



Variability in dissolved organic matter optical properties in surface waters in the Amerasian Basin

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Surface absorption and fluorescence measurements of Dissolved Organic Matter (DOM) were conducted along with hydrographic parameters in the Amerasian and Makarov Basins. Parallel factor analysis of DOM fluorescence identified four humic-like and one protein-like component in all 107 surface samples. Based on strong negative trends observed between the spectral slope in the 275–295 nm range and absorption at 370 nm, and four humic-like components C1–4, the DOM character was found to be basin-dependent. The Makarov Basin surface DOM was largely dominated by high molecular weight and humic-rich material whereas the Canada Basin surface DOM was more heterogeneous with a marked influence of *in situ* production. This study highlights that absorbing and fluorescing measurements can be used successfully to trace and differentiate DOM from diverse sources and across frontal zones, and as such can be convenient and complementary tools for the better understanding of marine biogeochemical cycling of carbon.

Keywords: CDOM, FDOM, Makarov Basin, Canada Basin, frontal zone, ice cover

Introduction

Colored dissolved organic matter (CDOM) is one of the major light attenuating components of natural waters: it is responsible for much of the ultraviolet and visible light attenuation in the water column (Hansell and Carlson, 2002). CDOM also plays an important role in the solubility of essential elements such as iron and copper in natural waters (e.g., Chen et al., 2013; Heller et al., 2013; Uchida et al., 2013). The optical properties of CDOM have been previously used to discriminate between terrestrial and marine DOM sources (Blough and Del Vecchio, 2002; and references therein) and to trace the mixing of terrestrially derived DOM in oceanic waters (e.g., Amon et al., 2003; Guéguen et al., 2011; Stedmon et al., 2011; Fichot et al., 2013). For example, the absorbance properties of CDOM have been used to differentiate the surface to 300 m waters of the Canada and Eurasian Basins (Stedmon et al., 2011). More recently the application of fluorescence excitation emission matrices (EEMs) with a multivariate statistical analysis such as parallel factor analysis (PARAFAC; Stedmon and Bro, 2008) allowed us to discriminate the main fluorescing components in the fluorescing DOM (FDOM) pool and to assess their dynamics in marine waters (Ishii and Boyer, 2007; Stedmon et al., 2007; Murphy et al., 2008; Walker et al., 2009; Yamashita et al., 2010; Jørgensen et al., 2011; Guéguen et al., 2011, 2012; Dainard and Guéguen, 2013).

The Arctic Ocean basin represents ~4% of global ocean area and receives approximately 2.5 times more inflow per unit area than the world average. This disproportionate share of global

river discharge into the Arctic Ocean and the high dissolved organic carbon concentrations in arctic rivers (up to 1600 μM ; Stedmon et al., 2011; Amon et al., 2012) highlight the importance of freshwater and terrestrial organic matter in the biogeochemical cycles of the Arctic Ocean. However, there is a limited amount of data available on how this allochthonous DOM mixes with autochthonous marine DOM.

The general circulation in surface oceanic waters in the western Arctic Ocean is dominated by two main water masses: saline Atlantic water, entering through eastern Fram Strait and the Barents Sea, flows eastward following the Arctic continental margin where Eurasian river runoff is incorporated; and fresher Pacific-derived waters entering through Bering Strait and modified in the Chukchi Sea and Mackenzie River-influenced Beaufort shelves. The Pacific-Atlantic boundary in the Arctic Ocean delineates the extent of the Pacific-origin water, and thus where the saltier Atlantic-origin water must subduct beneath the Pacific Water. Eurasian river water has been reported in the Beaufort Gyre (Yamamoto-Kawai et al., 2009) and Pacific-origin water has recently been found in the surface layer of the North Pole (Alkire et al., 2007, 2010). Historically, this boundary lies along the Lomonosov Ridge, at the boundary between the Eurasian and Amerasian Basins of the Arctic Ocean (Jones and Anderson, 1986; Anderson et al., 1994), but recent studies have showed Atlantic character in the Makarov Basin that lies in the northern Amerasian Basin (i.e., Makarov and Canada Basins) between the Lomonosov Ridge and the Alpha-Mendeleev Ridge (Guay et al., 1999; Nishino et al., 2008, 2013; Guéguen et al., 2012). This part of the ocean is largely under-sampled which makes difficult to assess the presence of such a barrier that constrains transfer of heat, solutes, and plankton (Guéguen and Kowalczyk, 2013; and references therein).

Here we report spectral absorbance and fluorescence properties of surface water samples collected along transects from waters in the Amundsen Gulf across the Canada Basin to the Makarov Basin and back again. Our aim is to provide insight on sources and mixing of CDOM and FDOM in surface waters (i.e., terrestrial, marine and *in-situ* production) and thus improve our understanding of the biogeochemical cycling of carbon in the ocean.

Materials and Methods

Sample Collection

Surface water samples were collected from 107 sites located between the Canadian Arctic coast and the Lomonosov Ridge during August 21–September 27, 2011, on board the icebreaker Canadian Coast Guard Ship Louis S. St-Laurent (Figure 1) during the Natural Resources Canada expedition to assess sovereignty rights in the deep Arctic Ocean as required by the United Nations Convention on the Law of the Sea (UNCLOS). Samples were collected from the water intake line (depth ~ 10 m) every 6 h when steaming. *In-situ* measurements of temperature and conductivity were obtained by a Seabird electronics SBE-21 probe mounted on the sampling line inside the icebreaker. *In situ* fluorescence chlorophyll Chla (WETStar fluorometer) and FDOM (WETLabs ECOFLD 370 nm excitation, 460 nm

emission; Guéguen et al., 2012) sensors were also mounted on the intake line. *In-situ* measurements were conducted every 5 s and 1-min-averaged during post-processing. Samples collected for absorbance and fluorescence analysis were immediately stored in pre-combusted amber-glass vials with little to no headspace at 4°C until the analysis is performed. All samples were measured within 6 months of collection. The preservation and storage conditions used in this study may have favored microbial activities (Spencer and Coble, 2014; and reference therein). For example, Baker et al. (2007) reported a significant decrease (mean 58%) in protein-like fluorescence when samples were not filtered. The *in situ* FDOM sensor was linearly correlated with a_{370} , and humic like components C2 and C3 (described below) measured in discrete samples ($r^2 = 0.80, 0.66, \text{ and } 0.67$, respectively; $n = 107$). Additionally, the survey was augmented by expendable CTD (XCTD; Lockheed-Martin-Sippican) deployment.

Absorbance Analysis

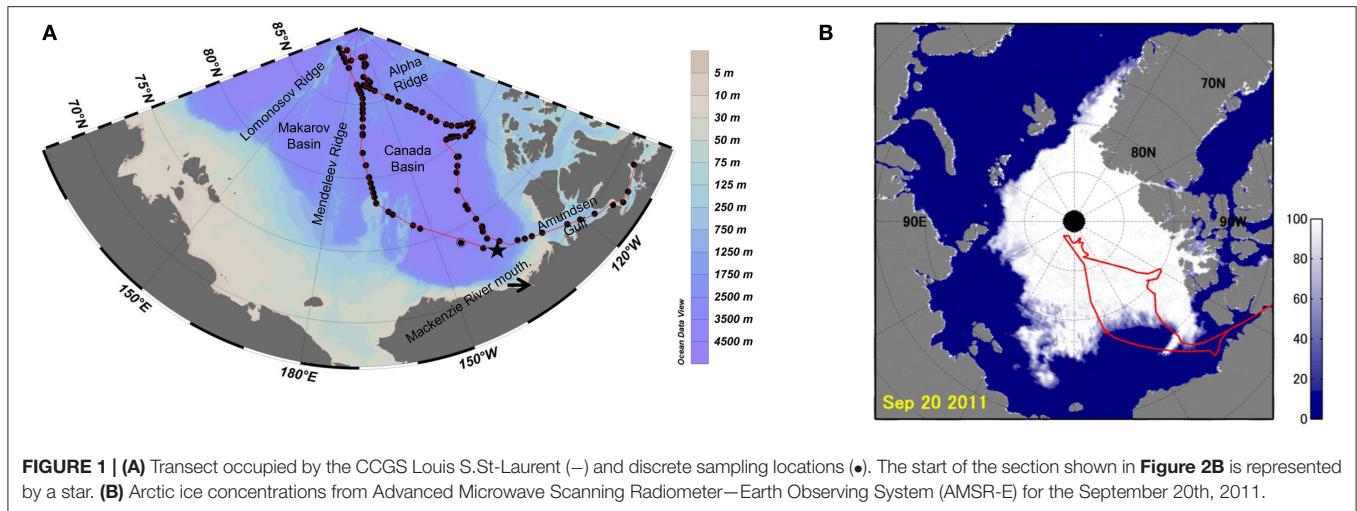
CDOM absorbance measurements of the collected water samples were made on a Shimadzu UV 2550 spectrophotometer over a wavelength region of 260–700 nm with a 10-cm quartz cuvette using Milli-Q water as a reference. In order to correct for instrumental drift and light scattering, an offset correction was applied by subtracting the average absorbance between 650 and 700 nm from the Milli-Q absorbance spectrum (Green and Blough, 1994; Guéguen et al., 2006, 2012). The measured absorbance at wavelength λ was converted to the absorption coefficient a (m^{-1}) according to the relationship:

$$a_{\lambda} = 2.303 \cdot A_{\lambda}/L,$$

where A_{λ} is the absorbance and L is the path-length of the optical cell in meters (here 0.1 m). The slope parameter S , a proxy for molecular weight (Helms et al., 2008), was calculated over a range of 275–295 nm ($S_{275-295}$; Helms et al., 2008) using non-linear least squares fitting procedures in Matlab. An increase in $S_{275-295}$ is indicative of decreasing aromaticity and molecular weight of the CDOM (Helms et al., 2008). S calculated over the 300–650 nm range ($S_{300-650}$; Stedmon and Markager, 2001; Retamal et al., 2007; Hancke et al., 2014) was used to compare previous mixing models (Stedmon and Markager, 2001; Hancke et al., 2014).

Fluorescence Analysis

Samples were warmed to room temperature and then analyzed with a Horiba Jobin-Yvon fluorometer (FluoroMax4) using a 1-cm quartz cuvette with excitation and emission slit widths each set to 5 nm. The quartz cuvette was checked for cleanliness between each sample at excitation wavelengths of 270 and 350 nm. The excitation-emission matrix (EEM) of each sample was recorded in signal-to-noise mode by collecting a series of emission wavelengths ranging from 250 to 600 nm at excitation wavelengths ranging from 300 to 450 nm, both in 5-nm increments. The post processing steps included: correction of instrument bias using the correction files provided by the manufacturer, subtraction of the EEM of Milli-Q water, and finally the fluorescence intensity was corrected to the area under the Milli-Q water Raman peak (excitation 350 nm) run



daily (Lawaetz and Stedmon, 2009). No significant variation in the integral of the Raman peak was observed during the investigation (1.2%). The fluorescence intensity was reported in equivalent water Raman units (r.u.). The Raman normalization is independent of the instrument design (Nelson and Coble, 2009; and references therein), allowing comparison with other studies.

PARAFAC decomposes the complex mixtures of FDOM fluorophores into their main components (Stedmon et al., 2003). PARAFAC analysis was carried out in Matlab using the DOMFluor toolbox (Stedmon and Bro, 2008), and outlier identification was performed using the OutlierTest function included therein. No samples with extreme leverage were found, indicating no extreme, and potentially outlying, EEMs in the dataset. Determination of the most suitable number of components was achieved by the split-half analysis and random initialization (Stedmon et al., 2003) whereby both halves were successfully validated. The model was constrained to non-negative values, and no systematic residual was found in the modeled EEMs. The spectra of the cross validated PARAFAC components were compared with those reported in earlier studies through online repository of published PARAFAC components (Murphy et al., 2014). The similarity of components was statistically identified as having Tucker congruence exceeding 0.95 (Murphy et al., 2014). The percent contribution of a given component was calculated as the ratio of the given component fluorescence intensity to the total humic component fluorescence intensity [i.e., $C1\% = C1/(C1 + C2 + C3 + C4)$].

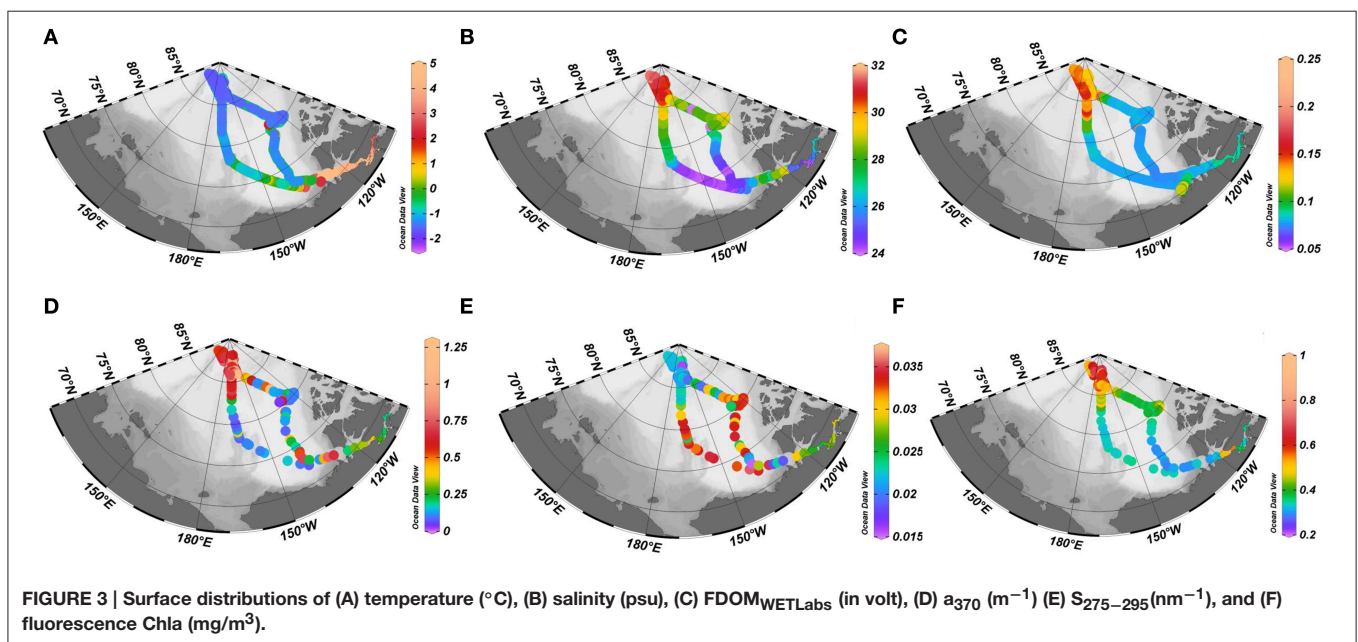
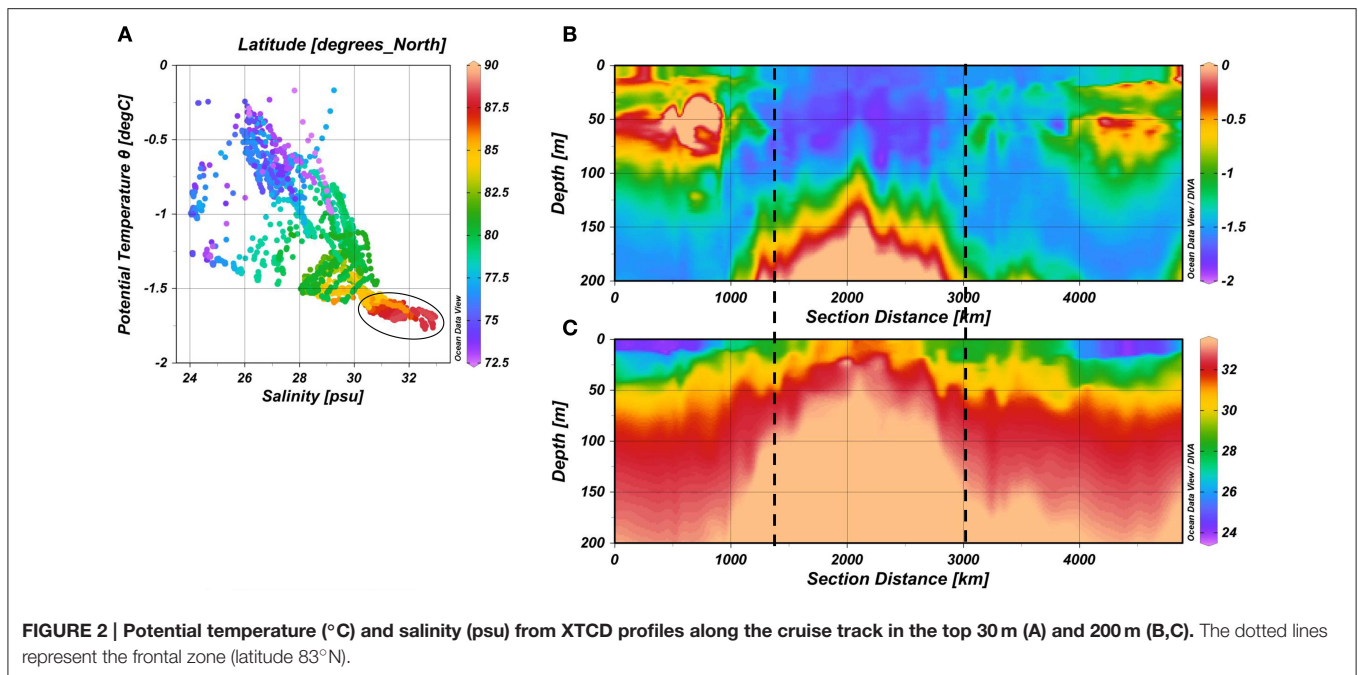
Results and Discussion

Salinity, Temperature, *in situ* FDOM, and CDOM Distributions

A marked frontal zone characterized by significant changes in physical and chemical characteristics was observed at latitude around 83°N (**Figures 2, 3**), coinciding with the latitude of the Alpha-Mendeleev Ridge that separates the Makarov and Canada Basins. Intrusion of warm Pacific Summer Waters around 50 m

and halo-stad around 50–125 m due to Pacific Summer and Winter Waters were found in the southern part of the UNCLOS area (<83°N; **Figure 2**). In contrast, higher latitude waters (>83°N) were ice covered ($98.9 \pm 1.8\%$; **Figure 1A**), colder (<-1.5°C) and saltier (>30; **Figure 2**), indicative of a stronger influence of Atlantic-origin water. The front at 83°N seems to be related to boundary of Atlantic and Pacific waters.

Surface distribution of salinity, temperature, FDOM_{WETLabs}, a_{370} , $S_{275-295}$ and Chla (**Figure 3**) showed contrasting characteristics across the frontal zone. The higher salinity (30.8 ± 0.3) observed north of 83°N suggest source waters of Atlantic origin while the high FDOM_{WETLabs} signal (0.14 ± 0.01 V), high a_{370} values (0.70 ± 0.15 m⁻¹) and low $S_{275-295}$ (0.0215 ± 0.001 nm⁻¹) reflect contribution from Eurasian Arctic rivers (Guay et al., 1999). Indeed, river influenced waters are typically enriched in larger molecular weight (low $S_{275-295}$; Helms et al., 2008) and humic-rich DOM (higher a_{370}). The near freezing temperature ($-0.75 \pm 0.53^\circ\text{C}$) north of 83°N reflects the presence of sea ice (**Figure 1B**). The >83°N region was also characterized by high Chla levels (>0.5 mg/m³). Nishino et al. (2013) have reported higher Chla in the Makarov Basin than in the Canadian Basin, likely due to sea ice retreat and shallower nutricline. On the other hand, the lower salinities were observed south of 83°N in the Canada Basin and the lowest salinities (as low as 24) were found in the central Canada Basin. These low salinities reflecting the contribution of sea ice melt and river inflow (Carmack, 1990; Yamamoto-Kawai et al., 2009). The highest $S_{275-295}$ values ($0.025-0.034$ nm⁻¹) coupled with lower FDOM_{WETLabs} (~ 0.8 V) and a_{370} values ($0.07-0.17$ m⁻¹) were found in the central Canada Basin (latitudes 72–83°N). The presence of lower molecular weight (high $S_{275-295}$) and less humified DOM is expected to primarily be the result of dilution of allochthonous CDOM (Granskog, 2012). Similar $S_{275-295}$ ranges were previously measured in the Arctic Ocean (Dainard and Guéguen, 2013; Fichot et al., 2013). The warmer (>0°C) and saltier waters (Salinity ~ 28) encountered in Amundsen Gulf were also characterized by low FDOM_{WETLabs} values, medium to high a_{370} values and low $S_{275-295}$. These values are indicative

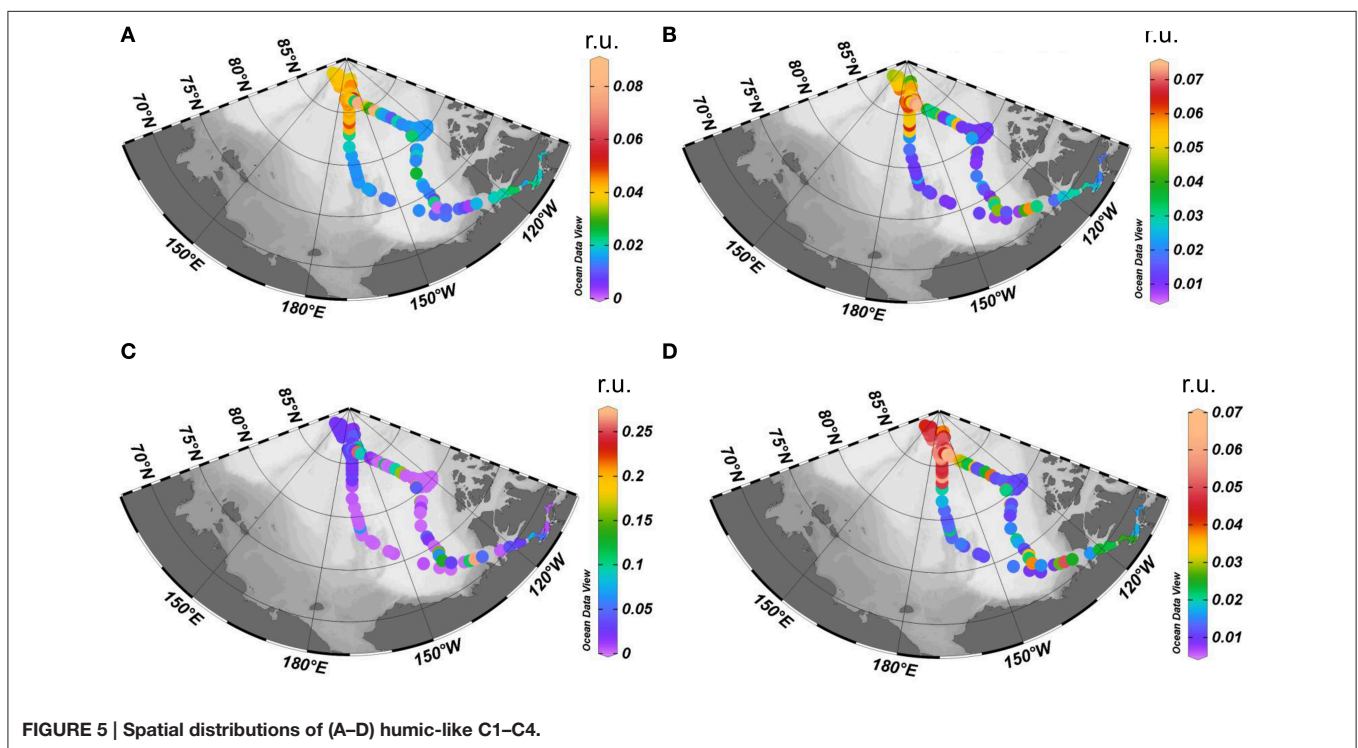
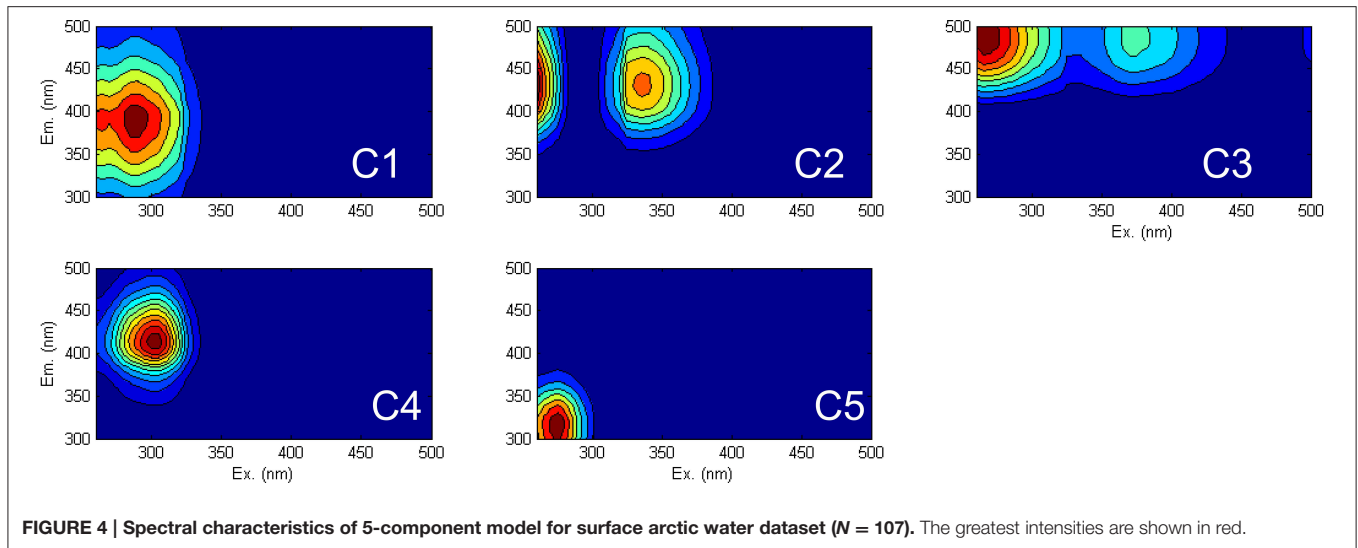


of river influence, but this will not be the Mackenzie River in 2011 as winds were easterly for the entire summer and fall which pushes the Mackenzie River water toward the west and offshore into the central Canada Basin (Yamamoto-Kawai et al., 2010).

FDOM Characterization and Distribution

Based on the PARAFAC modeling of EEM, five independent fluorescent components (C1–C5) were successfully validated. Their spectral characteristics were similar to components identified in previous studies (Figure 4; Figure S1). Components C1–C4 were characterized as humic-like, as they displayed an

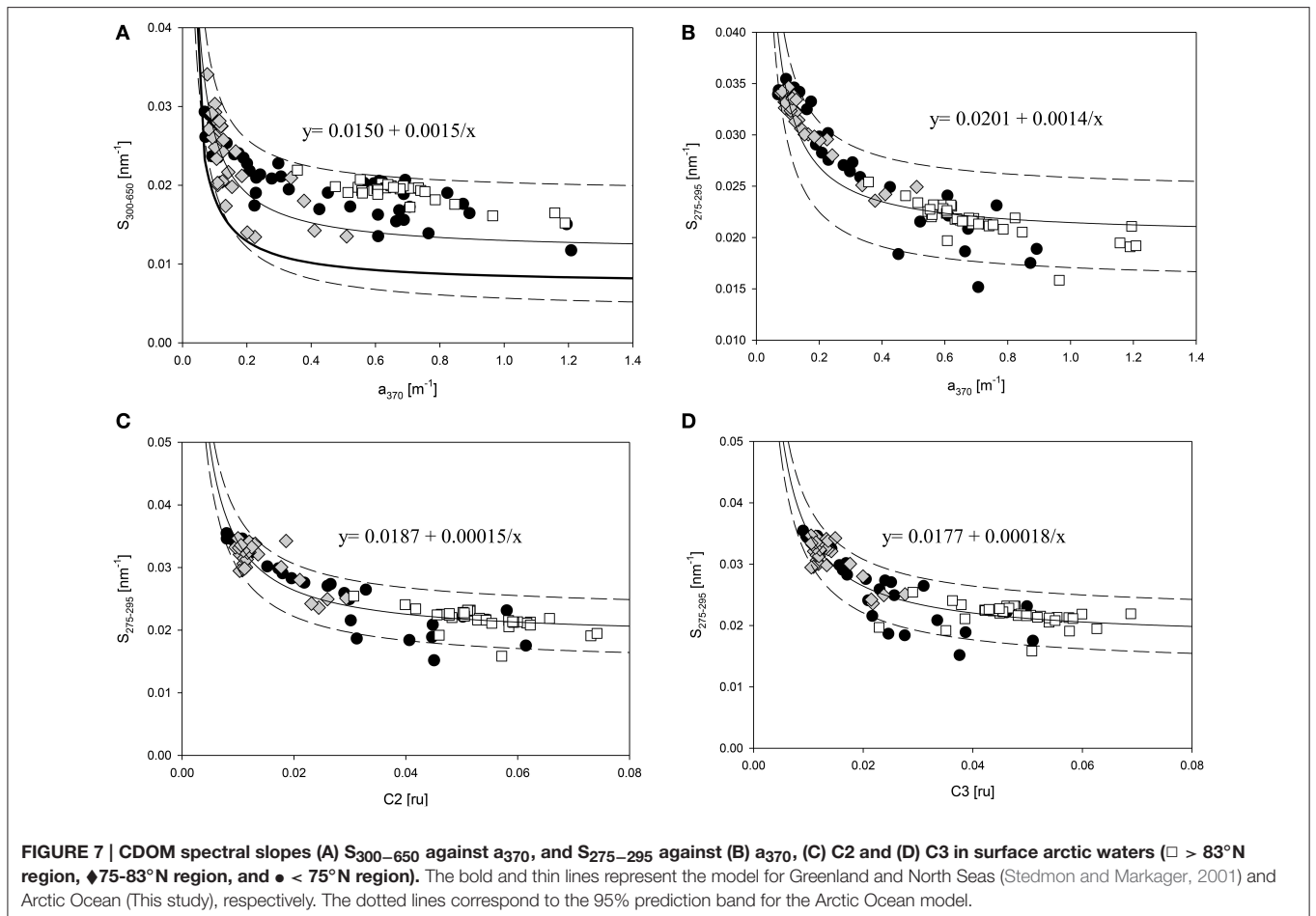
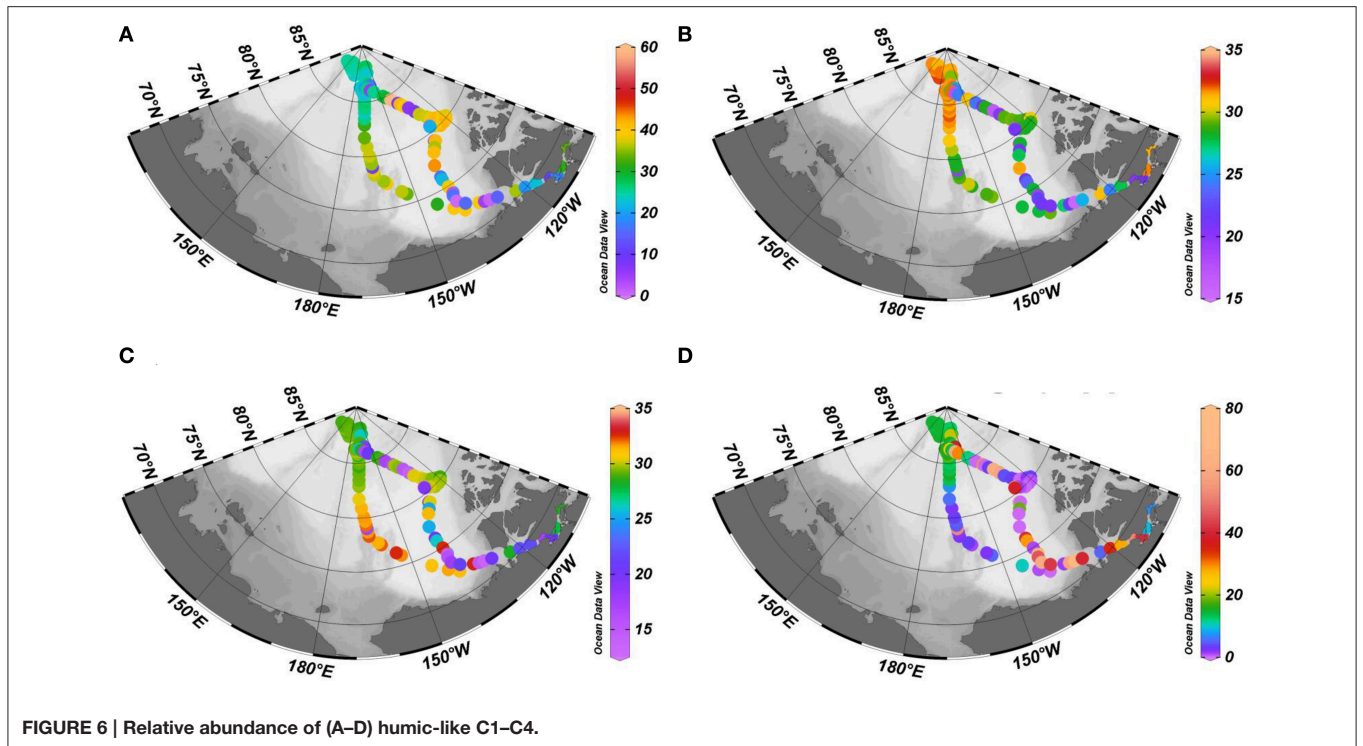
emission maximum around and above 400 nm. Components C1 and C4 showed an intensive peak at 285–305 nm excitation and 390–415 nm emission which was similar to the marine humic-like peak M traditionally defined (Coble, 1996). In recent studies, peak M has also been shown to be evident in DOM that has been altered by microbial reprocessing (Stedmon and Markager, 2005; Yamashita et al., 2008). Similar components were found in coastal and marine waters (Søndergaard et al., 2003; Yamashita et al., 2011; Dainard et al., 2015; Figure S1). The spectral features of C2 were similar to a component reported as humic-like peak A + M (Coble, 1996). Component C3 was assigned as the traditional



humic-like peak A + C (Coble, 1996) which is usually considered to be representative of fluorophores in terrestrial environments (Ishii and Boyer, 2007) and found in ocean waters (Søndergaard et al., 2003; Stedmon and Markager, 2005; Murphy et al., 2006, 2008; Stedmon et al., 2007; Walker et al., 2009; Kowalczyk et al., 2013; Brym et al., 2014). Component C5 was classified as protein-like since it displayed an emission maximum below 400 nm. This component was similar to those of a tyrosine-like component found in previous PARAFAC studies (Stedmon et al., 2007; Walker et al., 2009; Osburn and Stedmon, 2011; Cawley et al., 2012; Yamashita et al., 2013; Bianchi et al., 2014; Dainard

et al., 2015; Figure S1). As the samples were unfiltered, microbes could have altered the protein-like fluorescence via consumption of the original labile DOM and/or production of new protein-like fluorescence. Therefore, only results from the humic like components (C1–C4) were used for further data analysis in this study.

Strong humic-like C1–C4 signals were associated with the $>83^{\circ}\text{N}$ region (Figure 5). In terms of compositional distribution, microbially derived humic-like C1 was more abundant at intermediate latitudes in ice-edge (Figure 1B) and productive ($\text{Chla} \sim 0.4 \text{ mg}\cdot\text{m}^{-3}$; Figure 3F) waters ($75\text{--}83^{\circ}\text{N}$)



than in the $>83^{\circ}\text{N}$ region (32.4 ± 11.3 vs. $25.4 \pm 8.7\%$; **Figure 6A**). Greater abundance of microbially derived C1 agrees well with a more efficient microbial loop in the ice-edge bloom regions (Stanley et al., 2015). This contrasts with microbially derived C4 which was abundantly found in the $>83^{\circ}\text{N}$ region (15–20%; **Figure 6D**) and near the Mackenzie coast and in the Amundsen Gulf ($>40\%$; **Figure 6D**). Unlike the $>83^{\circ}\text{N}$ region, the coastal region (i.e., Mackenzie coast and Amundsen Gulf) was however not associated with high chl a (**Figure 3F**), suggesting that C4 was not always derived from freshly produced DOM. Humic-like C2–C3 (**Figures 6B,C**) showed higher abundances in the $>83^{\circ}\text{N}$ region ($\sim 15\text{--}32\%$); and along the western transect ($\sim 20\text{--}32\%$).

Mixing Processes

In marine environment, autochthonous CDOM can be discriminated from allochthonous CDOM by the relationship between spectral slope parameter and absorption coefficient (Stedmon and Markager, 2001). In our study, a significant inverse relationship was apparent between $S_{300-650}$ and a_{370} (**Figure 7A**), congruent with previous studies in the Arctic Ocean ($>100\text{ m}$ of depth; Stedmon et al., 2011), the Fram Strait (Granskog et al., 2012), and the Greenland and North Seas (Stedmon and Markager, 2001; Hancke et al., 2014). This contrasts with coastal waters (i.e., high terrestrial input) where S was shown to be independent of CDOM absorption coefficient (Babin et al., 2003). The majority of the points from $>83^{\circ}\text{N}$ waters had $S_{300-650}$ values around 0.019 nm^{-1} and deviated from the Stedmon and Markager (2001) model (thick line), likely the result of differences in DOM characteristics between Greenland/North Seas and western Arctic Ocean. $S_{275-295}$ was also used instead of $S_{300-650}$ as a better precision can be obtained in oceanic waters over the wavelength range 275–295 nm (Helms et al., 2008). A stronger relationship of $S_{275-295}$ vs. a_{370} was obtained compared to $S_{300-650}$ vs. a_{370} ($r^2 = 0.85$ vs. 0.65 , $n = 107$). However, this study, for the first time, reports a strong relationship between $S_{275-295}$ and FDOM (i.e., C2, $r^2 = 0.86$; C3, $r^2 = 0.85$; **Figures 7C,D**), suggesting mixing processes affecting C2 and C3 intensities in surface water.

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The behavior of CDOM and humic-like C2 and C3 was successfully modeled. Using **Figure 7**, one can clearly identify two different CDOM and FDOM pools: DOM from $>83^{\circ}\text{N}$, with an invariant S -value pool ($S_{275-295} = 0.0216 \pm 0.0017\text{ nm}^{-1}$), mixing with central Canada Basin waters ($75\text{--}83^{\circ}\text{N}$) with mean $S_{275-295}$ of $0.0312 \pm 0.0031\text{ nm}^{-1}$. Together these results show that the surface waters in the Canada and Makarov Basins differ significantly in composition.

Summary

This study shows that the combination of absorbance and fluorescence spectroscopies and multivariate statistics (i.e., PARAFAC) can be used to examine the source and mixing processes of DOM in the Canada and Makarov Basins. The presence of a frontal zone at latitude $\sim 83^{\circ}\text{N}$ was evidenced by marked changes in $S_{275-295}$ and a_{370} values and intensity and contribution of four humic-like PARAFAC components. High molecular weight (i.e., low $S_{275-295}$) DOM, dominated by terrestrially derived humic-like material, in the Makarov Basin was replaced by lower molecular weight DOM, with a reduced terrestrial character, in the Canada Basin. The CDOM and FDOM pools were concluded to be the result of mixing of two dominant sources (i.e., Eurasian and Central Canada Basins).

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Supplementary Material

The Supplementary Material for this article can be found online at: <http://journal.frontiersin.org/article/10.3389/fmars.2015.00078>

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Conflict of Interest Statement: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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