



1 Distribution, seasonality, optical characteristics, and fluxes of dissolved organic

2 matter (DOM) in the Pearl River (Zhujiang) estuary, China

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17 Abstract

18 Dissolved organic carbon concentration in the Pearl River estuary (PRE) of China was measured in 19 May, August, and October 2015 and January 2016. Chromophoric and fluorescent dissolved organic matter (CDOM and FDOM) in the latter three seasons were characterized by absorption and 20 21 fluorescence spectroscopy. Parallel factor analysis of the fluorescence spectra identified two protein-22 like, two humic-like, and one oxidized quinone-like FDOM components. The seasonality of average DOM abundance varied as follows: DOC: May (156 μ mol L⁻¹) > January (114 μ mol L⁻¹) \approx August 23 (112 µmol L^{-1}) > November (86 µmol L^{-1}); CDOM absorption at 330 nm: August (1.76 m⁻¹) > 24 November $(1.39 \text{ m}^{-1}) \approx$ January (1.30 m^{-1}) ; FDOM expressed as the sum of the maximum fluorescence 25 intensities of all FDOM components: November (1.77 R.U.) > August (1.54 R.U.) ≈ January (1.49 26 27 R.U.). Average DOM abundance in surface water was higher than in bottom water, their difference 28 being marginal (0.1-10%) for DOC in all seasons and for CDOM and FDOM in November and 29 January, and moderate (16–21%) for CDOM and FDOM in August. DOC showed little cross-estuary 30 variations in all seasons while CDOM and FDOM in January were higher on the west side of the 31 estuary than in the middle and on the east side. All three variables exhibited large variations and/or 32 rapid drawdowns at the head of the estuary (salinity <5) due to multiple freshwater endmembers and/or 33 biotic losses. In the saltier zone, they declined linearly with salinity except relatively constant DOC in 34 May and November. The decrease in FDOM was 5–35% faster than that in CDOM, which in turn was 35 2–3 times faster than that in DOC. Salinity and CDOM absorption coefficients can serve as indicators of DOC in August and January. Absorbance- and fluorescence-based indices demonstrate that 36 37 freshwater endmembers in all seasons mainly contained fresh, protein-rich DOM of microbial origin, 38 though the proportion of humic-like components was somewhat higher in August. Protein-like materials were preferentially consumed in the low-salinity section but the dominance of the microbial 39 signature was maintained throughout the saltier zone. Exports of DOC and CDOM (in terms of a_{330}) 40





into the South China Sea were estimated as 195×10^9 g and 266×10^9 m² for the PRE, and 362×10^9 g and 493×10^9 m² for the entire Pearl River Delta. Compared to other world major estuaries, the PRE presents the lowest concentrations and export fluxes of DOC and CDOM. Nonetheless, DOM delivered by the PRE is protein-rich and thus may significantly impact the local ecosystem.

45

46 **1 Introduction**

47 **1.1 Overview of DOM**

48 Dissolved organic matter (DOM) in the ocean drives major biogeochemical cycles involving carbon, 49 nutrients, trace metals, and trace gases (Miller and Zepp, 1995; Cauwet, 2002; Wells, 2002; Cobble, 50 2007). The chromophoric component of DOM (CDOM), which absorbs solar ultraviolet (UV) and 51 visible radiation (Blough et al., 1993; Nelson et al., 1998; Siegel et al., 2002), affects ocean optics and 52 generates various photoreactions (Mopper and Kieber, 2002; Zafiriou, 2002; Zepp, 2003). The 53 biogeochemical and optical significance of DOM depends on both its abundance and quality (i.e. 54 chemical composition), with the latter strongly linked to its origin of formation (Repeta, 2015; Lønborg 55 et al., 2016, Massicotte et al., 2017).

Coastal waters, particularly those impacted by river plumes, contain higher contents of DOM 56 57 relative to the open ocean. DOM in river water originates from soil leaching (terrigenous DOM, or 58 tDOM) and in situ microbial production. tDOM, enriched with lignin phenols (Opsahl and Benner, 59 1997), differs substantially from microbial-derived DOM, enriched with proteins and aliphatic 60 hydrocarbons (Martínez-Pérez, et al., 2017; Brogi et al., 2018), in optical property and biological and 61 photochemical lability (Hansen et al., 2016; Sulzberger and Arey, 2016). Consequently, river runoff 62 can profoundly impact coastal ecosystem functioning by increasing the quantity and altering the quality 63 of DOM in coastal waters.





64 The loads of terrigenous and microbial DOM and their proportions in a river rely on many factors, 65 among which precipitation and the vegetation type of the catchment area are key players. Forestcovered soils leach tDOM, while agricultural land boosts microbial DOM production by delivering to 66 67 rivers fertilizers that fuel biological activity. High precipitation during wet seasons flushes more tDOM 68 from soils into rivers compared to low precipitation during dry seasons (Fichot et al., 2014; Li et al., 69 2015). On the other hand, the residence time of river water during wet seasons is shorter than that 70 during dry seasons, which may decrease autochthonous DOM production (Taylor et al., 2003). 71 Furthermore, DOM in rivers may be subject to physical (e.g. flocculation and coagulation, Asmala et 72 al., 2014), biological (e.g. microbial uptake, Benner and Kaiser, 2011), and chemical (e.g. 73 photobleaching, Vecchio and Blough, 2002) removals during estuarine mixing, thereby reducing its 74 abundance and modifying its chemical and optical properties before being exported to coastal seas. 75 Conversely, biological production can add organic matter to the DOM pool transported downstream from the rivers (Bianchi et al., 2004; Fellman et al., 2010; Benner and Kaiser, 2011; Deutsch et al., 76 77 2012). In highly urbanized areas, industrial and residential sewages can be a significant contributor of 78 DOM to river systems (Baker, 2001; Guo et al., 2014). Pollutions not only directly bring in 79 anthropogenic DOM but also carry nutrients that enhance biological DOM production.

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81 1.2 The Pear River estuary

The Pearl River extends for 2214 km and has a catchment area of 450,000 km² (Lloyd et al., 2003; Zhang et al., 2008), with its entire drainage basin located south of 27°N in the subtropical zone. After entering the delta area, the Pearl River becomes a complex water network because of the continuous bifurcation of three main tributaries (the West, North, and East Rivers) and other smaller rivers (Fig. 1). The Pearl River system is connected to the South China Sea via three estuaries, Lingdingyang,





Modaomen, and Huangmaohai. The Lingdingyang estuary, the principal estuary of the Pearl River, is
commonly referred to as the Pearl River estuary (PRE).

The PRE stretches for ~70 km, covers an area of ~2000 km², and has an average depth of 4.8 m (Dong et al., 2004). Its topography is featured by three shoals (the east, west, and middle shoals; depths < 2 m) separated by two channels (the east and west channels; depths >5 m) which merge in the upper reach of the estuary near Humen (Wai et al., 2004) (Fig. 1). Tides in the PRE are irregular and semidiurnal; the mean tidal range is 0.86–1.7 m, increasing landward and reaching >3 m at Humen (Zhao, 1990).

95 Ranked the second largest in China and the thirteenth largest in the world (Zhang et al., 2008), the Pearl River discharges a freshwater volume of 285×10^9 m³ year⁻¹ to the South China Sea. The West 96 River is the largest tributary, contributing 73% of the Pearl River's total freshwater discharge, followed 97 98 by the North River (14%), and the East River (8%) (Wei and Wu, 2014). The PRE receives 50-55% of 99 the Pearl River's total freshwater flow from four major water outlets, namely Humen, Jiaomen, 100 Hongqimen, and Hengmen (Mikhailov et al., 2006), with Humen providing 35% of the freshwater 101 input to the PRE, followed by Jiaomen (33%), Hengmen (20%), and Honggimen (12%) (Kot and Hu, 102 1995). About 70% to 80% of the freshwater discharge in the Pearl River occurs in the wet season 103 (April–September) and only 20–30% in the dry season (October–March) (Wei and Wu, 2014).

Freshwater in the PRE tends to flow seaward along the west side, while coastal saline water intrudes landward along the east channel, causing large cross-estuary salinity gradients (Dong et al., 2004). Seawater intrusion can reach 20–25 km downstream of Humen in the wet season and beyond Humen in the dry season. The water column is strongly stratified during the wet season due to the large freshwater input but well-mixed or far less stratified during the dry season (Wai et al., 2004; Ou et al., 2009).





The Pearl River delivers 30.64×10^6 tons of sediment per year into the PRE, with 92–96% of this discharge taking place during the wet season (Xu et al., 1985; Wai et al., 2004). The suspended sediment concentration ranges from 40–300 mg L⁻¹ in the wet season and 20–190 mg L⁻¹ in the dry season and reaches >100 mg L⁻¹ in turbidity maxima occurring at several locations of the estuary (Xu, 1985; Zhao, 1990; Wai et al., 2004).

Phytoplankton biomass in the PRE is generally higher in the wet season than in the dry season but lower than expected from the high concentrations of dissolved inorganic nitrogen (Yin et al., 2000; Harrison et al., 2008; Lu and Gan, 2015; Li et al., 2017). Phytoplankton blooms develop only on local scales, usually in the mid-estuary during the dry season and in the lower part of the estuary during the wet season (Lu and Gan, 2015).

120 Mountainous and hilly landscapes dominate the drainage basin of the Pearl River with almost no 121 forest (Luo et al., 2002), leading to relatively low dissolved organic carbon concentrations ([DOC]) $(117-132 \text{ }\mu\text{mol }L^{-1})$ upstream of the Pearl River Delta (Shi et al., 2016). On the other hand, the Pearl 122 River Delta, a highly urbanized and industrialized region, delivers 5.8×10⁹ tons of industrial and 123 domestic sewage per year into the PRE (Lu et al., 2009), which is considered the principal source of 124 125 DOC in the upper reach of the PRE (Lin, 2007; He, 2010). A number of studies have determined [DOC] and the abundance of CDOM ([CDOM]) (in terms of fluorescence or absorption coefficients) in 126 127 the PRE (e.g. Dai et al., 2000; Callahan et al., 2004; Chen et al., 2004; Hong et al., 2005; He, 2010; Lei 128 et al., 2018). These studies show no consistent seasonality and estuarine mixing behavior of [DOC] and [CDOM] and no correlation between the variables except one occasion for the mid-salinity (5-20) 129 130 section of the estuary (Callahan et al., 2004).

The lack of seasonality and consistent estuarine mixing behavior of [DOC] and [CDOM] suggests complex processes controlling their transport, production, and loss in the PRE; it could, however, also result in part from the difference in spatiotemporal coverage of the stations sampled by different





134 studies. As previous DOC and CDOM data were collected over a span of 12 and 15 years, respectively, 135 the possibility of interannual variability cannot be ruled out. In addition, none of the past DOC studies surveyed all four seasons and many of them chose two different months to represent the wet and dry 136 137 seasons, though [DOC] and its mixing behavior may change on smaller time scales. The more limited 138 number of CDOM absorption surveys only sampled a single season with no winter visits. Concerning 139 the spatial coverage, some studies differ substantially in the distribution of sampling stations (e.g. Hong 140 et al., 2005 vs. Lei et al., 2018) and many did not cover the upper reach of the estuary (e.g. Chen et al., 141 2003; Chen et al., 2004; Wang et al., 2014; Lei et al., 2018).

142 Compared with the quantitative information on DOC and CDOM, much less is known about the 143 seasonality and mixing behavior of their qualitative aspects. He et al. (2010) examined the DOC 144 compositions (monosaccharides vs. polysaccharides and dissolved free amino acids vs. dissolved 145 combined amino acids) along a longitudinal salinity-gradient transect in the PRE. Hong et al. (2005) 146 determined the fluorescence excitation-emission matrices (EEMs) on samples collected in the dry 147 season and suspected that CDOM in the PRE bears a microbial signature derived from sewage 148 effluents. Besides, spectral slope coefficient (Hong et al., 2005; Lei et al., 2018) and [DOC]-normalized 149 fluorescence intensity (Callahan et al., 2004) have been sporadically used to assess the quality of 150 CDOM in the PRE. Finally, only a few studies have estimated the DOC export flux from the Pearl 151 River to the South China Sea (Lin, 2007; Ni et al., 2008; He et al., 2010), often with limited seasonal 152 coverage. The estimate made by Lin (2007) is almost two times that by Ni et al. (2008). No estimates 153 of CDOM export are available.

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155 **1.3 Hypothesis and objectives**

Given the large volume and seasonality of the freshwater discharge of the Pearl River, we hypothesize that DOM in the PRE presents substantial seasonal variability in terms of both abundance





158 and chemical composition and that the PRE is an important source of DOM to global oceans. To test 159 this working hypothesis, the present study sampled the same locations in different seasons within a 12month period, with the objectives of 1) evaluating the seasonality and estuarine mixing behavior of 160 161 DOC and CDOM in the PRE both quantitatively and qualitatively; 2) improving the estimate of DOC 162 export to the South China Sea; 3) providing the first assessment of seaward export of CDOM in the 163 PRE based on absorption coefficient measurements. Results from this study further increase our 164 understanding of DOM cycling in human-impacted estuarine waters and their contribution to the DOC 165 and CDOM budgets in coastal oceans.

166

167 **2 Methods**

168 2.1 Sample collection

169 The sampling area covered the entire PRE, stretching from ~30 km upstream of Humen to the outer 170 limit of the estuary (Fig. 1). Ten stations (M01–M10) were distributed across the main longitudinal axis 171 of the estuary, together with two shorter along-estuary transects, each having four stations on the east 172 (E01-E04) and west (W01-W04) sides. The coordinates of the stations alongside other sampling 173 information are shown in Table 1. Water samples were collected in duplicate from the surface (\sim 1m) 174 and near the bottom (1-2 m above the seabed) using a 5-L plexiglass sampler between 8-12 May, 7-11 August, 16–19 November 2015 and 10–14 January 2016 for [DOC] measurement and in the last three 175 176 seasons for CDOM analysis. The samples were filtered through 0.2-µm polyethersulfone (PES) filters (Pall Life Sciences) under low vacuum and the filtrates were transferred into 20 mL (DOC) and 100 177 178 mL (CDOM) clear-glass bottles with Teflon-lined screw caps. DOC samples were acidified to pH \sim 2 with 2 N HCl (Reagent grade, Merck). All samples were stored in the dark at 4°C until being analyzed 179 180 in a land-based laboratory within two weeks after water collection. Prior to use, the glass filtration 181 apparatus and the sample storage bottles were acid-cleaned and combusted at 450°C for 4 h, and the





PES filters were thoroughly rinsed with Milli-Q water and sample water. Water temperature and
salinity were determined with a SBE-25 conductivity-temperature-depth (CTD) profiler.

184

185 **2.2 Sample analysis**

[DOC] was determined in triplicate using a Shimadzu TOC-L_{CPH} analyzer calibrated with potassium hydrogen phthalate. The performance of the analyzer was checked, at intervals of 10 consecutive sample analyses, against Hansell's low carbon ([DOC]: $1-2 \mu mol L^{-1}$) and deep Florida Strait ([DOC]:

189 41–44 μ mol L⁻¹) reference waters. The coefficient of variation on five replicate injections was < 2%.

190 CDOM absorbance spectra were scanned from 800 nm to 200 nm at 1-nm intervals with a Shimadzu 191 UV-2550 dual beam spectrophotometer fitted with 10-cm quartz cells and referenced to Nanopure 192 water. The samples were allowed to warm up to room temperature in darkness before analysis. A 193 baseline correction was made by subtracting the mean absorbance value over 683-687 nm from all spectral values (Babin et al., 2003). The Napierian absorption coefficient, a_{CDOM} (m⁻¹), was calculated 194 195 as 2.303 times the absorbance divided by the light pathlength of the cell in meters (0.1 m). The 196 analytical uncertainty of a_{CDOM} measurement was assessed by analyzing six pairs of duplicate samples collected from the August cruise. Average a_{CDOM} at 330 nm (a_{330}) was 2.19 m⁻¹ (range: 1.19–4.37 197 m⁻¹); the average difference in each pair was 0.07 ± 0.05 m⁻¹, or $3.0\% \pm 1.4\%$. 198

Fluorescence excitation-emission-matrices (EEMs) were acquired using a Hitachi F-4600 fluorescence spectrophotometer fitted with a 1-cm quartz cuvette to characterize the FDOM composition (Coble, 1996; Boehme et al., 2004). Again, samples were warmed up to room temperature before analysis. Emission spectra were scanned from 230 nm to 600 nm at 2-nm intervals over excitation wavelengths between 200 nm and 450 nm at 5-nm increments. Raman scattering was removed by subtracting Nanopure water EEMs that were scanned on the same day as those for the samples. The spectral fluorescence intensities were normalized to Raman Units (R.U.) following the





Raman Scatter Peak correction reported by Lawaetz and Stedmon (2009). Potential inner-filtering effects were corrected using the obtained absorbance spectra (Ohno, 2002), even though self-shading should be insignificant since the absorption coefficient at 254 nm (a_{254}) was less than 15 m⁻¹ for all samples. Major peaks in the EEMs were identified according to the peak definitions proposed in Coble (2007).

211 PARAFAC analysis was performed to decompose the EEMs into a set of underlying fluorescent 212 components (Bro, 1997; Stedmon et al., 2003; Stedmon and Bro, 2008). The analysis was fed with 117 213 EEMs from all three seasons sampled for CDOM (Sect. 2.1). To reduce the dominance of high 214 fluorescence intensity signals, the EEMs were first scaled to a unit of variance within the sample mode 215 to construct the calibration model (Bro, 1997). PARAFAC models from 2 to 7 components with 216 constraints of non-negativity in all modes were successively conducted with MATLAB (version 2008b; 217 MathWorks 2008) using DOM Fluorescence Toolbox (DOM Fluor version 1.6) and validated as 218 described by Stedmon and Bro (2008). The parameters obtained from the PARAFAC model were used to calculate an approximate abundance of each component, expressed as F_{max} in Raman's unit (R.U.), 219 220 which corresponds to the maximum fluorescence intensity for a particular sample. Based on analysis of 221 triplicate samples from Sta. M01, M08, and M10, the uncertainty of F_{max} for each modeled component 222 was <2%.

223

224 3 Results

For brevity, seasons and/or water layers for a property are added as a superscript to the symbol or abbreviation denoting that property. For example, $[DOC]^{surf/Aug}$ stands for [DOC] in surface water in August and $a_{330}^{btm/Jan}$ for CDOM absorption coefficient at 330 nm in bottom water in January. The slope of linear regression of a property against salinity will be denoted by SLP (Sect. 3.4), with a superscript added to designate the specific variable and/or season. For instance, SLP^{[DOC]/May} denotes





- the slope of [DOC] vs. salinity in May. [DOM], [CDOM], and [FDOM] stand for the abundances of
- 231 DOM, CDOM, and FDOM, respectively, and names of PARAFAC components signify their F_{max} as
- well. Finally, symbols and abbreviations are used as both singular and plural forms.

233

234 **3.1 Hydrological settings**

The average freshwater discharges of the Pearl River for the sampling months were obtained from the Ministry of Water Resources of the P. R. of China (http://www.mwr.gov.cn/zwzc/hygb/sqnb). Assuming that 54% of the total discharge of the Pearl River went into the PRE (Mikhailov et al., 2006) giving 8.9×10^3 m³ s⁻¹ in May, 5.7×10^3 m³ s⁻¹ in August, 6.7×10^3 m³ s⁻¹ in November, and 5.0×10^3 m³ s⁻¹ in January. The freshwater discharge was 15% lower in August than in November due to an atypically dry weather in summer and a relatively higher precipitation in autumn. The precipitation in January was also above average, leading to a higher-than-normal freshwater discharge in that month.

Surface water temperature averaged 27.2°C in May, 30.0°C in August, 25.2°C in November, 18.8°C 242 in January. Water temperature in August was higher in the inner than in the outer estuary, whereas a 243 reverse trend was seen in the other sampling seasons (Fig. S1). Cross-estuary gradients occurred in all 244 245 four seasons often with irregular patterns. Yet, the east transect showed the highest temperatures in 246 November and the west transect displayed the lowest temperatures in January. The difference between the surface and bottom water was minor in January (0-1.5%) and minor to moderate in May (0-11.9%)247 248 and November (0.08-2.5%) except a few stations near the mouth of the estuary. In August, the bottom 249 temperature was substantially lower (3-14%) than the surface temperature at many stations and the 250 difference increased towards the sea.

Surface water salinity (SWS) ranged from 0.2–30.3 (mean: 9.7) in May, 0.2–20.6 (mean: 8.0) in August, 0.2–26.9 (mean: 8.3) in November, and 0.2–32.6 (mean: 17.0) in January. SWS was very low (range: 0.15–0.66) and remained fairly constant upstream of Sta. M05 in May and November (Fig. S2).





254 Saltwater intruded farther upstream to Sta. M03 in January, in line with the lower tides (Fig. S3) and 255 lower freshwater discharge at that time. Despite August showing the lowest estuary-wide mean SWS 256 among the four seasons, its SWS values in the upper reach of the estuary (Sta. M01-M05) were 257 considerably higher than those in May and November, and the value at Sta. M02 even surpassed that in 258 January. This phenomenon could be partly attributed to most stations in the upper reach being sampled 259 at high tides in August (Fig. S3). Seaward of the upper reach, SWS increased rapidly, albeit with 260 fluctuations likely linked to tidal cycles and passage of salinity fronts (Dong et al., 2004). Consistent 261 with published results (Dong et al., 2004), SWS exhibited cross-estuary gradients, often increasing 262 from the shallow water on the west side to the deeper water on the east side, which was particularly 263 evident in January and May (Fig. S2). Bottom water salinity at most stations was nearly identical to 264 SWS in January, slightly greater in May, moderately elevated in November, and much higher in August 265 (Fig. S2). Based on the salinity difference between the two layers, the water column was mostly well 266 mixed in January, weakly stratified in May, modestly stratified in November, and strongly stratified in 267 August. Remarkable exceptions were certain shallow stations along the west (Sta. W01–W03) and east (Sta. E01) transects at which the water column was well mixed in November and May and weakly 268 stratified in August. In addition, the water column in the low-salinity zone (Sta. W01-W05 in May and 269 270 November; Sta. W01–W02 in August and January) was essentially homogenous in all four seasons.

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272 **3.2 Distributions of quantitative DOM variables**

The quantitative DOM variables reported here are [DOC], a_{CDOM} , and PARAFAC-derived FDOM components. a_{330} is chosen as an indicator of [CDOM] (Osburn et al., 2009; Xie et al., 2012; Song et al., 2017). a_{CDOM} at other commonly used wavelengths and the spectral slope coefficient between 300 nm and 500 nm are presented in Table S1. The residual and split-half PARAFAC analyses validated five distinct FDOM components (Fig. 2), which explained 99.75% of the variance and thus adequately





278 modeled the different FDOM profiles in the dataset. Based on published spectral characteristics of 279 PARAFAC-modeled components (e.g. Stedmon et al., 2003; Cory and McKnight, 2005; Yamashita and Jaffé, 2008; Murphy et al., 2008; Santín et al., 2009; Massicotte and Frenette, 2011), components 1 280 281 (C1) and 5 (C5) as tyrosine-like and tryptophan-like fluorophores, components 2 (C2) and 4 (C4) were 282 assigned as terrestrial or ubiquitous humic-like DOM fractions, and component 3 (C3) as oxidized 283 quinone-like moieties, respectively. As C1 and C2 are highly correlated with C5 and C4, respectively 284 (Table S2), the sum of C2 and C4 (C2+C4 hereafter) and of C1 and C5 (C1+C5 hereafter) will be used 285 to describe the quantitative distributions of the humic-like and protein-like fractions. C3 is better correlated to C2+C4 ($R^2 = 0.953$) than to C1+C5 ($R^2 = 0.738$), suggestive of its humic character. 286

Table 2 summarizes the ranges and averages of all quantitative variables in different seasons and 287 water layers. The mean $[DOC]^{surf}$ was in descending order of May > January > August > November. 288 The mean a_{330}^{surf} , (C2+C4)^{surf}, and C3^{surf} exhibited the same seasonality of August > November > 289 January, differing from that of [DOC]. The seasonal trend of the mean (C1+C5)^{surf} followed November 290 291 > January > August, inconsistent with the two trends noted above. The seasonality in bottom water was 292 the same as that in surface water for the mean values of all these variables save a_{330} which was equal 293 between November and January in the bottom. The mean values in bottom water were lower than in 294 surface water for all variables and all seasons (Table 2). The seasonal trend of the absolute percent 295 difference between the bottom and surface followed August > May > November > January for [DOC] 296 and August > November > January for the CDOM and FDOM variables, conforming to the 297 successively weakening water column stratification from summer to autumn to winter (Sect. 3.1). The 298 average vertical differences ranged from 6.5–21.0% in August, 1.0–11.9% in November, and 0.1–5.5% 299 in January depending on the variable in question, with [DOC] showing the smallest disparities (Table 2). Despite the overall small vertical gradients, certain stations, often with the deepest water depths 300 301 (Table 1), did exhibit larger differences (>20%, Fig. S4).





302 All variables displayed similar along-estuary distribution patterns characterized by overall declining 303 abundances with increasing seaward distance (Figs. S4,5). Two features are noted here. First, [DOC] in May remained nearly constant from Sta. M01 to M03, consistent with the observation of He et al. 304 305 (2010) in April 2007. As Sta. M01-M03 all had near-zero salinities (0.18-0.27, Fig. S2) and were 306 distributed across the three entrances of the East River (Fig. 1), freshwater input from this river appeared to have little influence on [DOC] in May. Second, the declines of [DOC]^{surf} across the entire 307 308 main transect (Fig. 1), 40-42% in May, August, and November, and 54% in January, were 309 considerably lower than those of a_{330} and C1+C5, 70–74% in August, 80–84% in November, and 92– 310 93% in January. The parallel declines of C3 and C2+C4 were somewhat inferior in August (53-57%) 311 but comparable in November (72–76%) and January (92%).

312 Unlike the substantial cross-estuary salinity gradients noted earlier, lateral variations in [DOC] were 313 generally small in all seasons, with one prominent exception at ~54 km downstream of Sta. M01, where [DOC] in May on the east transect (Sta. E01: 192 μ mol L⁻¹) was 47% higher than that on the 314 main transect (Sta. M05: 131 μ mol L⁻¹; Fig. S4a). Systematic cross-estuary variations in a_{330} (Fig. 315 S4e-g) and FDOM components (Fig. S5) were not evident in August and November, while values in 316 317 January were consistently higher along the west transect (Fig. S4g, Fig. S5c,f,i) echoing the 318 substantially lower salinities on the west side (Fig. S2d). Large lateral differences in C1+C5 and 319 C2+C4 were again observed between Sta. E01 on the east transect and Sta. M05 on the main transect in 320 November (Fig. S5b,e).

321

322 **3.3 Distributions of qualitative DOM metrics**

The qualitative metrics reported here are the E_2/E_3 quotient (hereafter E_2/E_3), biological index (BIX), humification index (HIX), and percentages of C1+C5 (%(C1+C5)), C2+C4 (%(C2+C4)), and C3 (%C3) relative to the sum of C1–C5. E_2/E_3 , defined as the ratio of a_{250} to a_{365} , serves as a proxy for





326 the average molecular weight (MW) and aromaticity of CDOM, with lower values indicating higher 327 MW and higher aromaticity (Peuravuori and Pihlaja, 1997; Lou and Xie, 2006; Li and Hur, 2017). BIX, the ratio of fluorescence intensity at 380 nm to that at 430 nm with excitation at 310 nm, indicates 328 329 the relative contribution of fresh, autochthonous FDOM; higher BIX values signify higher 330 contributions of freshly produced FDOM of microbial origin (Huguet et al., 2009). HIX, the ratio of the 331 fluorescence intensity integrated over 435–480 nm to that over 300–345 nm with excitation at 254 nm, 332 is a surrogate of the extent of FDOM humification, with higher values denoting higher degrees of 333 humification (Ohno, 2002). %(C1+C5), %(C2+C4), and %C3 represent the relative contents of protein-334 like, humic-like, and quinone-like components in the total FDOM pool.

Table 3 summarizes the ranges and averages of all qualitative metrics for each sampling season and water layer. Bottom–surface differences were minor in all seasons and for all metrics. C1+C5 on average accounted for 50.2–66.4% of the total FDOM components and thus exceeded C2+C4 and C3 in all seasons. The seasonal trends of HIX and %(C2+C4) were both August > November > January and opposite to those of E_2/E_3 , BIX, and %(C1+C5) (i.e., January > November > August). Note that the difference in E_2/E_3 was marginal between August and November and in all other metrics between November and January. %C3 was highest in August and equal between November and January.

 E_2/E_3 in all three seasons increased gradually down-estuary (Fig. S6a–c) by up to 59% in August, 60% in November, and 76% in January. BIX in August and November dropped briefly within the first 10 km and then augmented slowly farther seaward; it remained, however, roughly constant in January (Fig. S6d–f). Mild convex curves with maxima located at mid-estuary characterized the longitudinal HIX distributions in all three seasons (Fig. S6g–i). %(C2+C4) presented an along-estuary distribution pattern (Fig. S7d–f) similar to that of HIX and inverse to that of %(C1+C5) (Fig. S7a–c). With a few exceptions, %C3 increased nearly monotonously from land to sea irrespective of seasons (Fig. S7g–i).

349 Cross-estuary gradients were generally minor (Figs. S6 and S7). An important exception was the 350 west transect giving lower E_2/E_3 in January, lower %(C1+C5) in August, and higher HIX and





351 %(C2+C4) in both August and January, with the gradients all diminishing seaward. Bottom–surface 352 differences at individual stations were mostly marginal (<10%). Certain stations, particularly those 353 with relatively deeper water depths (Table 1), showed considerably larger differences (>20%), as noted 354 in Figs. S6 and S7.

355

356 **3.4 Relationships between quantitative DOM variables and salinity**

357 Surface and bottom data for each variable in each season form a consistent property-salinity pattern (data not shown) and are thus treated as a single dataset. All variables displayed large variations and/or 358 359 sharp decreases at salinity <5 but remained rather constant ([DOC] in May and November) or declined 360 linearly (all other cases) at salinity >5 (Figs. 3 and 4). Results of linear regressions for the saltier zone 361 are summarized in Table 4. At a 95% confidence level, both the slopes (SLP) and intercepts are 362 statistically no different between August and January for [DOC] and a_{330} and between all three seasons 363 for C3 and C2+C4, indicating that the multi-season data on each of these occasions can be combined into a single dataset. SLP^{a330/Nov} is, however, $\geq 32\%$ lower than those in August and January. SLP^{C1+C5} 364 presents significant seasonal variations, with the value in January 23% and 89% higher than those in 365 366 November and August, respectively.

The percent decrease of each variable per unit increase of salinity across the saltier zone was calculated using the known regression equations shown in Table 4. a_{330} decreased 2.1 and 2.7 times faster than [DOC] in August and January, respectively (Table S3). [FDOM] declined faster than [CDOM] but their difference was much smaller than that between [CDOM] and [DOC]. The percent decreases in the FDOM components were 5–35% higher than those in a_{330} , with November showing the largest difference (25–35%) followed by August (5–21%) and January (<10%) (Table S3).

373

374 **3.5** Relationships between qualitative DOM metrics and salinity





375 As for the quantitative variables, the surface and bottom data of the qualitative metrics can also be 376 treated as a single dataset in relation to salinity (data not shown). In August and November, E_2/E_3 increased with salinity quickly in the restricted low-salinity section (salinity: <1.3) and slowly in the 377 378 saltier zone (Fig. 5). In January, the surge at low salinities was less obvious and a gradual rise of E_2/E_3 379 in the saltier zone was observed up to salinity 28.5 beyond which the trend curved up. All three seasons 380 gave consistent E_2/E_3 vs. salinity patterns from salinity 1.3 to 28.5. In the saltier zone, the data for each 381 season roughly followed the respective theoretical mixing line defined by the maximum- and 382 minimum-salinity E_2/E_3 values in the corresponding season (Fig. 5).

383 Between salinity 0 and 5, %(C1+C5) in August decreased by approximately 10% (Fig. 6a). In the 384 saltier zone, the west transect displayed an increasing (C1+C5) with salinity but was constantly 385 below the main and east transects which formed a coherent %(C1+C5) vs. salinity pattern featured by a 386 rebound from salinity 3 to 13 and a continuous decline at salinity >13. In November, (C1+C5)387 between salinity 0 and 10 ($63.9\% \pm 5.8\%$) is more scattered than that for salinity from 10 to 27 (65.1%388 \pm 2.1%) but the average values for the two sections are very similar. %(C1+C5) for the two most 389 marine samples dropped by 5-10% compared with the average over salinity 10-27 (Fig. 6a). A pan 390 shape characterized the distribution of (C1+C5) in January, revealing higher (C1+C5) values at 391 both the lowest and highest salinities and relatively lower values across a wide range of salinities in 392 between (Fig. 6a). The distributions of %(C2+C4) vs. salinity approximately mirrored those of 393 %(C1+C5) (Fig. 6b). %C3 in August decreased with salinity along the west transect whereas it increased linearly (Y = 0.19*X + 15.61, R² = 0.867, n = 28) along the main and east transects 394 395 combined (Fig. 6c). %C3 in January stayed rather constant $(13.5\% \pm 0.8\%)$ until an abrupt 14% decline 396 at salinity >32. The distribution of %C3 in November resembled that of %(C2+C4).

Except a few larger scatters at the lowest (November) and highest (November and January) salinities, BIX displayed little dependence on salinity in all three seasons (Fig. 7a). The HIX vs. salinity patterns (Fig. 7b) corresponded to those of %(C2+C4), leading to a strong linear relationship between





the two variables (Fig. S8a). HIX is also positively correlated to %C3 (Fig. S8b), despite a weaker
correlation than that of HIX to %(C2+C4), again suggestive of the humic character of C3 (Sect. 3.2).
No significant correlation was seen between BIX and %(C1+C5) (Fig. S8c).
In spite of the certain seasonal and spatial variations of the qualitative metrics noted above, the
overall changes of these variables in the saltier zone, after excluding several extreme values for E₂/E₃

405 and BIX, were fairly limited, ranging from 4.8–9.1, 0.94–1.36, 0.54–2.04, 43.4%–70.3%, 16.5%–

406 35.9%, and 10.4%–22.6% for E₂/E₃, BIX, HIX, %(C1+C5), %(C2+C4), and %C3, respectively.

407

408 **3.6 Relationships between [DOC] and** *a***CDOM and FDOM fluorescence**

409 [DOC] was linearly correlated to a_{330} for all three sampling seasons; the coefficient of determination 410 was, however, lower in November (Fig. 8a, Table 5). The fitted slope was in descending order of January $(32.0 \pm 2.0 \text{ m } \mu \text{mol } \text{L}^{-1}) > \text{August} (22.5 \pm 1.4 \text{ m } \mu \text{mol } \text{L}^{-1}) > \text{November} (18.8 \pm 2.2 \text{ m } \mu \text{mol})$ 411 L^{-1}). Similarly, [DOC] showed a strong, linear correlation with C1+C5 in August and January and a 412 413 relatively weaker one in November (Fig. 8b, Table 5). The fitted slopes in August and January were 414 comparable but ~ 2.8 (2.7–2.9) times that in November (Table 5). [DOC] was also significantly related to C2+C4 and C3 (Fig. 8c,d) but the correlations were considerably weaker than that with C1+C5 415 416 (Table 5).

417

418 4 Discussion

419 4.1 Sources of freshwater DOM endmembers

Large variations in [DOC] and [CDOM] in the freshwater section of the PRE have been observed previously (Chen et al., 2004; Lin, 2007; He, 2010; Wang et al., 2014; Lei et al., 2018). The present study confirmed this phenomenon in August ([DOC] only) and November ([DOC], [CDOM], and [FDOM]) when near zero-salinity (< 0.7) water was accessible down to Sta. M05 off Hongqimen (Fig.

424 1). This hefty fluctuation in DOM content is commonly ascribed to the presence of multiple freshwater





endmembers delivered by various water channels and outlets described in Sect. 1.2 (Cai et al., 2004;
Callahan et al., 2004; He et al., 2010). Because Humen holds most of the sewage discharge from
Guangdong Province (Pang and Li, 2001), which carries the highest DOM load, while the other
waterways on the west coast, less influenced by urbanization and industrialization, bear lower levels of
DOM (Callahan et al., 2004; Ni et al., 2008).

430 Although the existence of multiple "quantitative" endmembers in the PRE is well recognized, it 431 remains poorly understood if these endmembers differ qualitatively. Data published by Callahan et al. (2004) shows that [DOC]-normalized fluorescences of the freshwater endmembers in Jiaomen, 432 433 Hongqimen, and Hengmen differed little (c.v. = 4%) while the Humen endmember was 17% higher 434 than the mean of the other three endmembers in November 2002. Besides, fluorescence EEMs 435 collected upstream of Humen reveal tryptophan-like fluorophores to be the dominant FDOM fraction in 436 the Humen endmember which was suspected to originate from sewage effluents (Hong et al., 2005). The present study has analyzed by far the largest number of qualitative metrics (i.e. E_2/E_3 , relative 437 438 abundances of FDOM components, BIX, and HIX) and thus offers a more comprehensive means to 439 assess the nature of the freshwater endmembers. E_2/E_3 in near zero-salinity samples fell in a rather small range from 5.5 to 6.8 that corresponded to a MW range from 0.83 kDa to 1.18 kDa estimated 440 from the MW vs. E₂/E₃ relationship proposed by Lou and Xie (2006). The higher MW values were 441 442 observed in the Humen channel, while the lower ones in water from Jiaomen and Honggimen, both 443 being close to the borderline separating the high- and low-MW CDOM (i.e. 1 kDa). %(C1+C5) varied 444 from 70% at Sta. M01 in the Humen channel to 56% off Honggimen, consistent with a stronger 445 signature of anthropogenic DOC in the Humen channel (He et al., 2010). Yet %(C1+C5) for all 446 endmembers were >50%, demonstrating that protein-like components dominated all freshwater FDOM 447 endmembers. BIX was slightly higher while HIX lower at Sta. M01 than at Sta. M05 (BIX: 1.28 vs. 448 1.00; HIX: 0.53 vs. 1.34); all BIX and HIX values were, however, well above 0.8 and below 5, 449 respectively, implying the dominance of fresh, microbial-derived FDOM in all freshwater endmembers





450 (McKnight et al., 2001; Birdwell and Engel, 2010; Sazawa et al., 2011). Taking into account all these 451 qualitative metrics and the linear correlations between [DOC] and the FDOM components (Sect. 3.6). 452 we can conclude that all three freshwater DOM endmembers in November mainly comprised fresh, 453 relatively low-MW (~1 kDa) organic material of microbial origin, with the microbial nature in the 454 Humen endmember somewhat stronger. The sewage influence could be depressed due to a rapid 455 bacterial mineralization of the sewage-derived DOM between the point sources of pollution in the 456 Guangzhou area and the sampling stations downstream (He et al., 2010). Note that the three 457 endmembers also bore a perceptible terrigenous character, since the humic-like C2 and C4, albeit lower 458 in abundance than the protein-like C1 and C5, were still a significant fraction of the total FDOM pool 459 (Fig. 6). The values of the qualitative metrics at Sta. M01 in August and January were comparable to 460 those in November (Figs. S6,7), indicating that the Humen DOM endmembers in summer and winter 461 were also of microbial origin.

Based on an estimate of the relative contributions of land-, sewage-, and phytoplankton-derived 462 DOC, He (2010) and He et al. (2010) proposed that the land component is the dominant source of the 463 total DOC pool in the lower reach of the Humen channel. In this estimation, the authors assigned the 464 "natural background" [DOC] in the three major tributaries of the Pearl River (range: 114–125 umol 465 L^{-1} ; mean: 119 µmol L^{-1}) as "land-derived". Our result suggests that, apart from terrigenous DOC 466 leached from soil, this "land-derived" DOC contains an ample amount of river-born DOC of microbial 467 468 origin. This is consistent with the poorly forested watershed of the Pearl River (Luo et al., 2002) and 469 with the study of Ni et al. (2008) showing the molar carbon-to-nitrogen ratios of suspended particulate 470 organic matter in all major runoff outlets of the Pearl River Delta (7.2-9.3) to be close to those for 471 phytoplankton and bacterial biomass (5-8).

472

473 **4.2 Estuarine mixing and transformation of DOM**





474 Sharp decreases in [DOC], [CDOM], and [FDOM] in the low-salinity section of the PRE have been 475 previously observed and postulated as a result of adsorption, flocculation, biodegradation, and/or incomplete mixing of multiple freshwater endmembers (Callahan et al., 2004; Chen et al., 2004; Lin, 476 477 2007; He et al., 2010). The present study confirmed the earlier observations but more importantly 478 provided additional qualitative metrics that are instrumental for constraining the principal processes 479 causing this swift drawdown of [DOM]. The increases in %(C2+C4) and HIX and decreases in 480 %(C1+C5) and BIX in the low-salinity section (Figs. 6 and 7) indicate a bacterial preferential uptake of 481 protein-rich materials and hence a key role of biodegradation in controlling the loss of DOM. Our 482 result corroborates the finding of He et al. (2010) showing higher fractions of biodegradable DOC and 483 higher DOC bio-uptake rates in the low-salinity section than in the saltier zone. Note that the more 484 scattering of the qualitative metrics data in November (Figs. 6 and 7) likely reflects an incomplete 485 mixing of the multiple freshwater endmembers stated earlier. This partial-mixing effect may smear or 486 even entirely overshadow the biodegradation signal.

487 In the saltier zone, the linear decreases in [DOC] (see exceptions below), [CDOM], and [FDOM] 488 with salinity point to the absence of net removal and input of these constituents and physical dilution 489 being the principal mechanism dictating their estuarine mixing behaviors. The two extreme cases of 490 near-constant [DOC] vs. salinity in May and November indicate that the loss of DOC in the low-491 salinity section reduced its content to the level comparable to the marine endmember and again that the 492 removal of DOC in the saltier zone, if any, was roughly balanced by the input. Potentially important 493 DOM loss processes in the PRE are bacterial (He et al., 2010) and photochemical (Callahan et al., 494 2004) degradation. The significance of these processes relies on both their rates and the residence time of freshwater in the PRE. Using the volume of the estuary $(9.6 \times 10^9 \text{ m}^3)$ and the freshwater discharge 495 496 rate for each sampling season (Sect. 3.1), we estimated the residence time of freshwater in the top 1-m 497 laver to be 3.1 d in May, 4.9 d in August, 4.1 d in November, and 5.6 d in January. The value for May 498 is essentially identical to that previously reported for the wet season (Yin et al., 2000). Here the volume





of the estuary was obtained from the published average depth (4.8 m) and total area (2×10^9 m²) of the 499 500 estuary (Dong et al., 2004). The bacterial uptake rate of DOC in surface water of the saltier zone has been reported to be 0.04 μ mol L⁻¹ h⁻¹ in spring and 0.07 μ mol L⁻¹ h⁻¹ in summer, giving a consumption 501 of 3.0 μ mol L⁻¹ and 8.2 μ mol L⁻¹, respectively, when multiplied by the corresponding residence time 502 for May and August. Our unpublished data indicates that photodegradation in August could at most 503 reduce [DOC] by 0.76 μ mol L⁻¹ and a_{330} by 0.11 m⁻¹, after considering the attenuation of solar 504 505 radiation and the competition for light absorption by particles in the water column (Wang et al., 2014). The combined photochemical and bacterial DOC degradation in summer was thus ~9 μ mol L⁻¹, ~8% of 506 507 the initial [DOC] in the saltier zone. The parallel photobleaching loss of a_{330} was 7%. Such small losses 508 could be readily compensated for by DOM input from in situ primary production, sediment 509 resuspension, and/or freshwater discharge farther downstream. Notably, chlorophyll a concentration maxima of up to 11.0 μ g L⁻¹ and turbidity maxima of up to 154 mg L⁻¹ were spotted in the mid- and 510 511 lower estuary during our cruises (Xu et al., unpublished data). There was, however, no co-variation of 512 [DOM] (i.e. [DOC], [CDOM], and [FDOM]) with chlorophyll a or suspended particle concentration 513 (data not shown). This observation, in conjunction with the linear [DOM] vs. salinity relations, 514 demonstrates that autochthonous production was unlikely a major source of DOM and that adsorption 515 and flocculation were not a major sink of DOM in the saltier zone. The short residence time of 516 freshwater likely minimized the influences of these processes.

The completely different behaviors of [DOC] and [CDOM] with respect to salinity in the saltier zone in November (Fig. 3c,f) led to a decoupling of the two variables. This phenomenon has also been observed for summer by Chen et al. (2004). In fact, the disconnection of [DOC] and [CDOM] is an extreme case of the higher salinity-based [CDOM] gradient relative to that of [DOC] seen in August and January (Sect. 3.4). The difference in estuarine mixing behavior between [DOC] and [CDOM] arose mainly from two factors. First, a large portion of the freshwater DOM endmember was nonand/or weakly colored, as implied by its abundant fresh microbial constituents. Second, the marine





endmember's [DOC]-normalized a_{CDOM} was lower than the freshwater endmember's: 0.60 vs. 2.18 L mg⁻¹ m⁻¹ in August, 0.71 vs. 2.32 L mg⁻¹ m⁻¹ in November, and 0.26 vs. 1.71 L mg⁻¹ m⁻¹ in January at 330 nm.

527 The overall small variations of the qualitative metrics (Figs. 5-7 and Sect. 3.5) across the saltier 528 zone suggest that the chemical composition of CDOM and FDOM remained generally stable during 529 estuarine mixing, consistent with the marginal photochemical and microbial breakdown of DOM 530 elaborated above. The higher values of %(C2+C4), %C3, and HIX in August than in November and 531 January (Figs. 6 and 7) point to FDOM in summer containing a larger fraction of humic-like 532 fluorophores. The divergence in August of the west transect from the main and east transects with 533 respect to the FDOM metrics (save BIX) distributions vs. salinity (Figs. 6 and 7) suggests a different 534 freshwater mass on the west shoal enriched with humic-like FDOM and possibly originating from 535 Hengmen (Fig. 1).

536

537 **4.3 Indicators of** *a*_{CDOM} and [DOC] in the saltier zone

538 Salinity is a useful proxy of a_{CDOM} in light of their linear relationships in the saltier zone for all three sampling seasons (Fig. 3). Furthermore, a common equation (Y = -0.048*X + 1.99) can serve as a 539 predictive tool of a_{330} in August and January, given essentially the same statistics for each of these two 540 541 months (Table 4). For [DOC], salinity can be used as an indicator in August and January but not in 542 May and November (Fig. 3). Similar to the a_{CDOM} -salinity case, the August and January [DOC] data can be combined to formulate a single [DOC]– a_{CDOM} relationship (Y = 40.7*X + 75.6). Hence, [DOC] 543 544 in summer and winter can in principle be retrieved from remote sensing-based a_{CDOM} data (Siegel et al., 545 2002; Johannessen et al., 2003; Mannino et al., 2008). Absorption coefficients and fluorescence 546 intensities at the excitation and emission maximum wavelengths of C1 and C5 are also good indicators of [DOC] in August and January (Fig. 8). 547





Caution should be exercised when applying the [DOC] and a_{CDOM} predictive tools established here, since interannual variability and other factors may limit their applicability on broader time and space scales. For example, Hong et al. (2005) arrived at an a_{CDOM} -salinity relationship of $a_{355} =$ -0.045*salinity + 1.81 for November 2002, which is different from ours in the saltier zone ($a_{355} =$ -0.021*salinity + 0.98). Concurrent measurements of [DOC] and a_{CDOM} in the PRE are rare but Chen et al. (2004) reported no significant correlation between the two variables in July 1999.

554

555 4.4 Fluxes of DOC and CDOM

556 The fluxes of DOC and CDOM exported from the PRE to the South China Sea were estimated as 557 follows (Cai et al., 2004; Lin, 2007; He et al., 2010):

558
$$\mathbf{F} = \mathbf{Q} \times \mathbf{C}^* \tag{1}$$

where F denotes the flux of DOC or CDOM, Q the freshwater discharge rate, C^{*} the effective [DOC] 559 ([DOC]^{*}) or $a_{330} (a_{330}^*)$. C^{*} is the y-axis intercept of the regression line of [DOC] or a_{330} vs. salinity in 560 561 the saltier zone (Table 4). For May and November when [DOC] remained roughly constant across the saltier zone, C^{*} signifies the average [DOC] over this zone. Monthly fluxes were computed using 562 freshwater discharge rates for the sampling year and those averaged over 2006-2016 563 564 (http://www.mwr.gov.cn/zwzc/hygb/sqnb), under the assumption that the [DOC] or a_{330} obtained for 565 May, August, November, and January represents the entire spring (March, April, May), summer (June, July, August), autumn (September, October, November), and winter (December, January, February), 566 respectively. As no CDOM data was collected in May, the a_{330}^* for spring $(1.99 \pm 0.19 \text{ m}^{-1})$ was 567 derived from the mean of the [DOC]^{*}-normalized a_{330}^* in January (1.31 L mg⁻¹ m⁻¹) and August (1.36 568 L mg⁻¹ m⁻¹) multiplied by the $[DOC]^*$ in May (124.5 µmol L⁻¹). This treatment, with unknown 569 uncertainties, was based on the relatively small variations of the $[DOC]^*$ -normalized a_{330}^* among the 570 three CDOM sampling seasons (range: $1.31-1.50 \text{ Lmg}^{-1} \text{ m}^{-1}$). 571





572 Flux estimates for the sampling year are comparable to those for the 10-year period for spring and 573 summer, whereas the former is approximately twice the latter for autumn and winter due to aboveaverage freshwater discharge rates during the low-flow season of the sampling year (Table 6). 574 Aggregation of the fluxes for all four individual seasons arrives at an annual export of 240×10^9 g C 575 (sampling year) or 195×10^9 g C (10-year period) for DOC and of 329×10^9 m² (sampling year) or 266 576 $\times 10^9$ m² (10-year period) for CDOM in terms of a_{330} . As the PRE receives ~54% of the total Pearl 577 River freshwater discharge to the South China Sea (Mikhailov et al., 2006), including the rest 46% 578 gives a grand annual export of 362×10^9 g C of DOC and 493×10^9 m² CDOM, respectively, assuming 579 580 that the fluxes from the PRE are applicable to the entire Pearl River Delta.

581

582 4.5 Comparison with previous studies and other major estuaries

583 The [DOC]s obtained by this study in all four seasons are within the ranges previously reported for 584 the PRE (Table 7). DOC stock in the PRE thus did not seem to undergo large changes over the 7-year span from the last survey in 2008 to our study in 2015, suggesting that the gross inputs and losses of 585 DOM remained stable during this period. Compared to DOC, previous a_{CDOM} measurements are far 586 587 fewer and none of them was made in wintertime. The summer and autumn a_{330} values from this study are, however, comparable to those published (Table 7). Our DOC flux estimate for spring 2015 (5.8 \times 588 10^8 g C d⁻¹) is close to that reported by He et al. (2010) for spring 2007 (5.3 × 10^8 g C d⁻¹). The 589 summer 2015 value $(9.0 \times 10^8 \text{ g C d}^{-1})$ is, however, only 60% of the summer 2007's (He, 2010) due to 590 a much lower river runoff in 2015 (7174 m³ s⁻¹ vs. 25060 m³ s⁻¹). The DOC flux for the entire Pearl 591 River Delta estimated by this study $(362 \times 10^9 \text{ g C year}^{-1})$ is comparable to that $(380 \times 10^9 \text{ g C year}^{-1})$ 592 reported by Ni et al. (2008) but 44% lower than that $(650 \times 10^9 \text{ g C year}^{-1})$ obtained by Lin (2007). The 593 estimate by Ni et al. (2008) was based on monthly [DOC] measurements at eight major runoff outlets 594 595 of the Pearl River Delta from March 2005 to February 2006. Lin (2007) derived the estimate from data





596 collected during three cruises carried out in winter (February 2004), early spring (March 2006), and 597 summer (August 2005). Part of the difference between our study and Lin's could result from the 598 different temporal coverages. The main difference, however, stems from the much greater [DOC]^{*}

599 obtained by Lin (2007) (147 μ mol L⁻¹ for the wet season and 254 μ mol L⁻¹ for the dry season).

600 [DOC] and [CDOM] in the PRE are the lowest among the major world rivers (Table 8). The low 601 DOM load in the PRE could be associated with a deficiency of organic matter in soil of the Pearl 602 River's watershed having almost no forest (Luo et al., 2002). Moreover, although sewage effluents may bring in large amounts of DOM, a big portion of it can be rapidly bio-degraded before reaching the 603 head of the estuary (He et al., 2010). The lack of correspondence between $[DOC]^*$ and a_{330}^* and the 604 freshwater discharge rate (Fig. S9) suggests that [DOM] in the PRE be controlled by both soil leaching 605 606 and pollution input. In contrast, DOM in the majority of large rivers is predominantly terrigenous 607 (Bianchi, 2011; Raymond and Spencer, 2015) and the abundance of DOM in many rivers increases 608 with the river flow rate (Cooper et al., 2005; Holmes et al., 2013). Note that the absence of a link 609 between [DOC] and the freshwater discharge rate in the PRE observed by this study differs from the 610 anti-covariation of the two variables reported by Lin (2007) and Ni et al. (2008). Based on this anti-611 variation, Lin (2007) proposed that the PRE is a typical point source-regulated system in terms of DOC 612 concentration and distribution. It remains to be confirmed if our results imply a fundamental shift of the 613 PRE within ~10 years from a pollution-dominated system to a system jointly controlled by pollution 614 and soil flushing.

Owing mainly to the very low [DOC], our DOC export estimate for the Pearl River is the lowest among the 30 largest rivers worldwide (Raymond and Spencer, 2015), though the Pearl River is ranked the 13th largest river by discharge volume. The Pearl River value of 362×10^9 g C year⁻¹ only accounts for 0.14% of the global riverine DOC flux estimate of 250×10^{12} g C year⁻¹ (Raymond and Spencer, 2015). The estimate for CDOM export from the Pearl River is also the lowest among the limited





number of estimates available for the major world rivers (Table 9). It is worth noting that, despite its
small contribution on global scales, DOM delivered by the Pearl River is rich in labile, proteinaceous
constituents that can be readily utilized by microbes, thereby exerting a potentially important impact on
the local ecosystem.

624

625 5 Conclusions

626 The saltier zone of the PRE manifests smaller temporal variabilities and lower spatial gradients of 627 DOM than expected for a sizable estuary with a marked seasonality of river runoff. Several factors 628 functioning in concert lead to this phenomenon. First, a combination of the poorly forested watershed, 629 rapid degradation of pollution-derived DOM in the upper reach, and short residence time of freshwater 630 diminishes [DOM] and seasonal variations in both DOM quantity and quality. Second, the small 631 difference between the low-salinity and marine DOM endmembers tends to lessen the vertical and 632 lateral gradients in DOM again both qualitatively and quantitatively, notwithstanding the larger vertical 633 and cross-estuary salinity gradients. Both the concentrations and seaward exports of DOC and CDOM 634 in and from the PRE are the lowest among the major world rivers. However, as DOM undergoes 635 marginal processing during its transit through the estuary, the Pearl River delivers protein-rich, labile 636 organic matter to the continental shelf of the South China Sea where it may fuel heterotrophy.

637

638 Author contributions. GS and HX designed the study. HX and GS interpreted the results and prepared

the manuscript with input from PM. YL performed sample analysis and data processing. YL, GS, FY,

and RL carried out field sampling. PM conducted PARAFAC modeling.

641

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





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896	Figure captions
897	
898	Figure 1. Map of sampling stations in the Pearl River Estuary. Station names starting with letters M,
899	W, E designate the main, west, and east transects, respectively. See Table 1 for coordinates of the
900	stations. HM: Humen; JM: Jiaomen; HQM: Hongqimen; HeM: Hengmen; MDM: Maodaomen; HMH:
901	Huangmaohai.
902	
903	Figure 2. Excitation-emission contours of five components identified by PARAFAC modeling (left
904	panels) and split-half validations of excitation and emission loadings (right panels). Excitation/emission
905	maximum wavelengths are: C1: 275/320 nm; C2: <240(335)/426 nm; C3: 245/378 nm; C4:
906	255(370)/464 nm; C5: <240(290)/348 nm.
907	
908	Figure 3. DOC concentration and a_{330} versus salinity in the PRE. Red circles denote samples collected
909	in the low-salinity section where DOC and a_{330} showed rapid decreases or large variabilities with
910	salinity. Blue circles denote the samples collected in the saltier zone. Solid lines in panels a and c
911	represent means of the blue circles. Solid lines in the other panels denote linear fits of the blue circles.
912	Dashed lines signify the 95% confidence intervals. See Table 4 for fitted equations and statistics.
913	
914	Figure 4. Same as in Figure 3b,d,e-g except for FDOM components C1+C5, C2+C4, and C3.
915	
916	Fiure 5. E_2/E_3 versus salinity for each cruise. Lines denote conservative mixing lines defined by the
917	lowest- and highest-salinity points in the saltier zone.
918	
919	Figure 6. Percentages of FDOM components versus salinity for each cruise. Red solid circles denote
920	samples collected along the west transect (see Figure 1) in August.
921	
922	Figure 7. BIX (a) and HIX (b) versus salinity. Red solid circles denote samples collected along the
923	west transect (see Figure 1) in August.
924	
925	Figure 8. DOC concentration versus a_{330} (a), C1+C5 (b), C2+C4 (c), and C3 (d). Solid lines denote
926	linear fits of data for each cruise. See Table 5 for fitted equations and statistics.
927	

 $\bigcirc \text{Author}(s) \text{ 2018. CC}$









surf denotes surface and bt	m bottom.				
	DOC	a_{330}		FDOM (R.U.)	
	$(\mu mol L^{-1})$	(m^{-1})	C1+C5	C2+C4	C3
		May			
Surface	160 (110–243)	NA	NA	NA	NA
Bottom	155 (114–234)	NA	NA	NA	NA
((btm-surf)/surf)*100	-3.2	NA	NA	NA	NA
		Augus	st		
Surface	117 (96–167)	1.92 (1.07-4.35)	0.90 (0.43–2.02)	$0.51\ (0.26-0.80)$	0.30(0.17 - 0.46)
Bottom	109 (78–166)	1.60 (0.56-4.27)	0.71 (0.16–1.97)	0.41 (0.11–0.87)	$0.24\ (0.08-0.51)$
((btm-surf)/surf)*100	-6.5	-16.4	-21.0	-19.2	-18.6
		Novem	ber		
Surface	83 (77–133)	1.42(0.54 - 3.35)	1.21 (0.47–2.30)	0.40(0.14-0.61)	0.24(0.10 - 0.37)
Bottom	82 (70–100)	1.29 (0.60–3.40)	1.10(0.20 - 2.28)	0.36 (0.08–0.67)	$0.21 \ (0.06 - 0.38)$
((btm-surf)/surf)*100	-1.0	-8.7	-8.7	-9.8	-11.9
		Januar	y		
Surface	118 (71–194)	1.34 (0.29–3.98)	1.03 (0.24–3.09)	0.30 (0.06–0.69)	$0.20\ (0.04-0.51)$
Bottom	118 (66–207)	1.29 (0.33-4.11)	0.98 (0.23–2.95)	0.29 (0.06–0.72)	0.19(0.04-0.53)
((btm-surf)/surf)*100	-0.1	-1.2	-5.5	-4.1	-3.6

Table 2 Means (randow) of DOC α_{220} and intensities of fluorescent commonents in surface and hoftom waters for each cruise. Here

931 932

933 934





surf denotes surface an	d btm bottom.					
	E_2/E_3	BIX	HIX	%(C1+C5)	%(C2+C4)	%C3
			August			
Surface	6.84 (4.76–8.29)	1.04 (0.95–1.14)	1.56 (1.01–2.36)	51.6 (40.2-62.2)	30.5 (23.6–39.1)	18.0 (14.2–20.7)
Bottom	6.98 (5.18–8.74)	1.07 (0.94–1.25)	1.60 (0.97–2.28)	50.2 (41.2-62.8)	31.0 (23.0–38.3)	18.8 (14.2–22.6)
((btm-surf)/surf)*100	2.0	3.4	2.4	-2.6	1.8	4.4
			November			
Surface	6.89 (5.48–9.13)	1.13 (1.02–1.30)	1.01 (0.53–1.51)	64.3 (54.2–71.1)	22.2(18.5–29.7)	13.5 (10.7–16.1)
Bottom	7.17 (6.02–8.44)	1.09 (0.68–1.30)	1.03 (0.86-1.50)	64.0 (52.4–70.3)	22.6 (18.9–31.9)	13.5 (10.4–16.5)
((btm-surf)/surf)*100	4.0	-3.8	2.5	-0.6	1.7	0.1
			January			
Surface	7.60 (6.08–11.16)	1.15 (1.04–1.53)	0.90 (0.54–1.26)	66.4 (61.6–72.2)	20.2 (15.9–23.5)	13.5 (11.7–14.9)
Bottom	7.61 (6.10–10.38)	1.16(1.03-1.36)	0.88 (0.64–1.19)	66.3 (61.8–71.7)	20.3 (16.5–23.5)	13.4 (11.9–14.8)
((btm-surf)/surf)*100	0.2	0.5	-2.7	-0.1	0.5	-0.4





- **Table 4.** Results of linear regression (Y = a*X + b) of DOM quantitative variables
- 940 against salinity. SE denotes standard error.

	a±SE	b±SE	\mathbb{R}^2	р	
		DOC			
August	-1.31±0.16	121.3±2.4	0.72	< 0.0001	
January	-1.11±0.21	123.0±4.5	0.57	< 0.0001	
		a_{330}			
August	-0.042 ± 0.006	1.97±0.09	0.67	< 0.0001	
November	-0.029 ± 0.003	1.47 ± 0.05	0.80	< 0.0001	
January	-0.048 ± 0.003	1.93 ± 0.07	0.88	< 0.0001	
-		C1+C5			
August	-0.023 ± 0.004	0.94 ± 0.06	0.62	< 0.0001	
November	-0.035 ± 0.005	1.43 ± 0.10	0.67	< 0.0001	
January	-0.043 ± 0.004	1.61 ± 0.08	0.82	< 0.0001	
-		C2+C4			
August	-0.016 ± 0.001	0.60 ± 0.02	0.83	< 0.0001	
November	-0.014 ± 0.001	0.52 ± 0.01	0.94	< 0.0001	
January	-0.014 ± 0.001	0.53 ± 0.02	0.90	< 0.0001	
C3					
August	-0.008 ± 0.001	0.33±0.01	0.74	< 0.0001	
November	-0.008 ± 0.001	0.31±0.01	0.91	< 0.0001	
January	-0.008 ± 0.001	0.32 ± 0.01	0.86	< 0.0001	





- **Table 5.** Results of linear regression (Y = a*X + b) of [DOC] against a_{330} and FDOM
- 944 components. SE denotes standard error.

	a±SE	b ±SE	\mathbb{R}^2	р
		a_{330}		
August	22.5±1.4	72.5±2.9	0.89	< 0.0001
November	18.8 ± 2.2	61.8±3.3	0.68	< 0.0001
January	32.0±2.0	71.4±3.3	0.90	< 0.0001
		C1+C5		
August	43.5±2.1	76.2±2.0	0.93	< 0.0001
November	16.3±1.9	66.5±2.5	0.68	< 0.0001
January	40.7±2.3	72.6±3.0	0.91	< 0.0001
		C2+C4		
August	100.7±11.4	64.3±5.9	0.72	< 0.0001
November	52.2±9.8	65.2±4.0	0.47	< 0.0001
January	159.2±13.7	66.7±4.9	0.82	< 0.0001
		C3		
August	197.0±17.6	57.4±5.3	0.80	< 0.0001
November	106.8±16.3	61.1±3.9	0.56	< 0.0001
January	242.5±16.2	65.8±4.0	0.89	< 0.0001

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discharge rates for the sampling year and those averaged over a 10-year period from 2006 to 2016. Standard errors of the fluxes for the

Table 6. Estimates for DOC and CDOM (a_{330} -based) export from the Pear River to the South China Sea based on monthly freshwater

 





953 **Table 7.** DOC concentrations and a_{330} in surface water of the Pearl River estuary

Month	DOC (μ mol L ⁻¹)	Sampling Year	Reference
Jan.	71–194	2016	This study
Feb.	100–247 ^a	2004	Lin (2007)
Mar.	109–266	1997	Dai et al. (2000)
	103–229 ^a	2006	Lin (2007)
Apr.	84–278 ^b	2007	He et al. (2010) He (2010)
May	110-243	2015	This study
	58–160 ^c	2001	Callahan et al. (2004)
Jul.	109-315	1996	Dai et al. (2000)
	68–250	1999	Chen et al. (2004)
Aug.	96–167	2015	This study
	107–164 ^a	2005	Lin (2007)
	94–124 ^b	2008	He (2010)
Nov.	77–133	2015	This study
	82–187 ^c	2002	Callahan et al. (2004)
Month	$a_{330} (\mathrm{m}^{-1})$	Sampling Year	Reference
Jan.	0.29-3.98	2016	This study
May	$0.37 - 7.48^{d}$	2014	Lei et al. (2018)
Jul.	1.01-3.38 ^d	2013	Wang et al. (2014)
	0.54-1.98	1999	Chen et al. (2004)
Aug.	1.07-4.35	2015	This study
Nov.	0.54-3.35	2015	This study
	0.38-2.73	2002	Hong et al. (2005)

954 reported in the literature and this study.

955

- ^aRanges were estimated using the fitted [DOC]-salinity equations in Lin (2007) over
- 957 salinity 0-30.

958 ^bDOC concentrations upstream of Sta. M01 in the present study are excluded.

- 959 ^cValues were retrieved from figures 5a and 8b in Callahan et al. (2004).
- 960 ^dRanges were estimated using exponential decay equations established from data in table
- 961 1 in Lei et al. (2018).





River	DOM	References
DOC (μ mol L ⁻¹)		
Amazon	235	Raymond and Bauer (2001)
	277	Cao et al. (2016)
	307 (122-492)	Seidel et al. (2016)
Mississippi	489 (231–672)	Bianchi et al. (2004)
	417 ^a	Spencer et al. (2013)
Atchafalaya	331 ^a	Spencer et al (2013)
St Lawrence	307 (25–1333)	Hudon et al. (2017)
	231 ^a	Spencer et al (2013)
Mackenzie	375+100	Cooper et al. (2015)
WINCKCHZIC	347 (258-475)	Raymond et al. (2003)
	402 (250-576) ^b	Osburn et al. (2007)
	+02(250-570) 262(250,475)	Stadmon at al. (2007)
Vulton	503(230-473) 523+242	Steamon et al. (2011)
т икоп	333 ± 242	Cooper et al. (2005)
	509 (217–1258)	Kaymond et al. (2007)
	5/4-	Spencer et al. (2013)
	674 (200–1617)	Stedmon et al. (2011)
Kolyma	500±167	Cooper et al. (2005)
	594 (250–1025)	Stedmon et al. (2011)
Lena	724±283	Cooper et al. (2005)
	775 (542–1233)	Raymond et al. (2007
	948 (550–1600)	Stedmon et al. (2011)
Ob	733±167	Cooper et al. (2005)
	780 (458–1000)	Raymond et al. (2007)
	875 (375-1058)	Stedmon et al. (2011)
Yenisey	733±316	Cooper et al. (2005)
-	638 (242–1050)	Raymond et al. (2007)
	754 (208–1250)	Stedmon et al. (2011)
Yellow	202 (151–280)	Wang et al. (2012)
Yangtze	169 (137–228)	Wang et al. (2012)
Pearl River	149 (72–243) ^c	This study
Amazon	$a_{330} (\mathrm{m}^{-1})$	C_{22} at al. (2016)
Miggigginni	0.60^{a}	Ca0 et al. (2010)
Atabafalaya	11.55^{a}	Spencer et al. (2013)
St. Lawrence	0.65°	Spencer et al. (2013)
St. Lawrence	9.00 2.16 ^a	All et al. (2012)
Maalaansia	2.10 8.20 (5.10, 12.20) ^b	Spencer et al. (2013)
wiackenzie	$6.30(5.19-13.30)^{\circ}$	Osburn et al. (2009)
X7 1	0.04(3.01-9.03)	Steamon et al. (2011)
Y ukon	17.34	Spencer et al. (2013)
** 1	14.50 (2.65–37.84)	Stedmon et al. (2011)
Kolyma	13.63 (5.77–29.19)	Stedmon et al. (2011)
Lena	26.51 (15.48–52.94)	Stedmon et al. (2011)
Ob	22.43 (6.74–30.74)	Stedmon et al. (2011)
Yenisey	22.14 (3.50–44.79)	Stedmon et al. (2011)
Yangtze (Changjiang)	$2.60(2.29-3.02)^{1}$	Song et al. (2017)
Pearl River	$2.50(1.04-4.35)^{\circ}$	This study

^aRetrieved from DOC and CDOM fluxes and freshwater discharge rates in Spencer et al. (2013). 964

965 ^bFrom data at salinities <5

966

⁶From data at salinities <5. ^dRetrieved from the spectral slope and a_{350} at Sta. 10 in Cao et al. (2016) 967

968 ^eAverage value at Sta. SL1 and SL2 in Xie et al. (2012).

969 ^fAverage value at salinities <5.





- 970 **Table 9.** CDOM fluxes (a_{330} -based) from major world rivers to the ocean reported in the
- 971 literature. The flux estimated for the Pearl River by this study is also included for
- 972 comparison.

River	Flux ($\times 10^9$ m ² year ¹)	Reference
Mississippi	5070	Spencer et al. (2013)
Atchafalaya	2750	Spencer et al. (2013)
St. Lawrence	490	Spencer et al. (2013)
Mackenzie	1550	Stedmon et al. (2011)
Yukon	3520	Spencer et al. (2013)
	3260	Stedmon et al. (2011)
Kolyma	1340	Stedmon et al. (2011)
Lena	17100	Stedmon et al. (2011)
Ob	7350	Stedmon et al. (2011)
Yenisey	12600	Stedmon et al. (2011)
Pearl River	266	This study

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Figure 1. Map of sampling stations in the Pearl River Estuary. Station names starting
with letters M, W, E designate the main, west, and east transects, respectively. See Table
1 for coordinates of the stations. HM: Humen; JM: Jiaomen; HQM: Hongqimen; HeM:
Hengmen; MDM: Maodaomen; HMH: Huangmaohai.







Figure 2. Excitation-emission contours of five components identified by PARAFAC
modeling (left panels) and split-half validations of excitation and emission loadings (right
panels). Excitation/emission maximum wavelengths are: C1: 275/320 nm; C2:
<240(335)/426 nm; C3: 245/378 nm; C4: 255(370)/464 nm; C5: <240(290)/348 nm.







Figure 3. DOC concentration and a_{330} versus salinity in the PRE. Red circles denote samples collected in the low-salinity section where DOC and a_{330} showed rapid decreases or large variabilities with salinity. Blue circles denote the samples collected in the saltier zone. Solid lines in panels a and c represent means of the blue circles. Solid lines in the other panels denote linear fits of the blue circles. Dashed lines signify the 95% confidence intervals. See Table 4 for fitted equations and statistics.













Figure 5. E_2/E_3 versus salinity for each cruise. Lines denote conservative mixing lines 1000 defined by the lowest- and highest-salinity points in the saltier zone.







1002

Figure 6. Percentages of FDOM components versus salinity for each cruise. Red solid circles
denote samples collected along the west transect (see Figure 1) in August.







1006

1007 Figure 7. BIX (a) and HIX (b) versus salinity. Red solid circles denote samples collected

along the west transect (see Figure 1) in August.







1011 Figure 8. DOC concentration versus a_{330} (a), C1+C5 (b), C2+C4 (c), and C3 (d). Solid lines

1012 denote linear fits of data for each cruise. See Table 5 for fitted equations and statistics.