

Characterizing the origins of dissolved organic carbon in coastal seawater using stable carbon isotope and light absorption characteristics

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Abstract. In order to determine the origins of dissolved organic matter (DOM) occurring in the coastal seawater of Sihwa Lake, a water body semi-enclosed by a dyke along the western coast of South Korea, we measured the stable carbon isotope ratios of dissolved organic carbon (DOC- $\delta^{13}\text{C}$) and the optical properties (absorbance and fluorescence) of DOM in two different seasons (March 2017 and September 2018). The concentrations of DOC were generally higher in lower salinity waters in both periods, and excess of DOC was also observed in 2017 in high-salinity waters. The higher concentrations of DOC in the lower salinity waters originated mainly from marine sediments of tidal flats, based on the DOC- $\delta^{13}\text{C}$ values ($-20.7\pm 1.2\text{‰}$) and good correlations among the DOC, humic-like fluorescent DOM (FDOM_H), and NH_4^+ concentrations. However, the origins of the excess DOC observed in 2017 appears to be from two different sources: one mainly from marine sources such as biological production based on the DOC- $\delta^{13}\text{C}$ values (-19.1‰ to -20.5‰), and the other mainly from terrestrial sources by land-seawater interactions based on its depleted DOC- $\delta^{13}\text{C}$ values (-21.5‰ to -27.8‰). This land-derived DOM observed in 2017 was likely associated with DOM on the reclaimed land, which experienced extended exposure to light and bacterial degradation as indicated by the higher spectral slope ratio (S_R) of light absorbance and no concurrent increases in the FDOM_H and NH_4^+ concentrations. Our study demonstrates that the combination of these biogeochemical tools can be a powerful tracer of DOM sources in coastal environments.

1 Introduction

Dissolved organic carbon (DOC), a major component of dissolved organic matter (DOM), is the largest reduced carbon pool in the ocean (Benner et al., 1992; Raymond and Spencer, 2014). DOC plays a significant role in coastal carbon dynamics and biogeochemical cycles and thus it is important to understand the sources and characteristics of DOC (Vetter et al., 2007; Carson and Hansell, 2015). In the coastal ocean, DOM sources include (1) in-situ biological production (Carlson and Hansell, 2015), (2) terrestrial organic matter such as soils and plant matters (Opsahl and Benner, 1997; Bauer and Bianchi, 2011), and (3) anthropogenic sources such as industrial and agricultural wastewaters (Tedetti et al., 2010; Griffith and Raymond, 2011).

A part of DOM is known as colored dissolved organic matter (CDOM), which is the light-absorbing fraction of organic matter (Coble, 2007; Kim and Kim, 2016; Kim and Kim, 2018). The major fraction of CDOM, which emits fluorescence after absorbing light energy, is referred to as fluorescent DOM (FDOM) (Coble, 1996; Coble, 2007; Kim and Kim, 2016). Rivers are known as the major source of the humic-like FDOM (FDOM_H) in the coastal ocean (Stedmon and Nelson, 2015; Kim and Kim, 2016), while aerobic microbial remineralization of sinking organic matter is the major source of FDOM_H in the deep sea (Jørgensen et al., 2011; Catalá et al., 2015; Kim and Kim, 2016). On the other hand, Kim and Kim (2016) reported that anaerobic production in the bottom sediment of the deep East Sea (Japan Sea) accounted for approximately 10% of the total production of FDOM_H in a deep water column. Recently, the anaerobic process in the bottom sediment of coastal oceans was suggested as an important source of FDOM_H (Kim and Kim, 2018).

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In order to decipher the sources of FDOM_H and DOC in coastal waters, various approaches including optical properties and stable carbon isotope ratios ($\delta^{13}\text{C}$) have been developed. The humic-like component and protein-like component of FDOM were identified based on an excitation-emission matrix (EEM) spectroscopy method combined with a parallel factor analysis (PARAFAC) model (Bro, 1997; Coble, 2007; Kim and Kim, 2016; Kim and Kim 2018). The molecular weight and photochemical processes of DOM were characterized by the absorption spectra and spectral parameters of CDOM. For example, the absorption spectral slope ratio (S_R) negatively correlates with the DOM molecular weight and increases upon irradiation (Helms et al., 2008; Hansen et al., 2016). The stable carbon isotopic composition of DOC (DOC- $\delta^{13}\text{C}$) has been used to differentiate terrestrial versus marine DOC (Gearing, 1988; Wang et al., 2004; Lee and Kim, 2018; Lee et al., 2020). In general, $\delta^{13}\text{C}$ values of terrestrial sources such as C₃ and C₄ plants range from -23‰ to -34‰ and -9‰ to -17‰, respectively, while those derived from marine phytoplankton were found to range from -18‰ to -22‰ (Gearing, 1998).

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In this study, we used DOC- $\delta^{13}\text{C}$, FDOM, and S_R values together to characterize different sources of DOM in Sihwa Lake, South Korea, one of the most dynamic coastal settings in terms of salinity changes, hypoxia, metal pollution, and eutrophication (Kim et al., 2009; Ra et al., 2011; Lee et al., 2014; Kim and Kim, 2014; Lee et al., 2017; Kim and Kim, 2018).

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2 Materials and methods

2.1 Study area and sampling

Sihwa Lake (126.6 °E; 37.3 °N) is located on the western coast of South Korea, with an area of 57 km² and average depth of 3.2 m (maximum depth: 18 m). This lake was originally constructed as a land reclamation project planned by the government to provide agricultural land and water for the nearby metropolitan area during the 1980s and 1990s (Bae et al.,

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2010) (Fig. 1). Freshwater runs through six small streams into the lake, and four waterways connect the lake to the Banwol industrial complex (Fig. 1). Because the lake experienced serious deterioration of its water quality owing to wastewater discharges from industrial complexes under limited water circulation, sluice gates were constructed and opened twice a day for water exchanges between the lake and the Yellow Sea since 2012. The dyke is currently used as a tidal power plant (Lee et al., 2017) (Fig. 1). The total volume of Sihwa Lake water is $\sim 3.3 \times 10^8 \text{ m}^3$. The water exchange rate is approximately $3.4 \times 10^8 \text{ m}^3 \text{ y}^{-1}$, enough to replace the entire reservoir in a year (Lee et al., 2003; Lee et al., 2014).

Water samples for the entire water column were collected on board a ship (~ 1 ton) in March 2017 and September 2018. In 2018, only surface water samples were collected at shallow stations (station numbers 1–6) as the water level of the reservoir was lower than in 2017. In order to investigate the effect of wastewater from the industrial complex, an additional sample was collected near the Banwol waterway (station B4) in 2018 (Fig. 1). The temperature and salinity were measured using a conductivity-temperature-depth (CTD) instrument (Ocean Seven 304, INDONAUT Srl).

The water samples were filtered through a pre-combusted ($450 \text{ }^\circ\text{C}$ for 5h) GF/F filter (pore size: $0.7 \text{ }\mu\text{m}$; Whatman). The samples for the DOC and DOC- $\delta^{13}\text{C}$ analyses were acidified with 6M HCl (to a final sample pH of ~ 2) to avoid any bacterial activity and were stored in pre-combusted glass ampoules (Kim and Kim, 2010). Samples for the FDOM analysis were stored in pre-combusted amber vials in a refrigerator at 4°C . Samples for the dissolved inorganic nutrient analysis were stored frozen in polypropylene conical tubes.

2.2 Chemical analyses

Inorganic nutrient concentrations were measured using a nutrient auto-analyzer (New QuAAtro39, SEAL analytical). The analytical uncertainties were $<5\%$ for the reference materials for NO_x (KANSO Co., LTD). The dissolved oxygen (DO) concentration was determined using Winkler's method (Carpenter, 1965). The DOC concentration was measured using a high-temperature catalytic oxidation (HTCO) method using a total organic carbon (TOC) analyzer (TOC-V_{CPH}, Shimadzu) (Kim and Kim, 2010). The accuracy of the DOC measurements was $\pm 2 \text{ }\mu\text{mol L}^{-1}$ for a certified reference material of deep seawater (DSR; $\sim 43 \text{ }\mu\text{mol L}^{-1}$ DOC; University of Miami) (Hansell, 2005). The DOC- $\delta^{13}\text{C}$ values were measured using an isotope ratio mass spectrometer (IRMS; Isoprime, Elementar) connected to a TOC analyzer (Vario TOC cube, Elementar) (Panetta et al., 2008; Troyer et al., 2010). The DOC- $\delta^{13}\text{C}$ values were in very good agreement, within $<5\%$, with the reported values of the reference materials: IAEA-CH6 sucrose ($\delta^{13}\text{C} = -10.45 \pm 0.03\text{‰}$), Suwannee River Fulvic Acid (SRFA; $\delta^{13}\text{C} = -27.6 \pm 0.12\text{‰}$; International Humic Substances Society), and DSR (University of Miami) as reported by Lang et al. (2007) ($\delta^{13}\text{C} = -21.7 \pm 0.3\text{‰}$) and Panetta et al. (2008) ($\delta^{13}\text{C} = -21.4 \pm 0.3\text{‰}$).

2.3 Optical measurements

Fluorescence and absorbance spectra of the samples were measured using a spectrophotometer (Aqualog, Horiba). For the FDOM analysis, the emission and excitation wavelength ranges were set from 240 to 600 nm and from 250 to 500 nm, respectively, with 3 nm scanning intervals. The PARAFAC analysis for the EEM data was conducted using the Solo software (Eigenvector, 108 Inc.) (Han et al., 2020). The Raman and Rayleigh scattering signals, inner-filter effect, and blank subtraction were corrected with Milli-Q water (18.2 MΩcm) using the Solo software (Bro, 1997; Zepp et al., 2004; Stedmon and Bro, 2008; Han et al., 2020). The PARAFAC results were validated by a split-half analysis and random initialization (Stedmon and Bro, 2008). The fluorescence intensities of FDOM were normalized with the Raman peak area of water and were presented in Raman Unit (RU) (Lawaetz and Stedmon, 2009).

The PARAFAC model characterized one marine humic-like, one protein-like, and two terrestrial humic-like fluorescent components in Sihwa Lake, which are consistent with those of a previous study (Kim and Kim, 2018) (Fig. S1). The spectral shapes of the fluorescent components were compared with those in other results in the OpenFluor database (<https://openfluor.lablicate.com>) (Murphy et al., 2014). All components (C1 to C4) were matched with the major components from 36, 39, 62, and 19 studies, respectively, with similarity scores of 95%. The spectral characteristics of component 1 (FDOM_C; Ex/Em = 342/427 nm) and component 3 (FDOM_A; Ex/Em = 381/493 nm) are known to be associated with terrestrial humic-like DOM (Coble 2007). Component 2 (FDOM_M; Ex/Em = 297/388 nm) is known to be associated with marine humic-like DOM originating from microbial remineralization (Coble, 2007; Jørgensen et al., 2011). Component 4 (FDOM_P; Ex/Em = 282/322 nm) is characterized as a protein-like (tryptophan-like) DOM, which originates mainly from biological production (Coble, 2007). In this study, FDOM_C was used as a representative of humic FDOM (FDOM_H) since all humic-like components showed similar patterns.

UV-visible absorption spectra of the samples were measured with a scanning wavelength range of 240–700 nm. The absorption coefficient was calculated using the following equation:

$$a_{\lambda} = 2.303A_{\lambda}/l \quad (1)$$

where a is the absorption coefficient (m^{-1}), A_{λ} is the absorbance, and l is the optical path length of the quartz cuvette (m). The S_R value was calculated as the ratio of the spectral slope of shorter wavelengths ($S_{275-295}$) to that of longer wavelengths ($S_{350-400}$) (Helms et al., 2008; Han et al., 2020). The spectral slope (S) was calculated using the following equation:

$$a_{\lambda} = a_{\lambda_{ref}} e^{-S(\lambda - \lambda_{ref})} \quad (2)$$

where a is the Napierian absorption coefficient (m^{-1}), λ is the wavelength, and λ_{ref} is the reference wavelength (Twardowski et al., 2004; Helms et al., 2008).

3 Results

125 In 2017, the vertical distribution of salinity indicated a well-mixed water column (Fig. 2). Similarly, DO and NH_4^+
concentrations were vertically uniform (Fig. 2). Horizontally, salinity increased from 28 at the innermost station to 32 at the
outermost station (Fig. 2). The concentrations of DO and NH_4^+ were in the ranges of 7–13 mg L^{-1} (average: $10.1 \pm 2.4 \text{ mg L}^{-1}$)
and 0.1–25 $\mu\text{mol L}^{-1}$ (average: $8.7 \pm 8.1 \mu\text{mol L}^{-1}$), respectively (Fig. 2). The concentration of DO gradually increased with an
increase in the salinity along the transect, while the NH_4^+ concentration decreased with an increase in the salinity (Fig. 2).
130 The NH_4^+ concentration showed the lowest values ($<1 \mu\text{mol L}^{-1}$) between station 10 and station 13 (Fig. 2).

In 2017, the DOC concentrations were in the range of 97–349 $\mu\text{mol L}^{-1}$ (average: $184 \pm 76 \mu\text{mol L}^{-1}$). The distribution of
DOC concentrations was quite different from the salinity and DO distributions, with the highest concentrations in the surface
water samples of stations 3, 4, 5, 6, 7, 8, 9 and the bottom water samples of stations 3, 4, and 5 (Fig. 2). The DOC- $\delta^{13}\text{C}$
135 values ranged from -19.2‰ to -27.8‰ (average: $-21.8 \pm 1.9\text{‰}$) (Fig. 2). The most depleted values were observed in the
surface water samples of stations 5, 6, 7, 9, and 10 (-22.6‰ to -27.8‰) (Fig. 2). The concentrations of FDOM_C (terrestrial
humic-like component 1), FDOM_A (terrestrial humic-like component 2), FDOM_M (marine humic-like component), and
 FDOM_P (protein-like component) were in the ranges of 1.6–4.1 RU (average: $2.3 \pm 0.8 \text{ RU}$), 0.6–1.8 RU (average: 1.1 ± 0.3
RU), 1.0–2.4 RU (average: $1.5 \pm 0.5 \text{ RU}$), and 1.6–6.1 RU (average: $2.8 \pm 1.0 \text{ RU}$), respectively (Fig. 2). The concentrations of
140 all FDOM components were generally higher in the upstream stations and decreased with salinity (Fig. 2). The FDOM_P
concentration was slightly higher in the bottom water of station 10 (Fig. 2). The S_R values were in the range of 0.70–1.76
(average: 1.21 ± 0.20), with higher values in the surface water samples of stations 6, 8, 9, and 10 (Fig. 2).

In 2018, salinity showed a wide range, between 18 and 30, compared with that in 2017 (Fig. 3). Low-salinity waters
145 (salinity: 18–27) were observed from the innermost station to station 9 (Fig. 3). The concentrations of DO and NH_4^+ were in
the ranges of 6–11 mg L^{-1} (average: $8.2 \pm 1.6 \text{ mg L}^{-1}$) and 0.4–25 $\mu\text{mol L}^{-1}$ (average: $13.1 \pm 7.9 \mu\text{mol L}^{-1}$), respectively (Fig. 3).
The NH_4^+ concentrations in the outermost stations were lower than the detection limit (Fig. 3).

In 2018, the concentrations of DOC were in the range of 101–195 $\mu\text{mol L}^{-1}$ (average: $130 \pm 32 \mu\text{mol L}^{-1}$). The DOC
150 concentrations gradually decreased with an increase in the salinity (Fig. 3). The DOC- $\delta^{13}\text{C}$ values ranged from -19.1‰ to
 -21.5‰ (average: $-20.0 \pm 0.6\text{‰}$) (Fig. 3). The concentrations of FDOM_C , FDOM_A , FDOM_M , and FDOM_P were in the ranges
of 1.4–5.1 RU (average: $1.9 \pm 0.9 \text{ RU}$), 1.3–4.1 RU (average: $1.8 \pm 0.7 \text{ RU}$), 1.4–4.9 RU (average: $2.1 \pm 0.9 \text{ RU}$), and 1.1–2.5
RU (average: $1.6 \pm 0.4 \text{ RU}$), respectively (Fig. 3). All FDOM_H concentrations were higher in 2018 than they were in 2017
(Fig. 3). The FDOM_P concentrations were generally higher in the surface water samples, with a slightly higher value at
155 station 12, where the salinity is slightly lower (Fig. 3). The S_R values were in the range of 0.72–1.08 (average: 0.87 ± 0.10)
(Fig. 3). The S_R values were relatively constant at all sampling stations (Fig. 3).

4 Discussion

In both sampling periods, low-salinity waters showed higher DOC, lower DO, higher NH_4^+ , and higher FDOM_H (FDOM_C) concentrations (Figs. 2 and 3). As such, the DOC and FDOM_H concentrations exhibited significant negative correlations against salinity, indicating low-salinity-associated inputs of both DOM components (Figs. 4a and 4b). However, the patterns of the DOC distributions were very different from those of the FDOM_H distributions, indicating the decoupling of the major sources for both DOM components (Figs. 2 and 3). In this region, anthropogenic source inputs were found to be insignificant since there were no anomalous increases in DOC and FDOM_H concentrations in 2018 at station B4, where the waterway connects to the Banwol industrial complex (Fig. 3).

In both periods, the concentrations of FDOM_H showed significant correlations against salinities and NH_4^+ concentrations, with two negative slopes against salinity and a single positive slope against NH_4^+ (Figs. 4 and 5). This correlation trend suggests that the main source of FDOM_H was dependent on the supply of NH_4^+ , rather than freshwater inputs (Figs. 4a and 5a). In this lake, Kim and Kim (2018) hypothesized that FDOM_H is produced by the anaerobic decomposition of organic matter in bottom sediments in the freshwater-seawater mixing zone, based on good correlations among the salinity, NH_4^+ , and FDOM_H concentrations as well as low Eh values in samples with higher FDOM_H concentrations, although possible terrestrial inputs could not be excluded. Such correlations could also be observed if terrestrial fresh water, which has higher FDOM_H and lower NH_4^+ levels, passes through a salinity mixing zone where extremely high amounts of NH_4^+ and low FDOM_H are introduced from bottom sediments. However, our $\text{DOC}-\delta^{13}\text{C}$ values ($-20.0\pm 0.4\text{‰}$) observed in low-salinity (<28) water samples here exclude any possible significant contributions of terrestrial sources, indicating the effective degradation of terrestrial sources before they reached the mixing zone. In this tidal flat environment, the large inputs (i.e., seepage) of DOC and FDOM_H , together with NH_4^+ , may have happened from marine sediments without significant decreases in the DO concentration (Kim et al., 2012), as the shore water runs back and forth over a wide sediment-surface area during tidal cycles (Fig. 5). In this case, depending on the salinity ranges of the overlying waters, different slopes of FDOM_H against salinities could be observed for different seasons as observed in this study (Fig. 4b).

In 2018, the distribution pattern of the DOC concentrations was similar to that of FDOM_H (Fig. 3). During this period, the freshwater contribution was greater compared with that in 2017, with the lowest salinity being approximately 18. However, the $\text{DOC}-\delta^{13}\text{C}$ values in all samples in this period ranged from -19.1‰ to -21.5‰ (average: $-20.0\pm 0.6\text{‰}$), falling into the range of marine phytoplankton values (-18‰ to -22‰) (Gearing, 1988) (Fig. 4c). The S_R values were relatively low and constant (average: 0.86 ± 0.1) at all stations in 2018, indicating no significant changes in the input sources or the effects of photochemical and bacterial degradation (Moran et al., 2000; Helms et al., 2008) (Fig. 4d). Thus, we conclude that the main source of DOC in 2018 was from marine sediments, together with FDOM_H and NH_4 , as external terrestrial sources of DOC were insignificant or effectively degraded on the tidal flat (Kim et al., 2012).

In 2017, the sources of DOC were more complicated, showing large excess DOC concentrations independent of salinity (Fig. 2). Here, this excess denotes concentrations higher than that of the incoming seawater (background) from the Yellow Sea (highest salinity and lowest DOC concentrations) observed in 2018 (Fig. 4a). The samples with excess DOC levels observed in 2017 were nearly 69% higher than the DOC values of the background seawater ($\sim 110 \mu\text{mol L}^{-1}$) (Fig. 4a). The samples with higher DOC levels observed in 2017 were separated into two major groups (Group 1 and Group 2) based on their DOC concentrations, DOC- $\delta^{13}\text{C}$ values, and salinities (Fig. 2). Group 1 ($n=3$) includes the samples observed in stations 12 and 13 (Fig. 2). Group 2 ($n=11$) includes the samples observed in the surface waters of stations 4, 5, 6, 7, 8, 9, and 10, and in the bottom waters of stations 3, 4, 5, and 6 (Fig. 2).

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For the Group 1 samples, the DOC concentrations ranged from 144 to 223 $\mu\text{mol L}^{-1}$ (average: $175 \pm 42 \mu\text{mol L}^{-1}$) (Figs. 2 and 4a). The DOC- $\delta^{13}\text{C}$ values ranged from -19.1‰ to -22.1‰ (average: $-20.6 \pm 1.5\text{‰}$), which are close to the $\delta^{13}\text{C}$ values of marine organisms (Figs. 2 and 4c). The S_R values (average: 1.07 ± 0.03) in this group were relatively low and constant compared with that of Group 2 (Figs. 2 and 4d). Thus, the higher DOC concentrations observed in this group appear to be from in-situ biological production.

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For the Group 2 samples, the DOC concentrations ranged from 103 to 330 $\mu\text{mol L}^{-1}$ (average: $213 \pm 69 \mu\text{mol L}^{-1}$) (Figs. 2 and 4a). The DOC- $\delta^{13}\text{C}$ values ranged from -21.5‰ to -27.8‰ , which include the signature of terrestrial C_3 plants (-23‰ to -32‰) (Gearing, 1988) (Figs. 2 and 4c). For this group, increases in FDOM_H concentrations were not observed, although the S_R values (average: 1.26 ± 0.2) were higher than those at other stations, likely due to the influence of highly degraded low-molecular-weight DOM (Helms et al., 2008) (Fig. 4d). Thus, this terrestrial DOM observed in 2017 seems to be associated with non-fluorescent and low-molecular-weight DOM on the reclaimed land by direct land-seawater interaction through the tidal inundation of seawater. This could happen if the terrestrial DOM experienced intense light exposure and/or bacterial degradation on land, a condition likely facilitated through the process of land reclamation.

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Without the use of these multiple tracers, in this region, the higher DOC and FDOM_H concentrations occurring in areas with lower salinity waters could be attributed to inputs with terrestrial origins, while the higher DOC concentrations in high salinity areas could be attributed to in-situ biological production. However, these multiple tracers can lead to the better identification of different source inputs. Therefore, our results suggest that the combination of stable carbon isotope, FDOM , and S_R values can be a critical tool for deciphering the sources and characteristics of DOM in coastal waters where various DOM sources are present.

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5 Conclusions

225 Different sources and distributions of DOM were determined in different seasons using various tracers in Sihwa Lake, South
Korea. Our results reveal that the higher DOC concentrations occurring in low-salinity water, which could be regarded as
having terrestrial sources, were actually from marine sediment sources based on $\text{DOC}-\delta^{13}\text{C}$ values (-19.1‰ to -21.5‰)
together with significant correlations among the DOC, FDOM_{H} , and NH_4^+ concentrations. On the other hand, the excess
DOC concentrations occurring in high-salinity waters at some stations, which are generally believed to be from marine
sources, were found to be from non-fluorescent, low-molecular-weight, terrestrial DOM sources from the reclaimed land
230 area based on depleted $\text{DOC}-\delta^{13}\text{C}$ values (-21.5‰ to -27.8‰) and high S_R values (1.26 ± 0.2), without concurrent increases
in the FDOM_{H} and NH_4^+ concentrations. Our study demonstrates that the combination of these multiple DOM tracers can be
used successfully in other coastal waters where the sources and characteristics of DOM are complicated.

Data availability. All data are available upon request to the corresponding author.

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Author contributions. GK contributed to the conceptualization of the manuscript. HH and GK were involved in planning the
research. HH collected samples and performed the analyses. All authors were involved in analyzing the results and writing
the paper.

240 *Competing interests.* The authors declare that they have no conflict of interest.

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350

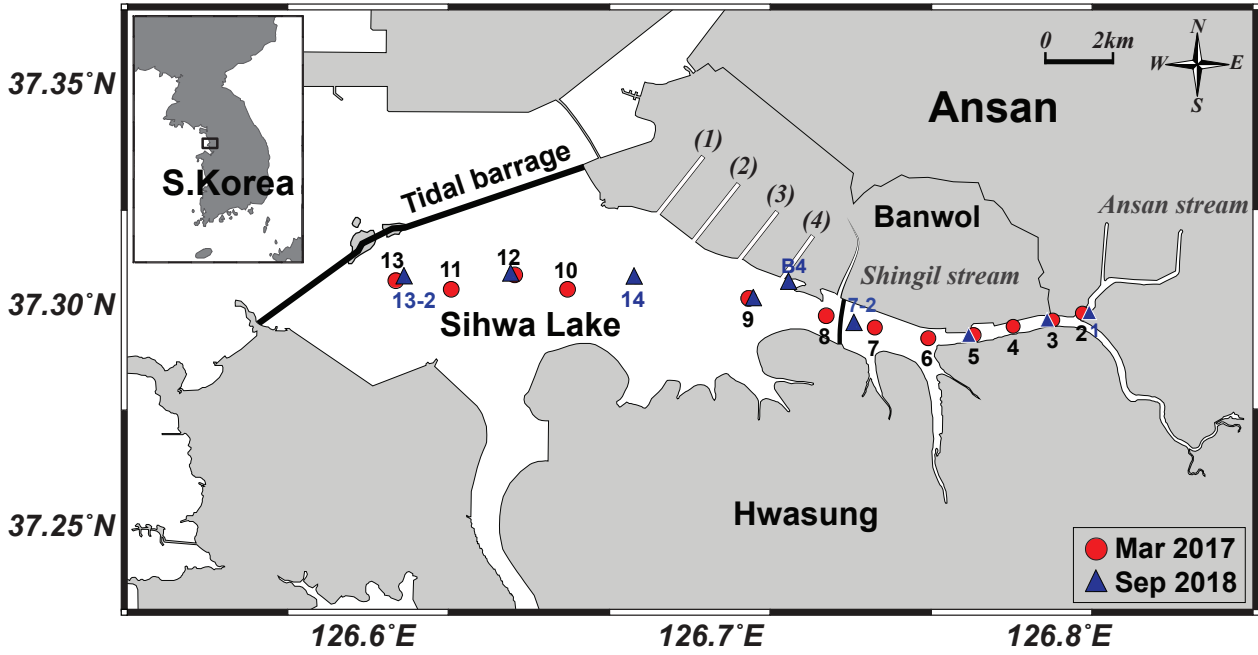


Figure 1: Map of sampling stations in Sihwa Lake, South Korea.

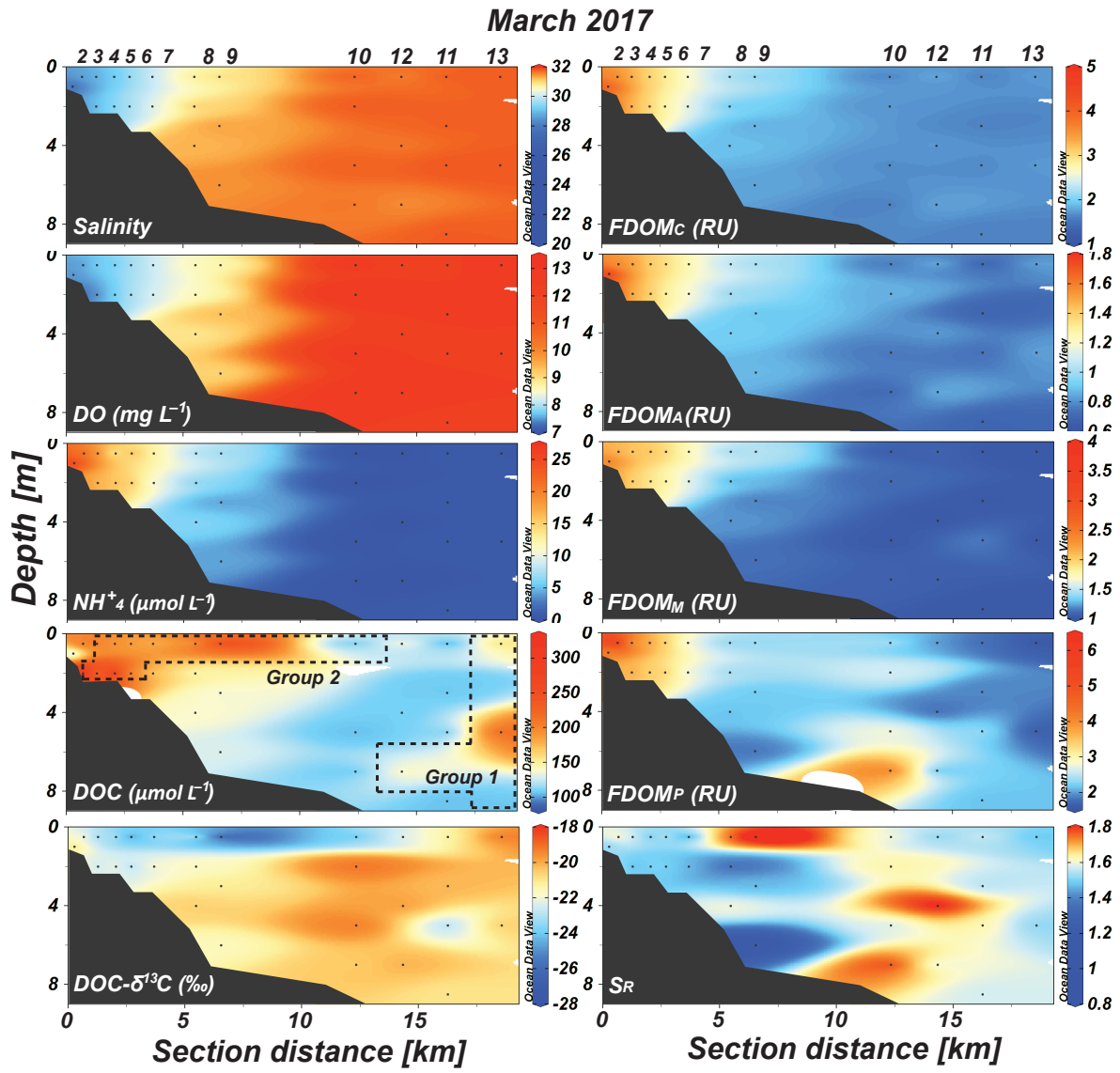


Figure 2: Vertical distributions of salinity, DO, NH_4^+ , DOC, $\text{DOC-}\delta^{13}\text{C}$, FDOM_C , FDOM_A , FDOM_M , FDOM_P , and S_R values in Sihwa Lake in March 2017. The dashed blocks represent stations belonging to the Group 1 and the Group 2, which were separated based on the distributions of DOC, $\text{DOC-}\delta^{13}\text{C}$, and salinity.

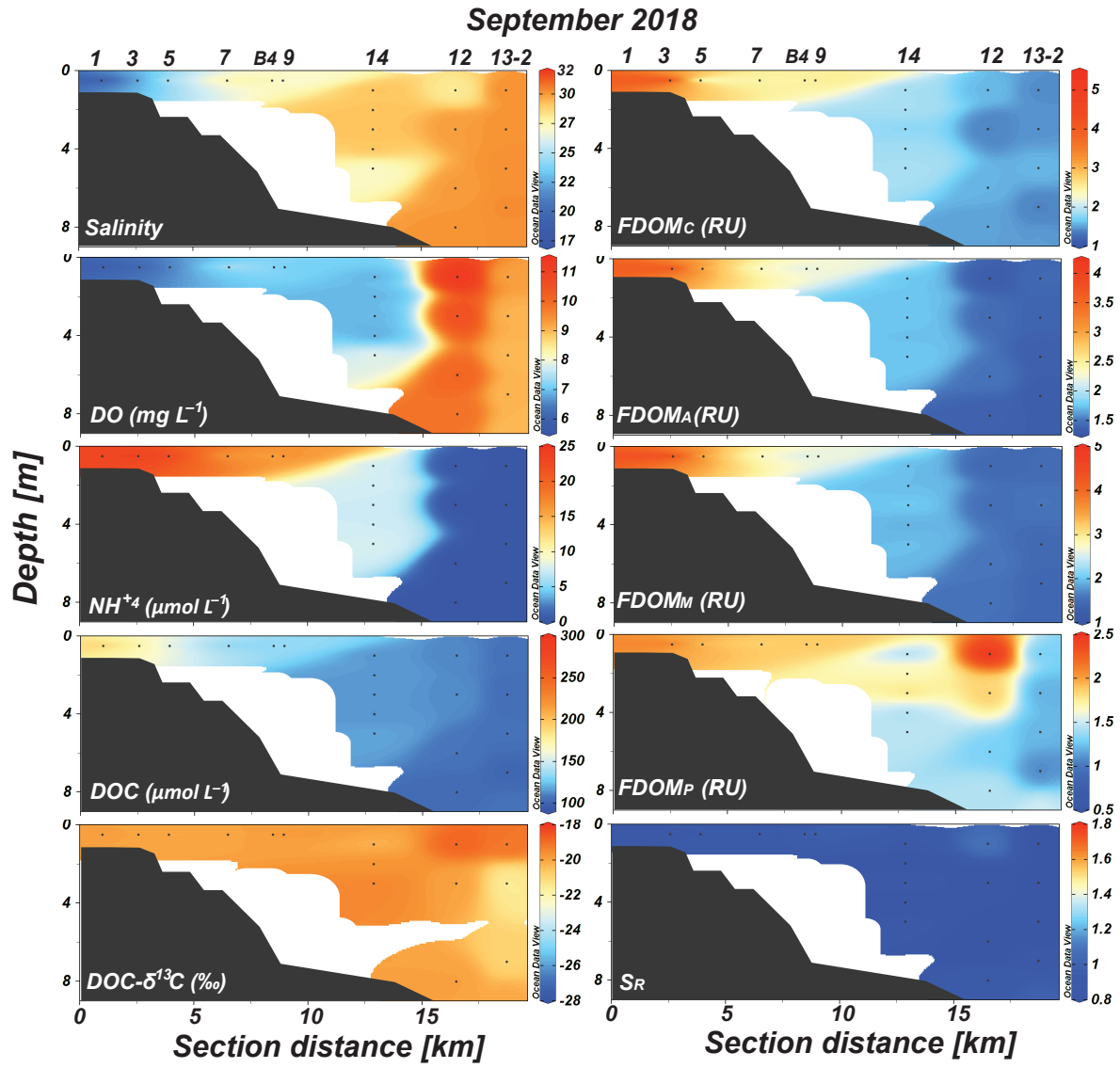
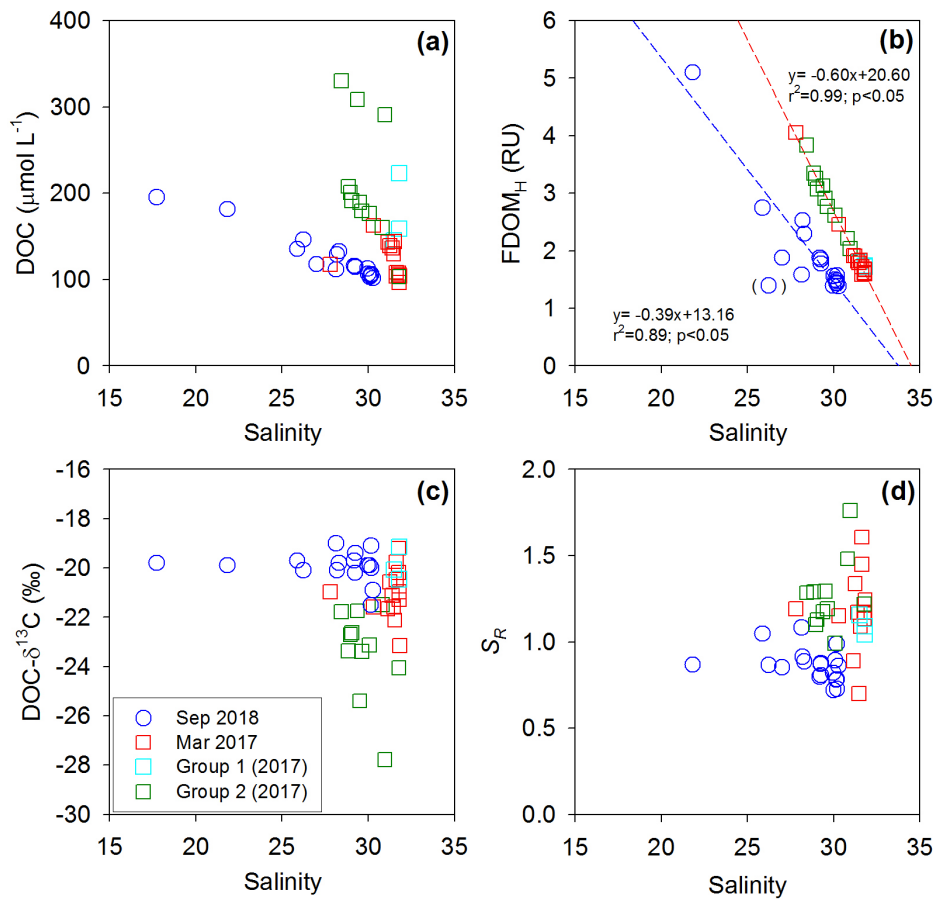


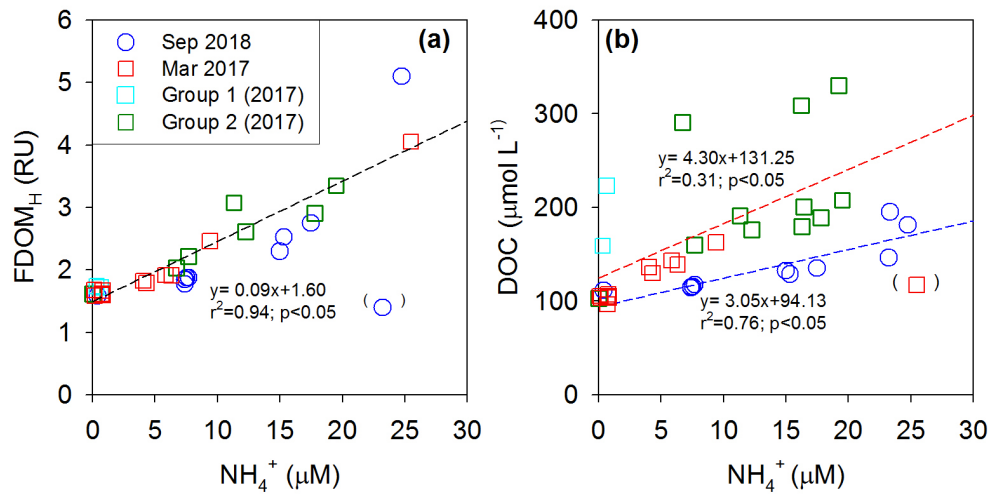
Figure 3: Vertical distributions of salinity, DO, NH_4^+ , DOC, $\text{DOC}-\delta^{13}\text{C}$, FDOM_C , FDOM_A , FDOM_M , FDOM_P , and S_R values in Sihwa Lake in September 2018.



385

Figure 4: Plots of salinity versus (a) DOC concentrations, (b) FDOM_H concentrations, (c) DOC- $\delta^{13}\text{C}$ values, and (d) S_R values in Sihwa Lake in March 2017 (red square) and September 2018 (blue circle). The dashed lines represent the regression lines.

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Figure 5: Plots of NH_4^+ concentrations versus (a) FDOM_H concentrations and (b) DOC concentrations in Sihwa Lake in March 2017 (red square) and September 2018 (blue circle). The dashed lines represent the regression lines.