polar Research, Research Organization of Information and Systems, 10-3 Midori-cho, Tachikawa, Tokyo 190-8518, Japan

<sup>3</sup> Department of Polar Science, SOKENDAI (The Graduate University for Advanced Studies), 10-3 Midori-cho, Tachikawa, Tokyo 190-8518, Japan

<sup>4</sup> Institute for Agro-Environmental Sciences, NARO, 3-1-3 Kannondai, Tsukuba, Ibaraki 305-8604, Japan

<sup>5</sup> School of Veterinary Medicine, Kitasato University, Towada, Aomori 034-8628, Japan

† Research Fellow of Japan Society for the Promotion of Science

‡ Present address: Research Group for Marine Geochemistry (ICBM-MPI Bridging Group), Carl von Ossietzky University of Oldenburg, Institute for Chemistry and Biology of the Marine Environment (ICBM), Carl-von-Ossietzky-Str. 9-11, 26129 Oldenburg, Germany

\* corresponding author e-mail: [morimaru.kida@people.kobe-u.ac.jp](mailto:morimaru.kida@people.kobe-u.ac.jp); Tell: +81-78-803-5846

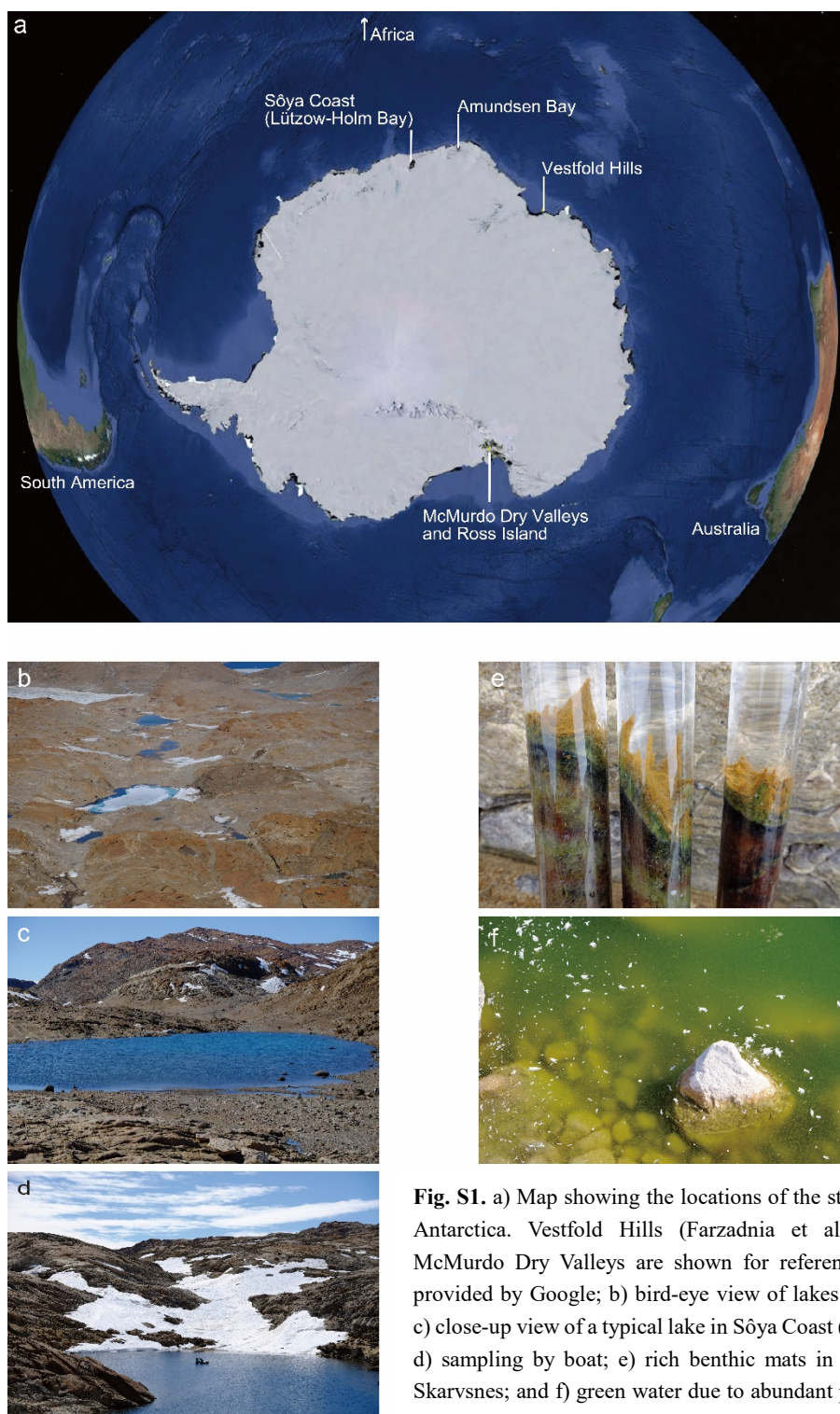
ORCID ID: 0000-0002-9908-2012 (M.K.)

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Figure S1: Map and pictures



**Fig. S1.** a) Map showing the locations of the studied areas in Antarctica. Vestfold Hills (Farzadnia et al., 2017) and McMurdo Dry Valleys are shown for reference. Map data provided by Google; b) bird-eye view of lakes in Skarvsnes; c) close-up view of a typical lake in Sôya Coast (Lake Oyako); d) sampling by boat; e) rich benthic mats in Lake Hotoke, Skarvsnes; and f) green water due to abundant phytoplankton (Amundsen Lake 3 “penguin lake”, Amundsen Bay).

## SI 1. Optical index

We calculated optical indices from the absorbance and EEM data to estimate the structural properties and origin of DOM. Spectral slope ( $S_{275-295}$ ), which has been related to apparent molecular weight (MW) and the degree of photodegradation (Helms et al., 2008) and used as a source differentiation (terrestrial vs marine) tool in estuarine studies (Fichot and Benner, 2012), was calculated using a linear fit to the natural log-transformed  $a_\lambda$  spectrum in the ranges of 275–295 nm (coefficient of determination  $R^2 > 0.97$ ; Fig. S3a). Typically, a higher (steeper)  $S_{275-295}$  value suggests a decrease in molecular weight and aromaticity of DOM and an increase in the degree of photodegradation (Helms et al., 2008). DOC-specific ultraviolet absorbance ( $SUVA_{254}$ ), which is related to aromaticity of humic substances (Weishaar et al., 2003) and can be used as a metric for %HPO (Kida et al., 2018) and the light absorbing ability of DOM, was calculated by dividing the UV absorbance ( $m^{-1}$ ) measured at  $\lambda = 254$  nm by the DOC concentration ( $mg\ C\ L^{-1}$ ). The lowest absorbance at 254 nm with a 5-cm path length cuvette (0.01) was sufficiently higher than the instrument measurement accuracy of 0.002 when the absorbance is  $< 0.5$  (manufacturer provided).

The measurement locations of the fluorescent indices are provided in Fig. S5 for visualization purposes. The fluorescence index (FI) was calculated as the ratio of emission intensities at 470 and 520 nm ( $I_{470}/I_{520}$ ) at 370 nm excitation (Cory et al. 2010). FI values are somewhat dependent on the instrument used even after instrument-specific response correction, and the FluoroMax gives the FI value of 1.42–1.55 for microbial end-member and 1.21 for soil end-member fulvic acids (Cawley et al., 2013; Cory et al., 2010). Four samples were found not to have humic fluorophores, and thus FI of these samples is not presented (Fig. S4). The biological index (BIX) was calculated as  $I_{380}/I_{430}$  at 310 nm excitation and it estimates autochthonous biological production and subsequent heterotrophic consumption by bacteria in water environments. High values ( $> 1$ ) of BIX correspond to DOM of predominantly autochthonous origin and to freshly released DOM in water, whereas lower values (0.6–0.7) correspond to low DOM production in natural waters (Huguet et al., 2009). The humification index (HIX) was calculated as the sum of  $I_{435-480}$  divided by the sum of  $I_{300-345} + I_{435-480}$  at 255 nm excitation and it ranges from 0 to 1 with increasing degrees of humification (Ohno, 2002).



## SI 2. PARAFAC

We analyzed the EEMs by parallel factor analysis (PARAFAC) to mathematically deconvolute them into distinct fluorescent components with independent spectral characteristics. The bottom water was collected at six lakes to increase the number of samples for PARAFAC, but their chemical properties are not discussed in this study because they were likely the mixture of the porewater and corresponding lake water. We used MATLAB R2017b with the *drEEM* toolbox (v 0.4.0) (Murphy et al., 2013). Despite the large fluorescence intensity differences among samples, EEMs were not normalized to total signals because normalization led to a worse and unstable model. During the exploratory phase, eight outliers were removed based on their leverage, and emission wavelengths below 318 nm were removed. PARAFAC was applied to the resulting EEM dataset (73 samples  $\times$  43 excitations  $\times$  117 emissions). The model was constrained to non-negative values. Models (5–7 components) were developed with a convergence criterion of  $10^{-10}$ . Each model was iterated 50 times after initializing with random starting values, and only the least squares solution was retained. The final seven-component model accounted for  $> 99.95\%$  of the measured spectral variation and had a low core consistency value (0.48%). The appropriate component number was determined using residual analysis and external validation against published models (Murphy et al., 2018). Due to the limited number of samples (Murphy et al., 2013), split-half validation with random splitting did not produce a stable result (not always validated). The external validation was achieved by quantitatively comparing the identified fluorescent components with the published data using an open-access spectral database (OpenFluor; Murphy et al., 2014). On OpenFluor, one can compare fluorescence datasets and determine the Tucker congruence coefficient (TCC) between pairs of excitation ( $TCC_{ex}$ ) and emission ( $TCC_{em}$ ) spectra. If an identified fluorescent component has a strong match ( $TCC_{ex \times em} > 0.95$ ), it means that the same and/or a spectrally indistinguishable validated component has been reported in the literature, and that the identified fluorescent component has legitimate spectral characteristics. The post-modeling residual analysis confirmed that the seven-component model adequately reproduced each EEM. This type of validation has been recently proposed for the one-sample PARAFAC approach (Murphy et al., 2018; Wünsch et al., 2017), and it was enabled here for the ‘traditional’ multi-sample PARAFAC owing to the open database. We note that a five-component model was also validated, but it often failed to adequately represent protein-like fluorescence and visible wavelength regions ( $Ex > 350$  nm,  $Em > 400$  nm). Also, three of the five identified components in the five-component model had much smaller (0–4) number of strong matches ( $TCC_{ex \times em} > 0.95$ ) on OpenFluor, implying that the five-component model was not adequate. After model building, the outliers were projected onto the validated model to obtain scores for them. However, four out of the eight samples were not adequately modeled (judged by residuals); thus, their PARAFAC results are not presented.

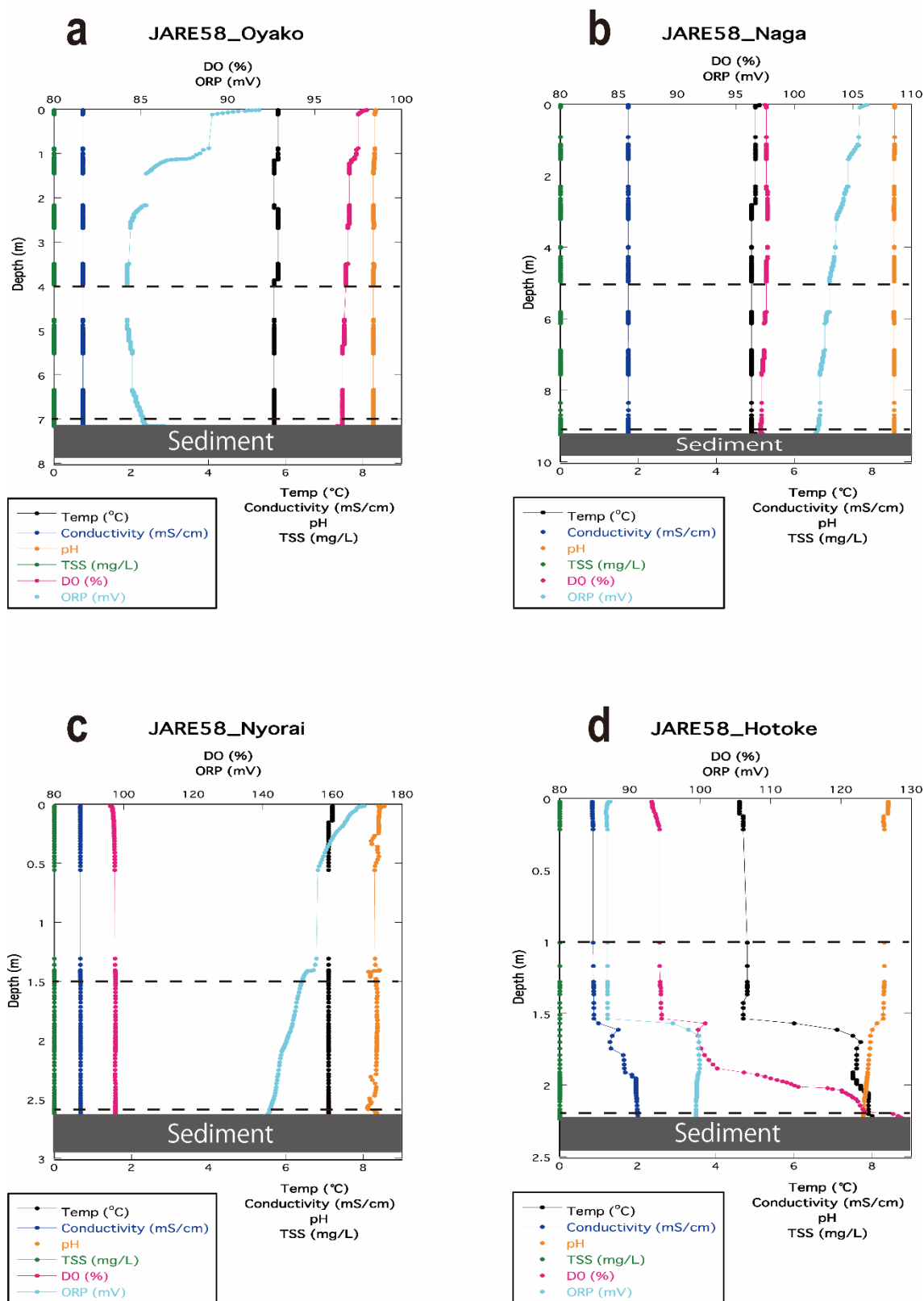
## Table S1: Stream data

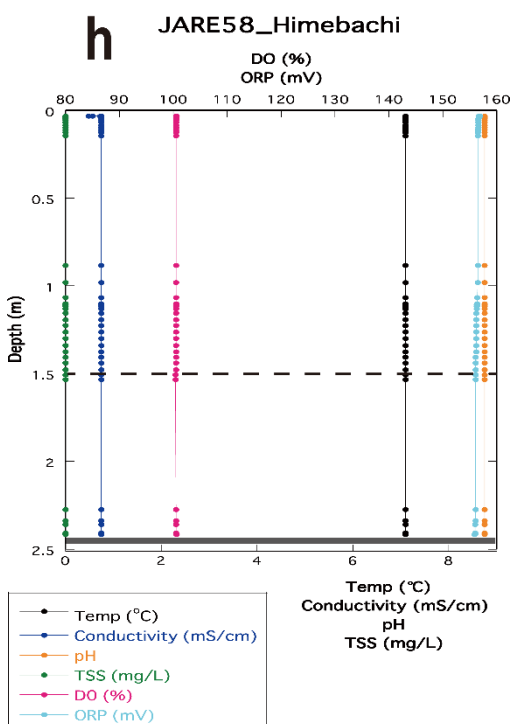
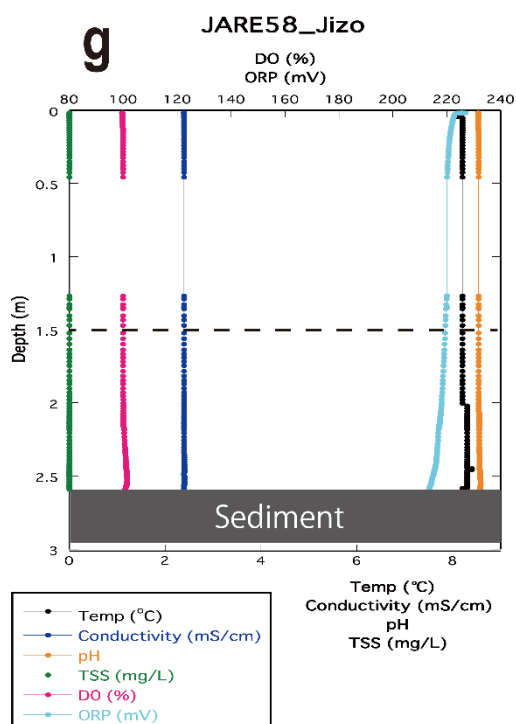
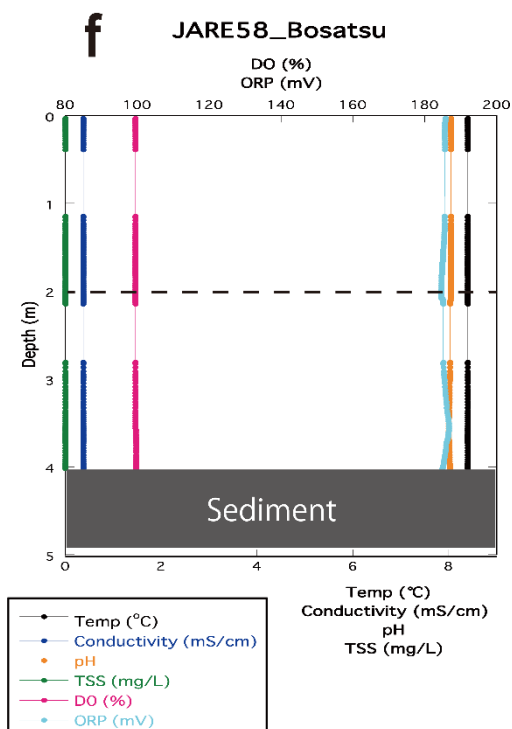
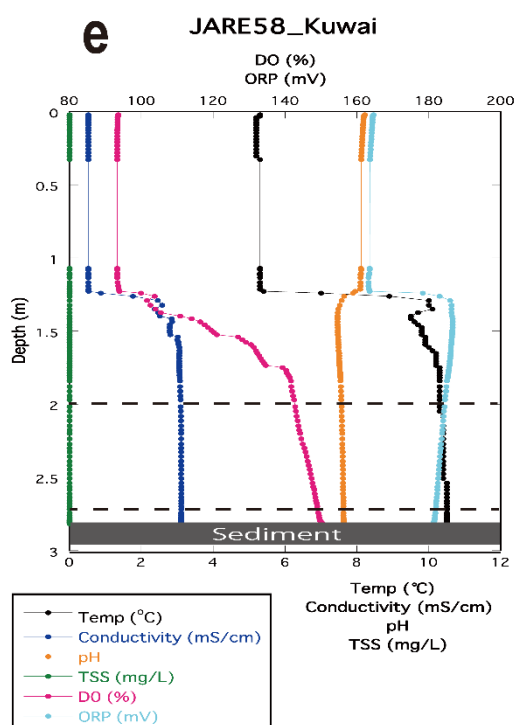
**Table S1.** The sampling locations and basic optical and chemical properties of DOM for samples collected from streams in Yukidori and Yatsude Valleys in Langhovde ice-free area.

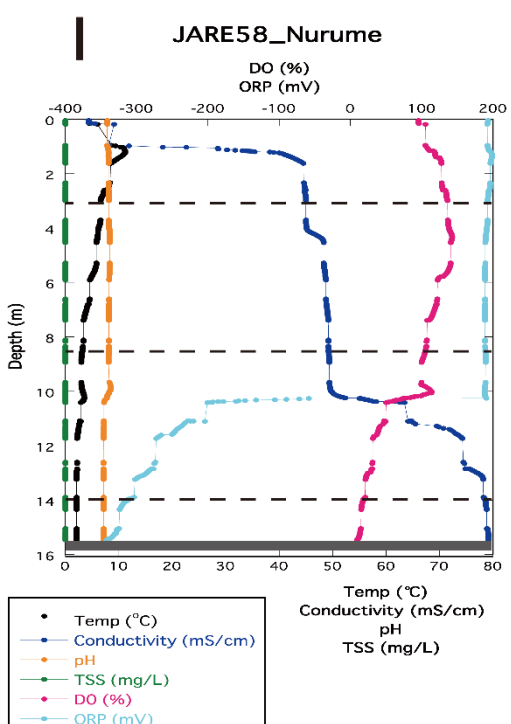
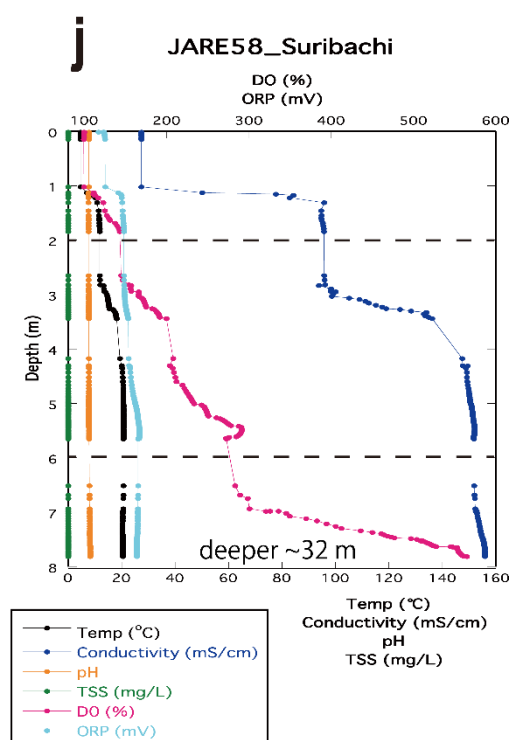
Stream name	Latitude (decimal)	Longitude (decimal)	Sampling date	Salinity	pH	Water temp. (°C)	DOC (mg C L <sup>-1</sup> )	%HPO (%)	$a_{254}$ (m <sup>-1</sup> )
Yukidori 1	-69.24105	39.77995	8 Jan. 2017	0.022	8.59	1.2	0.326	38.0	0.79
Yukidori 2	-69.24072	39.77785	8 Jan. 2017	0.041	10.0	1.5	0.724	44.0	1.65
Yukidori 3	-69.24052	39.77172	8 Jan. 2017	0.038	8.59	2.5	0.664	38.7	1.40
Yukidori 4	-69.24057	39.76347	8 Jan. 2017	0.041	8.10	3.6	0.553	50.5	1.21
Yukidori 6	-69.24075	39.75300	11 Jan. 2017	0.037	7.08	8.3	0.850	37.7	0.82
Yukidori 7	-69.24090	39.74454	11 Jan. 2017	0.043	7.06	4.8	0.481	51.3	0.91
Yukidori 8	-69.24174	39.73134	11 Jan. 2017	0.049	7.06	7.9	0.525	40.2	1.29
Yukidori 9	-69.24226	39.71842	11 Jan. 2017	0.048	6.78	8.0	0.677	39.8	1.75
Yatsude 1	-69.26201	39.78727	9 Jan. 2017	0.052	7.20	1.8	0.436	40.0	0.89
Yatsude 3	-69.25331	39.76625	9 Jan. 2017	0.040	7.11	1.0	0.413	36.3	0.81
Yatsude 4	-69.25268	39.76462	9 Jan. 2017	0.047	7.17	0.9	0.460	41.6	1.07
Yatsude 5	-69.25173	39.76064	9 Jan. 2017	0.049	7.20	1.1	0.564	42.0	0.90
Yatsude 6	-69.24942	39.75051	9 Jan. 2017	0.046	7.03	2.2	0.707	26.9	1.26
Yatsude 7	-69.24752	39.74376	9 Jan. 2017	0.048	6.93	2.1	0.690	29.1	1.30
Yatsude 8	-69.24531	39.73379	9 Jan. 2017	0.038	7.07	5.0	0.320	47.6	0.70
Yatsude 9	-69.24416	39.71998	16 Jan. 2017	0.058	7.13	3.0	0.643	60.5	0.69

Yukidori 5 = Lake Yukidori, Yatsude 2 = Lake Heito (See Table 1).

Figure S2: Water chemistry depth profile of the sampled lakes







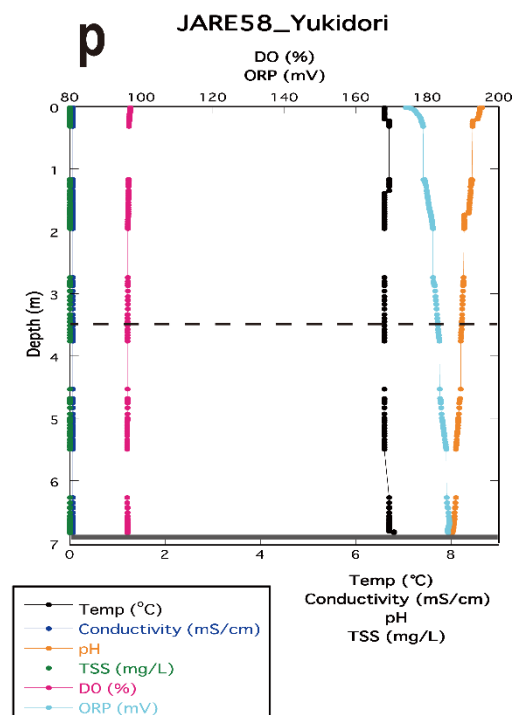
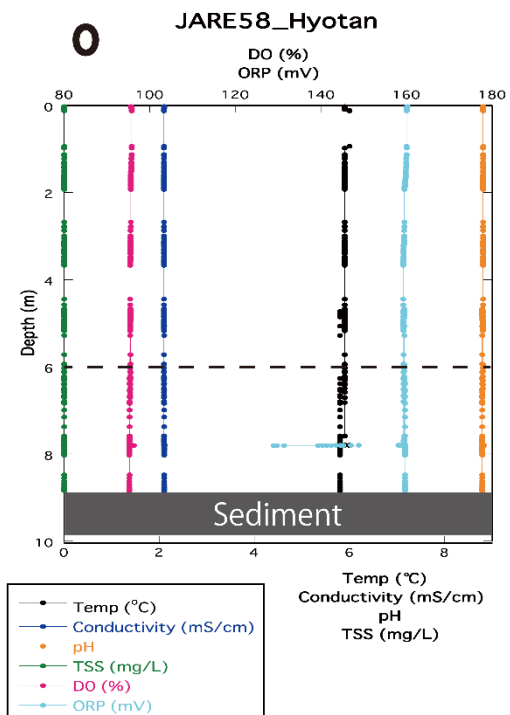
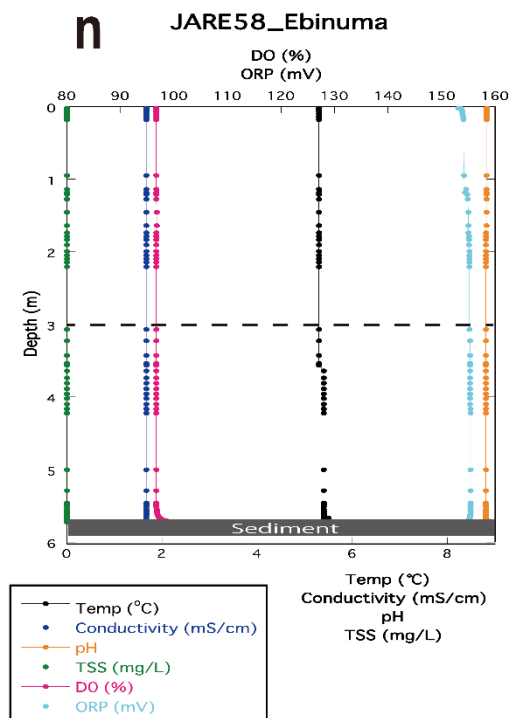
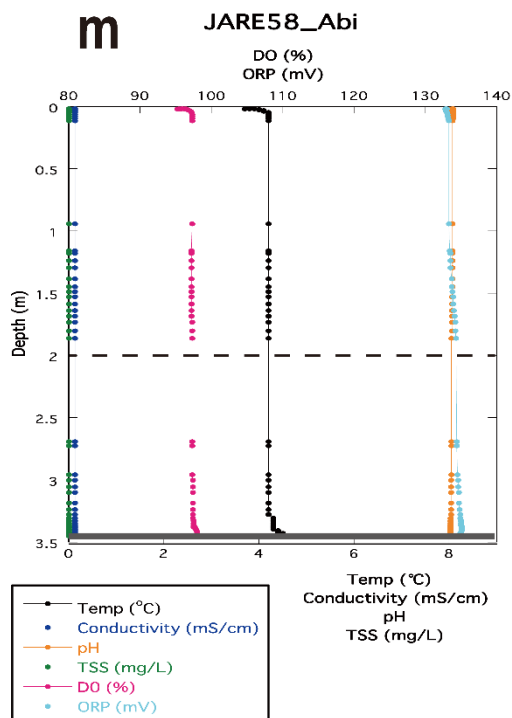
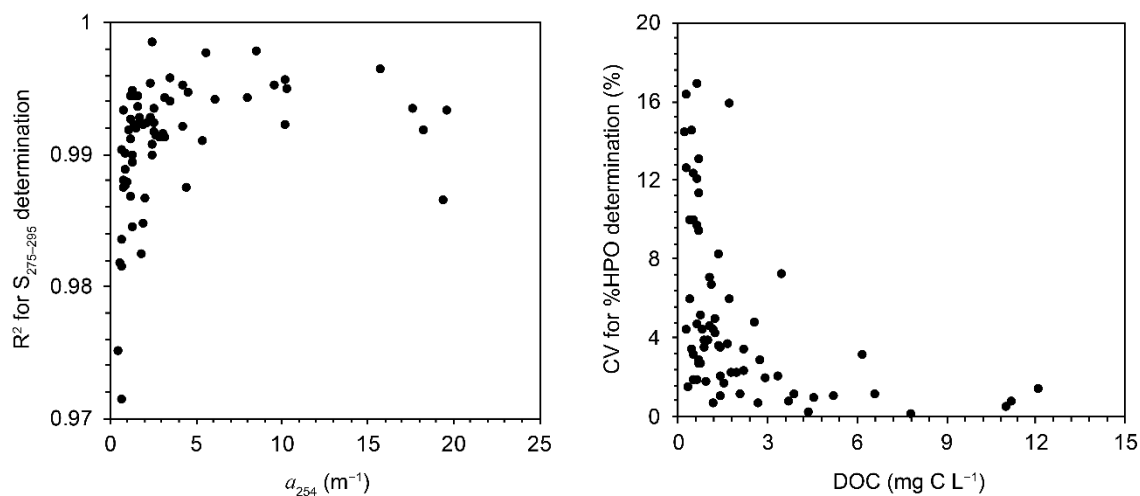




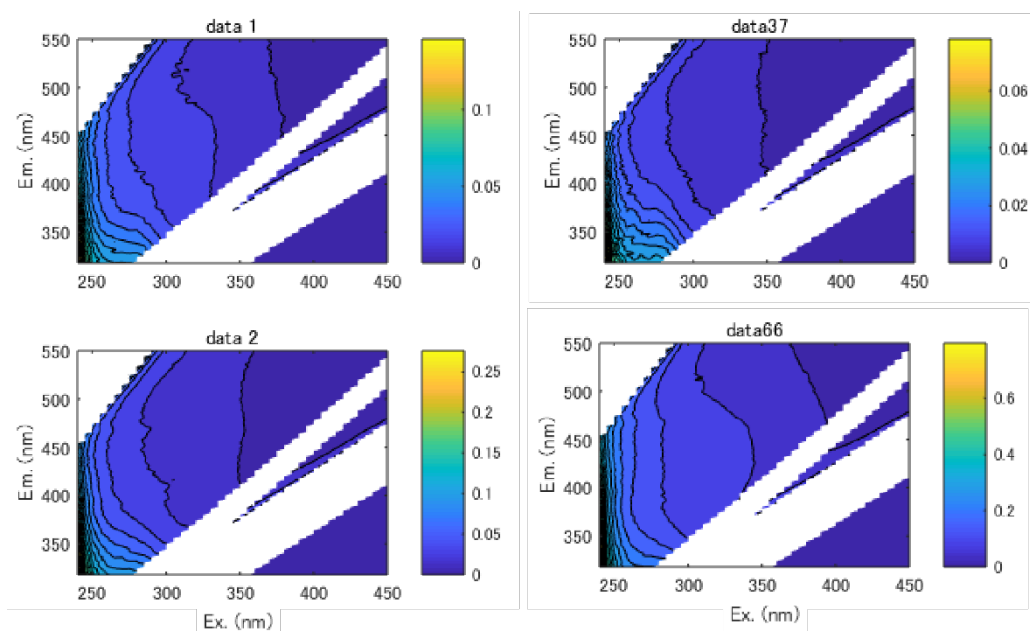
Figure S3: Quality assurance for the  $S_{275-295}$  and %HPO determination



**Fig. S3.** Quality assurance. Coefficient of determination ( $R^2$ ) for the  $S_{275-295}$  determination by linear regression of natural log-transformed  $a$  spectra as a function of  $a_{254}$  and coefficient of variation (CV) of the quadruplicate determination of the proportion of hydrophobic fraction (HPO) of DOM by the DAX-8 batch adsorption as a function of DOC.

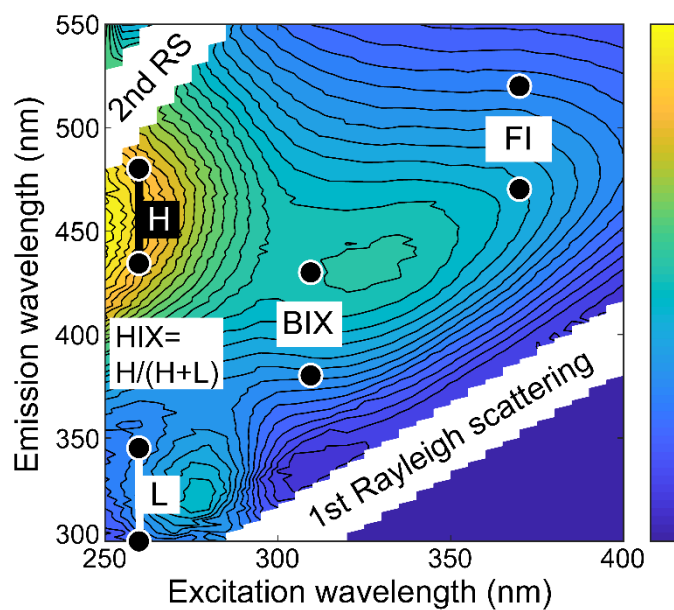


Figure S4: Samples without humic fluorophores



**Fig. S4.** Samples without humic fluorophores. Fluorescence index (FI) of these samples is not presented since  $FI \sim 1$  and meaningless due to the absence of fluorescence in the region where FI is calculated ( $Ex = 370$  nm). Also, the established PARAFAC model could not adequately represent EEMs of these samples, so their PARAFAC results are not presented. The samples are: glacial lake (data 1) and glacial water (data 2) in Skarvsnes, Maruwan-Oike in Rundvågshetta (data 37), and Amundsen Lake1 (data 66) in Amundsen Bay.

Figure S5: Measurement locations of FDOM indices



**Fig. S5.** Excitation emission matrix and measurement locations of fluorescent indices. Fluorescent intensity is presented in color (arbitrary unit). The two points used in the fluorescence index (FI,  $I_{470}/I_{520}$ ) and biological index (BIX,  $I_{380}/I_{430}$ ) are represented by filled circles at an excitation of 370 nm and 310 nm, respectively. The humification index (HIX) is calculated by area under an emission of 435–480 nm (H) divided by area under an emission of 435–480 nm (H) + 300–345 nm (L) at an excitation of 254 nm. In this study, HIX was calculated at an excitation of 255 nm.

Table S2: Nutrient data

**Table S2.** Nutrients concentrations of the studied lakes and streams. Blank cells indicate that nutrients concentrations were below the detection limit of the analyzer. See Table 1 for the detailed information of lakes.

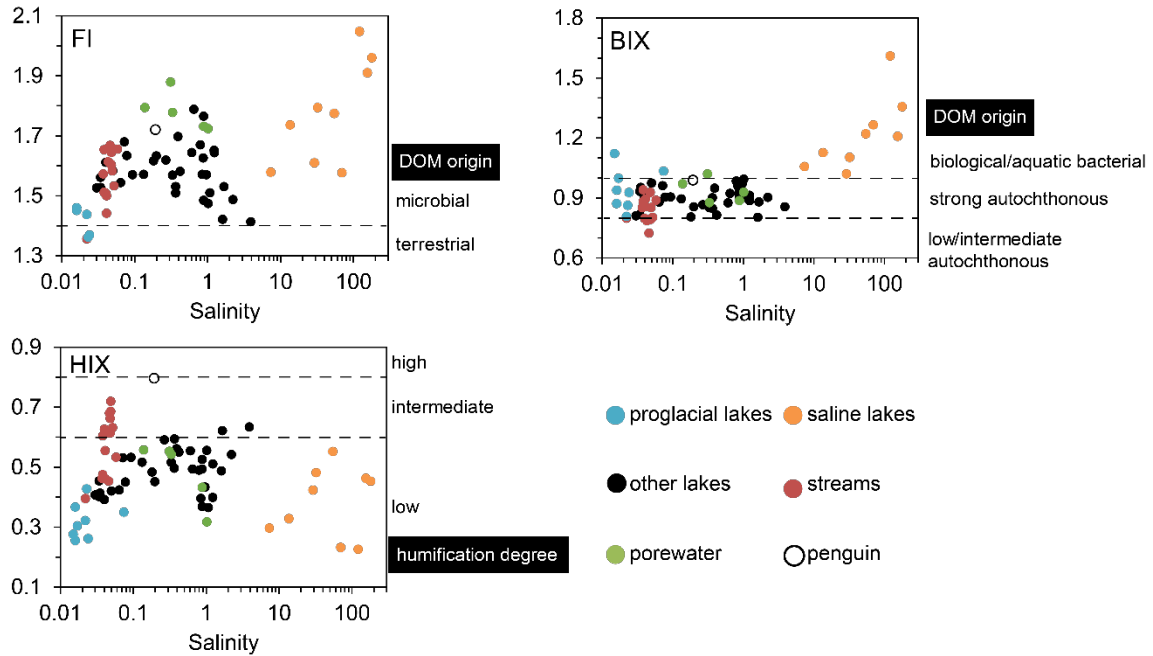
Area name	Lake/stream name	Depth	Salinity	NO <sub>2</sub> <sup>-</sup> μmol/L	NO <sub>3</sub> <sup>-</sup> μmol/L	NH <sub>4</sub> <sup>+</sup> μmol/L	%DON	PO <sub>4</sub> <sup>3-</sup> μmol/L	SiO <sub>3</sub> <sup>2-</sup> μmol/L
Amundsen	Amundsen Lake1	shore	0.04	0.08	18.23	1.08	44.0	4.35	73.17
Amundsen	Amundsen Lake2	shore	0.03			0.59	97.8	0.41	3.29
Amundsen	Amundsen Lake3	shore	0.19	3.82	276.34	11.94	29.3	31.70	173.49
Amundsen	Richardson	shore	0.02		Tr	0.35	98.5	3.17	7.77
Breivågnipa	Hiroe	4 m	0.02				100		2.60
Breivågnipa	Breivagnipa Lake1	shore	0.20			0.44	98.4		97.09
Langhovde	Yukidori (Yukidori 5)	3.5 m	0.03		2.33	0.27	75.6	Tr	16.17
Langhovde	Yukidori (Yukidori 5)	pore	0.31	NA	NA	NA	NA	NA	NA
Langhovde	Mitsuike Ue	shore	0.61	Tr	5.20	0.42	80.4		129.53
Langhovde	Mitsuike Naka	shore	0.79	Tr	3.19	0.25	83.2		204.53
Langhovde	Mitsuike Shita	shore	0.88	Tr	4.88	0.82	81.0		159.57
Langhovde	Nurume	3 m	29.11			0.17	99.2	0.24	6.01
Langhovde	Nurume	8.5 m	32.20	Tr		3.49	92.9	0.98	8.61
Langhovde	Nurume	14 m	54.73	0.08	1.22	479.10	NA	45.63	119.44
Langhovde	Heito (Yatsude 2)	shore	0.02	0.20	3.38	0.17	76.3		4.50
Rundvågshetta	Maruwan-Oike	9 m	0.02			0.06	99.4		2.56
Rundvågshetta	Maruwan Kita	shore	0.02		Tr	0.43	97.2		2.96
Rundvågshetta	Maruwan Minami	shore	0.02			0.09	99.2		14.93
Skallen	Skallen-Oike	3.5 m	0.13			0.20	98.4		73.86
Skallen	Skallen-Oike	pore	0.14	NA	NA	NA	NA	NA	NA
Skallen	Skallen Lake1	shore	0.07			0.17	98.2		30.08
Skallen	Skallen Lake2	shore	0.05		Tr	Tr	96.0		39.04
Skallen	Skallen Lake3	shore	0.08			0.16	98.9		52.10
Skallen	Skallen Lake4	shore	0.04			Tr	99.5		28.10
Skallen	Skallen Lake5	shore	0.03			Tr	98.9		15.95
Skallen	Glacial Lake	shore	0.07			Tr	99.8	Tr	12.69
Skallen	Glacial Water	stream	0.02		Tr	0.57	83.5	0.34	Tr
Skarvsnes	Oyako	4 m	0.36	NA	NA	NA	NA	NA	NA
Skarvsnes	Naga	5 m	0.88			Tr	100		91.26
Skarvsnes	Naga	pore	0.88	NA	NA	NA	NA	NA	NA
Skarvsnes	Nyorai	1.5 m	0.33			Tr	99.9		102.01
Skarvsnes	Nyorai	pore	0.33	NA	NA	NA	NA	NA	NA
Skarvsnes	Hotoke	1 m	0.42			Tr	99.4		87.62
Skarvsnes	Hotoke	pore	1.02	NA	NA	NA	NA	NA	NA
Skarvsnes	Kuwai	2 m	1.61			0.16	98.0		69.13

Table S2. Continued.

Area name	Lake/stream name	Depth	Salinity	NO <sub>2</sub> <sup>-</sup> μmol/L	NO <sub>3</sub> <sup>-</sup> μmol/L	NH <sub>4</sub> <sup>+</sup> μmol/L	%DON	PO <sub>4</sub> <sup>2-</sup> μmol/L	SiO <sub>3</sub> <sup>2-</sup> μmol/L
Skarvsnes	Bosatsu	2 m	0.18			0.17	99.7		61.17
Skarvsnes	Jizo	1.5 m	1.23			Tr	99.9		116.50
Skarvsnes	Himago	shore	3.90			0.15	99.6	Tr	21.46
Skarvsnes	Mago	shore	2.23			0.13	99.7		19.81
Skarvsnes	Sara	1.5 m	0.37			Tr	99.8		93.61
Skarvsnes	Tokkuri	shore	0.87			0.18	99.6		47.84
Skarvsnes	Kumogata	shore	0.65			0.60	98.2		74.74
Skarvsnes	Hyotan	6 m	1.07			Tr	100		11.55
Skarvsnes	Hamagiku	2 m	0.06			0.17	99.1		63.04
Skarvsnes	Ebi	3 m	0.84		Tr	Tr	99.4		71.51
Skarvsnes	Kikuno	shore	1.23			Tr	99.8		123.48
Skarvsnes	Hashioki	shore	0.27			Tr	99.8		171.60
Skarvsnes	Skarvsnes Lake2	shore	0.96		Tr	Tr	99.7		66.56
Skarvsnes	A-5	shore	0.39		0.57	0.78	96.5		207.49
Skarvsnes	Hechima	shore	1.67		Tr	Tr	98.9		169.75
Skarvsnes	Magobachi	shore	1.02		0.80	Tr	95.7		95.15
Skarvsnes	Misumi	shore	0.09		Tr	Tr	97.1		58.60
Skarvsnes	Kobachi	1.5 m	7.37			0.14	99.6	Tr	36.32
Skarvsnes	Kobachi	6 m	13.59			Tr	99.8	0.06	46.49
Skarvsnes	Suribachi	2 m	69.84			6.58	93.4		65.67
Skarvsnes	Suribachi	6 m	121.35			4.58	97.8	0.64	151.70
Skarvsnes	Funazoko	2 m	154.99			153.20	80.7		131.93
Skarvsnes	Funazoko	6 m	179.40		5.66	288.14	75.9	0.41	162.02
Langhovde	Yukidori 1	stream	0.02	0.57	12.18	0.22	28.2		6.16
Langhovde	Yukidori 2	stream	0.04	1.54	57.82	0.21	10.4	Tr	26.52
Langhovde	Yukidori 3	stream	0.04	1.00	57.53	0.18	11.4		19.62
Langhovde	Yukidori 4	stream	0.04	0.59	57.94	0.19	8.7	Tr	17.50
Langhovde	Yukidori 6	stream	0.04	Tr	3.06	0.15	72.9	Tr	23.70
Langhovde	Yukidori 7	stream	0.04		21.29	0.16	28.2	Tr	27.29
Langhovde	Yukidori 8	stream	0.05	Tr	26.31	0.16	27.4		42.12
Langhovde	Yukidori 9	stream	0.05	0.15	23.36	0.31	30.7		34.38
Langhovde	Yatsude 1	stream	0.05	0.13	5.96	0.14	40.8		50.30
Langhovde	Yatsude 3	stream	0.04	0.21	1.38		79.9		15.99
Langhovde	Yatsude 4	stream	0.05	0.25	16.62	0.18	22.9	Tr	17.79
Langhovde	Yatsude 5	stream	0.05	0.08	19.05	0.31	28.2	Tr	19.11
Langhovde	Yatsude 6	stream	0.05	Tr	44.26	0.14	21.5	0.07	19.37
Langhovde	Yatsude 7	stream	0.05	0.17	53.85	0.13	15.2	Tr	18.85
Langhovde	Yatsude 8	stream	0.04	0.08	40.22	Tr	18.5	Tr	11.95
Langhovde	Yatsude 9	stream	0.06	Tr	40.82	Tr	21.3		40.76

NA, not analyzed; Tr, below the quantification limit and thus values not shown

Figure S6: Distribution of FI, BIX, and HIX along salinity



**Fig. S6.** Distribution of fluorescence index (FI), biological index (BIX), and humification index (HIX) along salinity. Samples are color-coded according to the sampling category, and ‘penguin’ lake is a turbid, green-colored lake with abundant phytoplankton because of penguin guano inputs from nearby colonies (Amundsen Lake3, Fig. S1f). The interpretation of index values is presented on the right side of each figure along with the threshold lines to decide DOM origin and property (Cory et al., 2010; Cory and McKnight, 2005; Huguet et al., 2009; Ohno, 2002). FI and BIX values of saline lake samples were high. HIX values of stream samples and Amundsen Lake 3 were higher than those of others at the corresponding salinity.

Table S3: Assignments of the identified PARAFAC components

**Table S3.** Assignments of the identified PARAFAC components with the criterion of Tucker congruence coefficient ( $TCC_{ex \times em}$ )  $> 0.95$  using OpenFluor database. Red color indicates that an identified fluorescent component has  $TCC_{ex \times em}$  between 0.9 and 0.95, which is a fair threshold that takes into account variability that can arise from modeling and comparing two completely independent datasets (Lorenzo-Seva and ten Berge, 2006; Wünsch et al., 2017)..

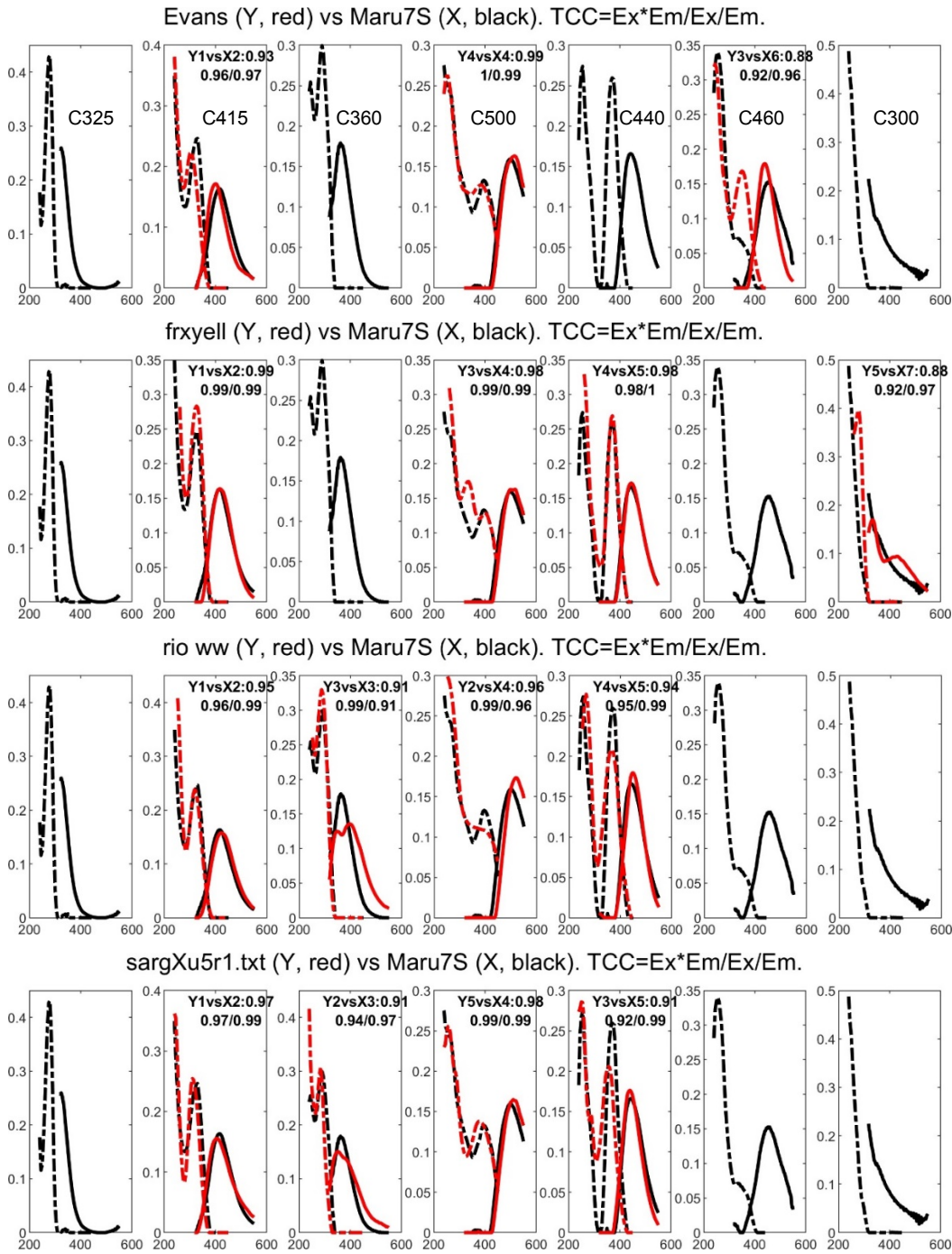
	Source	Data name	Comp No.	Description	References
<b>C1</b> <b>(C325)</b>	Autochthonous	IPY-C3O	C4	protein-like	(Guéguen et al., 2014)
		Kattegat	C6	tryptophan-like	(Osburn and Stedmon, 2011)
	Allochthonous	Not found			
	Ubiquitous	Not found			
<b>C2</b> <b>(C415)</b>	Autochthonous	TropicalRivers	C3	microbial humic-like	(Yamashita et al., 2010a)
		DarkOcean	C2	in situ microbial production of fluorescent humic-like materials	(Catalá et al., 2015)
	Allochthonous	DaecheongReservoir	C2	terrestrially derived humic substances	(Li et al., 2016)
		LiverpoolBay	C1	terrestrial humic-like	(Yamashita et al., 2011)
	Ubiquitous	Pantanal3_ww	C1	ubiquitous	(Murphy et al., 2018)
		Partners	C1	ubiquitous in aquatic environments	(Walker et al., 2013)
<b>C3</b> <b>(C360)</b>	Autochthonous	DW_Sweden_Gavle	C4	tryptophan-like	(Heibati et al., 2017)
		horsens2003	C5	combination of peaks N and T	(Stedmon et al., 2003)
	Allochthonous	Beringia	C4	not been reported previously/humic-like	(Walker et al., 2009)
	Ubiquitous	Not found			

**Table S3.** Continued

	Source	Data name	Comp No.	Description	References
<b>C4</b>	Autochthonous	Not found			
<b>(C500)</b>	Allochthonous	BalticIce	C2	terrestrially derived humic	(Stedmon et al., 2007)
		CoastalBEPOM	C2	terrestrial humic	(Brym et al., 2014)
	Ubiquitous	osPARAFAC_PonyLake	C6	one-sample decomposition of FDOM from Pony Lake	(Wünsch et al., 2017)
		Pantanal3_ww	C3	ubiquitous	(Murphy et al., 2018)
		Horsens2005	C2	fulvic acid fluorophore group. Present in all environments	(Stedmon and Markager, 2005a)
<b>C5</b>	Autochthonous	LiverpoolBay	C2	not defined by Coble et al/microbial humic-like	(Yamashita et al., 2011)
<b>(C440)</b>		TropicalRivers	C4	not defined by Coble et al/microbial humic-like	(Yamashita et al., 2010a)
	Allochthonous	Kattegat	C3	terrestrial humic A peak	(Osburn and Stedmon, 2011)
		Kauai	C8	Terrestrial humic substances A and C peak	(Murphy et al., 2008)
	Ubiquitous	GoMaine	C7	ubiquitous humic-like	(Cawley et al., 2012)
		Horsens2005	C4	fulvic acid fluorophore group. Present in all environments	(Stedmon and Markager, 2005a)
<b>C6<sup>†</sup></b>	Autochthonous	Bergen	C1	microbial processing of algae-derived DOM	(Stedmon and Markager, 2005b)
<b>(C460)</b>	Allochthonous	Horsens2005	C1	terrestrial humic fluorophore group, photodegradation restricted to UVB and UVC	(Stedmon and Markager, 2005a)
		FCE	C2	terrestrial humic-like, suggested as photo-refractory	(Yamashita et al., 2010b)
	Ubiquitous	Not found			
<b>C7</b>	Autochthonous	RecycleStM	G4	tyrosine-like	(Murphy et al., 2011)
<b>(C300)</b>		DarkOcean	C4	tyrosine-like	(Catalá et al., 2015)
		BalticIce	C5	tryptophan associated with proteins	(Stedmon et al., 2007)
	Allochthonous	Not found			
	Ubiquitous	Not found			

<sup>†</sup> A UVC humic-like component is often reported as accounting for the largest variability in the dataset, while C<sub>460</sub> had the second smallest explaining variability in this study. This may be because environments with a large contribution from terrestrial sources are often largely affected by a UVC humic-like component, while the Antarctic lakes and streams had little, if any, terrestrial inputs (such as aerial dusts).

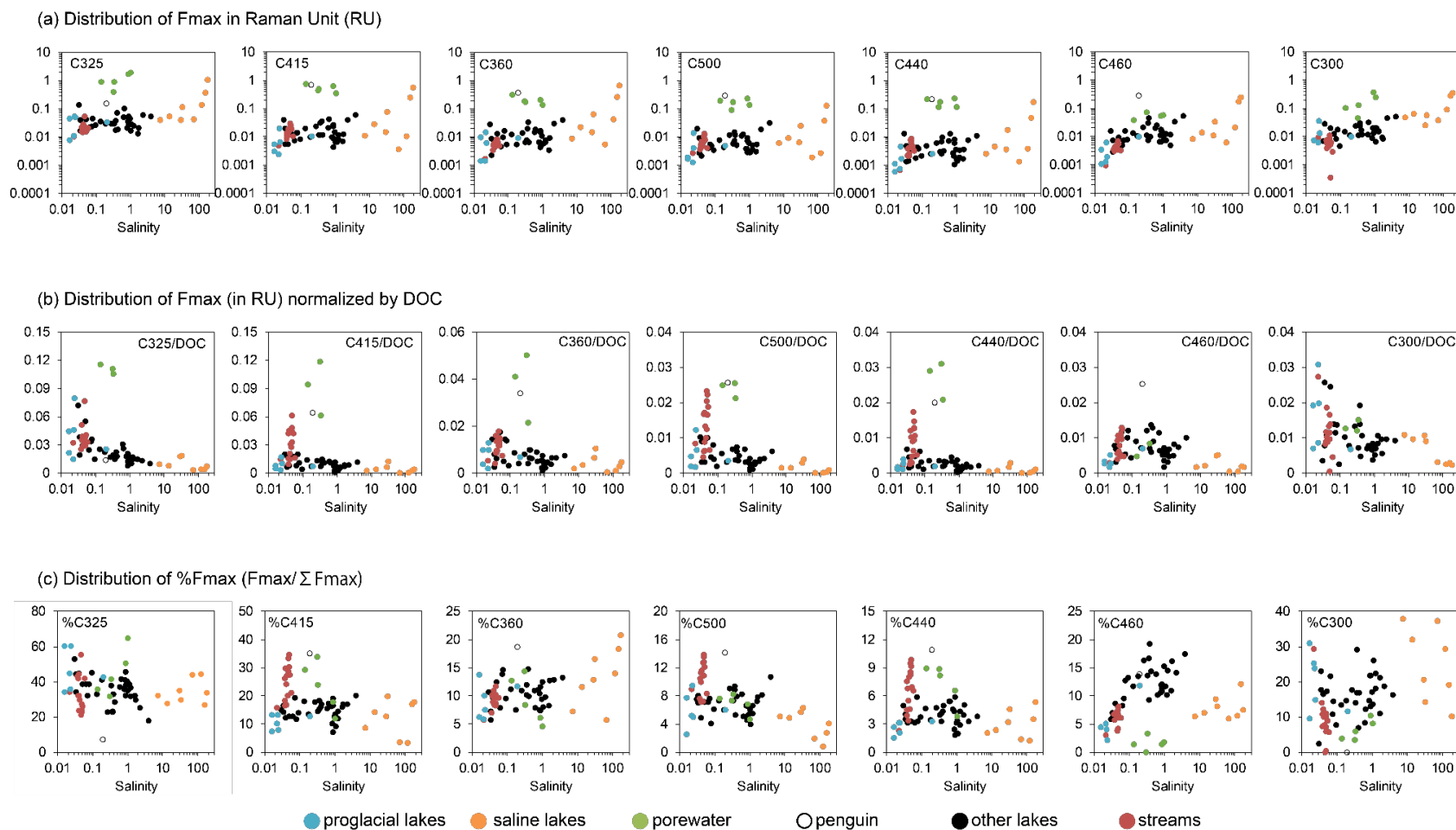
Figure S7: PARAFAC comparisons with ubiquitous models



**Fig. S7.** Comparisons of our PARAFAC model with PARAFAC models from the “ubiquity” paper (Murphy et al., 2018). Our 7-component model is shown in black, while 4- or 5-component ubiquitous models are in red. Pretty good matches are found for e.g. two Antarctic samples (coastal sample of Cape Evans and Lake Fryxell), Rio Negro (Brazil), and the deep Sargasso Sea (4534 m), especially for C<sub>415</sub>, C<sub>500</sub>, and C<sub>440</sub> which we considered as ubiquitous components.

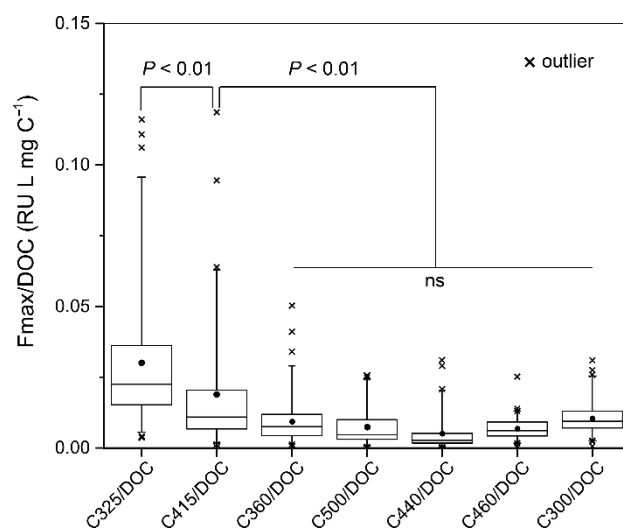


Figure S8: Distribution of the identified PARAFAC components along salinity



**Fig. S8.** Distribution of the identified PARAFAC components along salinity (log scale). See Fig. S6 for the sample color code interpretation.

Figure S9: Fmax/DOC of the identified PARAFAC components



**Fig. S9.** Box plot showing the distribution of Fmax/DOC of the identified PARAFAC components. The black solid circle and line represent the mean and median, respectively. The horizontal edges of the boxes denote the 25<sup>th</sup> and 75<sup>th</sup> percentiles, and the whiskers denote the 5<sup>th</sup> and 95<sup>th</sup> percentiles.

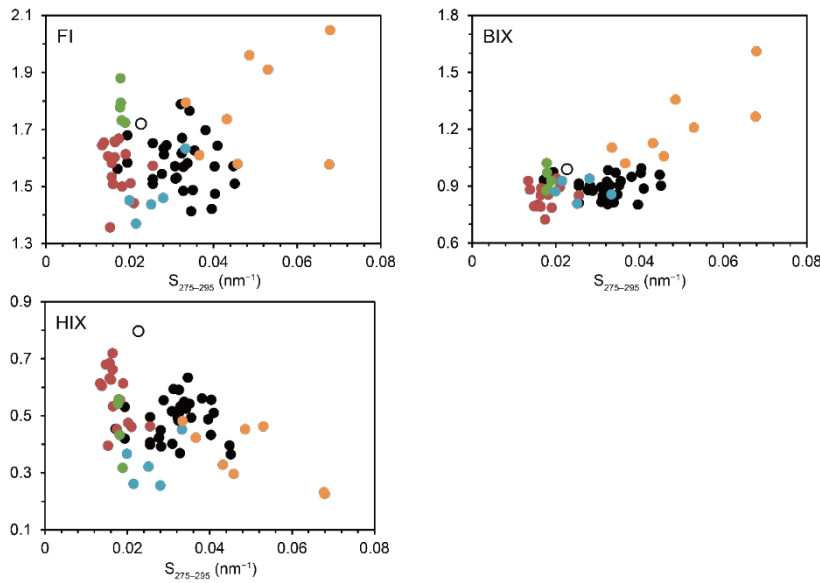
**Table S4.** Evaluation of salinity effects on CDOM and FDOM properties

Since our sample dataset encompassed large salinity differences ( $S = 0\text{--}179$ ), the observed differences in the optical properties may have been in part caused by the differences in salinity rather than DOM compositions between lakes. Thus, we evaluated salinity effects by changing the salinity of the solution containing the International Humic Substances Society (IHSS)'s reference sample Pony Lake fulvic acid (PLFA) at a constant pH of 8. Results showed that salinity effects are negligible on both CDOM ( $S_{275\text{--}295}$ ;  $< 3\%$ ) and FDOM indices (FI, BIX, and HIX;  $< 0.5\%$ ) (Table S4) that showed large variations along salinity in this study (Figs. 1 and S6). Prior research has shown that increases in the ionic strength increased the absorbance (mainly 300–500 nm) and decreased a spectral slope ( $S_{350\text{--}400}$ ) of reference DOM samples (Suwannee River humic and fulvic acids and PLFA) (Gao et al., 2015). However, the increase in the absorbance of PLFA was approximately one order of magnitude lower than that of the Suwannee River samples, and the influence of ionic strength variation was best explained with the changes of the protonation of the phenolic groups in the studied DOM (Gao et al., 2015). The non-sensitivity of PLFA to changes in salinity is consistent with the findings of Gao et al., and this indicates that effects of salinity were not quantifiable when comparing different salinity samples of predominantly autochthonous origin, and compositional change, rather than salinity, was the primary driver for variability in CDOM and FDOM properties. Similarly, the slight differences in pH ( $\sim 2$  unit) did not likely systematically influence our CDOM and FDOM dataset because there were no linear trends in the optical properties ( $S_{275\text{--}295}$ , FI, BIX, and HIX) along pH ( $P > 0.01$ ).

**Table S4.** Effects of salinity on CDOM and FDOM indices of Pony Lake fulvic acid (PLFA) at pH = 8. Salinity was changed by adding a desired amount of NaCl to PLFA solutions, and pH was adjusted by adding a small amount of NaOH or HCl solutions. Optical measurements were conducted after the 6-h equilibrium period.

Salinity	Optical index				Change in index values compared to $S = 0.188$ in percentage (%)			
	$S_{275\text{--}295}$	FI	HIX	BIX	$S_{275\text{--}295}$	FI	HIX	BIX
0.188	0.01436	1.638	0.8642	0.7722	-	-	-	-
0.452	0.01429	1.637	0.8657	0.7722	0.53	0.07	0.17	0.01
2.65	0.01434	1.635	0.8647	0.7743	0.15	0.17	0.06	0.27
25.0	0.01415	1.633	0.8683	0.7751	1.51	0.28	0.48	0.38
126	0.01396	1.630	0.8674	0.7752	2.82	0.47	0.36	0.39

Figure S10: Fluorescence-based indices versus  $S_{275-295}$



**Fig. S10.** Distribution of fluorescence index (FI), biological index (BIX), and humification index (HIX) along the spectral slope ( $S_{275-295}$ ). Color code according to Fig S6.

The fluorescence-based indices failed to adequately capture the changes in FDOM compositions as EEM-PARAFAC did. The fluorescence index (FI) did not show any trend along  $S_{275-295}$ , suggesting that the DOM compositional changes, neither photodegradation nor salinity, were responsible for the large variability in FI values (1.36–2.05, Figs. S6 and S10). There was a weak positive correlation between FI and %C<sub>360</sub> ( $R = 0.49$ ,  $P < 0.001$ ), but FI did not show the highest values in nutrient-rich saline lakes, implying that FI did not capture recent in-situ production as %C<sub>360</sub> did (Fig. 3b). The large variability in FI values of autochthonous DOM complicates its use as an indicator for DOM origin. The biological index (BIX)- $S_{275-295}$  plot showed a very similar distribution pattern as the BIX-salinity plot (Figs. S6 and S10). BIX was not sensitive in capturing the changes in FDOM compositions in water types other than saline lakes, and the higher values observed for saline lakes in Fig. S6 could be due to photodegradation (Fig. S10), and not due to a change in DOM origin. Thus, a frequently observed increasing BIX values during estuarine mixing, which has been interpreted as changes in DOM origins from terrestrial to recent in-situ production (Huguet et al., 2009), should be evaluated carefully. Murphy et al. (2018) also reported up to 37% increase in BIX values after photodegradation of DOM isolates. The humification index (HIX) showed an unclear trend along  $S_{275-295}$  (Fig. S10). HIX in this study had the very strong correlation with %C(ubiquitous humic + C<sub>460</sub>) ( $R^2 = 0.97$ ,  $P < 0.001$ ), meaning that HIX could not resolve distinct fluorophores ubiquitous humic-like components (C<sub>415</sub>, C<sub>500</sub>, and C<sub>440</sub>) and C<sub>460</sub>, which showed very different distribution patterns along  $S_{275-295}$  (Figs. 4c and 4d).

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