A comprehensive conceptual framework for signaling in-lake $CO₂$ through dissolved organic matter

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Highlights:

● Lake CO₂ and DOM turnovers follow common biogeochemical pathways

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- Subtropical lakes can be carbon sinks or sources due to varying $CO₂$ mechanisms
- DOM biodegradation governs temporal variations in CO₂ relative to terrestrial

inputs

- Photo-mineralization of activated aromatic compounds fuels lake $CO₂$
-

CO₂ levels are well modelled with 0.4%–2.1% overestimations *via* DOM signals

A comprehensive conceptual framework for signaling in-lake CO² through

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dissolved organic matter

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Abstract

Organic carbon (C) and $CO₂$ pools are closely interactive in aquatic environments. While there are strong indications linking freshwater $CO₂$ to dissolved organic matter (DOM), the specific mechanisms underlying their common pathways remain unclear. Here, we present an extensive investigation from 20 subtropical lakes in China, establishing a comprehensive conceptual framework for identifying $CO₂$ drivers and retrieving CO² magnitude through co-trajectories of DOM evolution. Based on this framework, we show that lake $CO₂$ during wet period is constrained by a combination of biogeochemical processes, while photo-mineralization of activated aromatic compounds fuels CO² during dry period. We clearly determine that biological degradation of DOM governs temporal variations in $CO₂$ rather than terrestrial C inputs within the subtropical lakes. Specifically, our results identify a shared route for the uptake of atmospheric polycyclic aromatic compounds and $CO₂$ by lakes. Using machine learning, in-lake $CO₂$ levels are well modelled through DOM signaling regardless of varying CO² mechanisms. This study unravels the mechanistic underpinnings of causal links between lake $CO₂$ and DOM, with important implications for understanding obscure aquatic $CO₂$ drivers amidst the ongoing impacts of global climate change.

Keywords: Inland water CO₂, drivers and magnitude, dissolved organic matter, subtropical lakes, carbon cycling, biogeochemical pathways.

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1. Introduction

Natural lakes act as dynamic reactors, emitters and sinks for aqueous $CO₂$ turnover, representing a substantial contribution to global carbon (C) cycle (Alin and Johnson, 2007; Pi et al., 2022). Most lakes (and reservoirs) show supersaturation in relation to atmospheric CO₂, with global CO₂ evasion estimated to range from 0.06 to 0.84 Pg C yr⁻¹ (Raymond et al., 2013). The emission spectrum is admittedly broad, demonstrating that CO₂ drivers for current lakes are highly variable (Finlay et al., 2015; Perga et al., 2016; Weyhenmeyer et al., 2015). The prevailing consensus is that inland water $CO₂$ are fuelled by two main mechanisms: terrestrially-derived C inputs (Heathcote et al., 2015; Li et al., 2018) and actual degradation of organic C (Allesson et al., 2021; Begum et al., 2023). Terrestrial fluxes of CO² and dissolved organic C (DOC) released from soils into surface waters, for instance, typically account for $CO₂$ excess: the former may rapidly escape to the atmosphere (Campeau et al., 2019), while the latter will participate in aquatic degradation within lakes (Maberly et al., 2013). Biological respiration and photochemical mineralization of DOC result in the actual production of in-lake $CO₂$ (Dempsey et al., 2020). Consequently, it is important to recognize that the evolution of $CO₂$ and DOC likely shares common pathways. Yet to date, a conceptual framework for unraveling in-lake $CO₂$ through these cotrajectories with DOC is not well established, despite mounting evidence of their specific associations driven by geographical and biological processes (Hessen et al., 2017; Lapierre and Giorgio, 2015; Lapierre et al., 2013).

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Dissolved organic matter (DOM), refers to the specific material form of DOC, involving the processes that govern organic C transport and transformation in natural waters (Lynch et al., 2019; Ni and Li, 2022). Lakes collect large quantities of allochthonous organic litters from surrounding shorelines, providing an important source, together with soil leaching, for DOM (Nakhavali et al., 2021; Wilkinson et al., 2013). Aquatic biology, meanwhile, functions as primary producers and contributors to autochthonous DOM pool in lakes (Hu et al., 2022). A large amount of natural DOM will undergo decomposition and eventually mineralize into CO₂ via photochemical and biological processes (Kellerman et al., 2015; Milstead et al., 2023). The DOM molecular composition, source and fate are interrelated within freshwater organic C pool and processing (Butturini et al., 2022). For example, inlake lignin and soil fulvic acid originate commonly from terrestrial inputs (Cory et al., 2007; Yang et al., 2021), whereas carbohydrates, lipids and proteins are indicative of biological metabolism (Zhang et al., 2020). Microbial degradation and photobleaching alter DOM structure, and further modify molecular weight and recalcitrance (Hansen et al., 2016; Helms et al., 2014; Logozzo et al., 2021). All these characterizations can be simply extracted through DOM spectroscopy and molecular data analysis (Chen et al., 2019; Xu et al., 2020). From this perspective, DOM potentially serves as a signaling catalyst of organic C evolution.

Natural DOM can be established as a nexus, encompassing data on DOC origins and turnover, while also decodes the allocation of $CO₂$ *via* common DOC pathways

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(Bodmer et al., 2016; Ni et al., 2023; Zhang et al., 2024). Specifically, the mechanistic underpinnings of the causal relationships between lake DOM and $CO₂$ involves: 1) the co-trajectories of lake CO² and organic C evolution (Begum et al., 2023; Lapierre et al., 2013); and 2) the inherent attributes of DOM that signal organic C source and fate (Kujawinski et al., 2009; Ni and Li, 2020). It is anticipated that dissolved $CO₂$ drivers can be identified through DOM signaling, as demonstrated by previous observations in aquatic environments (Kang et al., 2023; Ni et al., 2020b). However, these connections may be decoupled by spatiotemporal shifts of organic C pathways and mutual interferences among drivers (Winterdahl et al., 2016), making them not universally present. For instance, biologically derived connections between DOM and CO² could be decoupled due to inhibited respiration and strengthened geochemical connections e.g., terrestrial inputs and photochemical processes over space and time. Within this framework, the question arises whether there is a universal determination for retrieving both $CO₂$ levels and sources, in spite of dynamic $CO₂$ drivers and potential interferences. Furthermore, how can we figure out other $CO₂$ contributions that are unable to be gauged by explainable DOM variables.

In this study, we aim to identify and allocate in-lake $CO₂$ by linking with DOM signals, addressing the mechanistic underpinnings and ultimately providing a comprehensive conceptual framework for understanding these relationships. We collected 249 observations from 20 subtropical lakes in China, to quantify aqueous CO² levels, and analyse DOM spectroscopy and molecular information. Our

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hypothesis postulated that the common pathways involved in $CO₂$ and DOM will vary at timescales attributing to biogeochemical shifts in subtropical lakes. In testing this hypothesis, we specifically: 1) examined the dynamics of aqueous $CO₂$ magnitude in lakes; 2) characterized DOM molecular composition, origin and fate; and 3) identified $CO₂$ drivers and modelled $CO₂$ levels through DOM signaling. This study proposes a DOM-based path for capturing varying $CO₂$ mechanisms in lakes, which can be universal and transferable for other natural waters.

2. Materials and methods

2.1. Study area and sampling

During August to April from 2022 to 2023, we sampled 134 sites from 20 lakes in subtropical China, geographically extending from 23º24′ to 31º32′N in latitude and from 98º57′ to 106º34′E in longitude (Fig. 1). Our sampling encompassed the bulk of representative subtropical lakes across Yunnan-Guizhou Plateau and Sichuan Basin, representing a broad spectrum of lake surface areas $(<10-330 \text{ km}^2)$. We realized that CO² and organic C biogeochemistry is potentially constrained by sampling locations. For instance, terrestrial C and nutrient levels are typically higher along the shorelines in comparison to offshore areas (Biddanda and Cotner, 2002). Consequently, our investigation involved sampling from both shorelines and more distant offshore areas to ensure spatial representativeness of lake samples (Fig. S1). Most lakes were revisited to incorporate temporal patterns of wet (May–October) and dry periods (December–April) in the subtropical region, resulting in a total of 249 observations

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being collected. We acknowledge that geographical locations can affect regional climate and hydrology. Therefore, we carefully distinguish the wet and dry periods during the investigation. Specifically, it mostly includes mountainous lakes to minimize potential interferences from groundwaters and human activities. Being in the monsoonal climate zone, sampling locations exhibit mean annual temperature and precipitation varying within 10–20℃ and 700–1300 mm yr⁻¹, respectively. The terrain is composed largely of mountains and hills, with elevations ranging from 260 to 2700 m. Detailed information on sampling locations see Supplement Materials (Appendix Table A1 and Fig. S1).

Before water sampling, gas was collected *in-situ* from the atmosphere using gas sampling bag at each lake. Surface water $(\sim 10 \text{ cm})$ was collected at each site, and 1 L water samples were filtered through glass microfiber filters (GF/F 47, 0.7 μm, Whatman). The filtrates were refrigerated in 100 mL high-density polyethylene (HDPE) plastic containers at 4°C. *In-situ* pH and water temperature (T) were measured using a portable CyberScan PCD 650 multi-parameter system (Eutech, USA), while wind speed (U_0) was determined with a Testo410-1 anemometer (Testo, Germany). Total alkalinity was titrated with Alkalinity Test MColortestTM (Merck, Germany). Subsequent UV absorbance and fluorescence spectroscopy were conducted for all water sample $(n = 249)$. Given the high experimental cost, we mixed water samples (equal volumes) from each of the lakes as composite samples for Fourier transform ion cyclotron resonance mass spectrometry (FT-ICR MS)

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characterization to acquire comprehensive insights into molecular-level DOM. While this approach does not result in information loss regarding the DOM molecules, it presents a potential limitation in analysing the temporal variations of molecular DOM. Two small lakes were excluded due to insufficient filtrate volumes for FT-ICR MS, leaving a total of 18 composite samples. The subsequent solid-phase extraction (SPE) and FT-ICR MS analysis were commissioned to the China National Analytical Center, Guangzhou.

2.2. Laboratory analysis

Gas samples were determined for atmospheric $CO₂$ using gas chromatography-flame ionization detection (Fuli Analytical Instrument Co., Ltd., China). Water chemistry, UV absorbance and fluorescence spectroscopy were analysed for the filtered water samples. Lake DOC and dissolved inorganic carbon (DIC) concentrations were measured using a multi N/C 2100S analyzer (Analytik Jena, Germany) in triplicate. Chromophoric DOM was assessed using a double-beam scanning spectrophotometer (UV-5500 PC, Shanghai). The absorption spectra ranged from 200 to 700 nm with an interval of 1 nm. Fluorescence DOM was analysed using a RF-6000 Spectrophotometer (Shimadzu, Japan). Excitation-emission matrices (EEM) were scanned from 200 to 450 nm with a 5-nm interval for excitation wavelengths, and from 250 to 600 nm with a 1-nm interval for emission wavelengths.

Molecular composition of lake DOM was determined using FT-ICR MS (Bruker

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Solari X, Germany). For SPE, water samples were acidified with hydrochloric acid to pH 2.0, and passed through a PPL column (200 mg, 3 mL). The PPL column was activated using 30 mL methanol, and salt was removed using 30-mL Milli-Q water and hydrochloric acid. The flow rate was kept at $1-2$ mL min⁻¹, and samples were washed with 3 mL methanol (HPLC grade). Negative electrospray ionization (ESI) ion source was injected into a Bruker Solarix XR 7.0T FT-ICR MS system at a flow rate of 120 μ L h⁻¹. The injection and extraction voltages were was set to 4.0 kV and -500 V for capillary inlets. The mass-to-charge ratios (m/z) were within the range of 100–1200, with an ion accumulation time of 0.02 or 0.06 s. During FT-ICR MS analysis, $m/z = 369.119106$ served as the internal reference ion for real-time mass calibration. It acquired 300 single transients/scans to improve the signal-to-noise ratio (S/N) of the target peaks. For high-quality FT-ICR MS, Suwannee River natural organic matter (SRNOM) was used as the standard for calibrating mass axis.

2.3. Lake CO² calculation and DOM characterization

We defined aqueous partial pressure of CO_2 (pCO_2) as CO_2 level in lakes, which was calculated from pH, water temperature, and total alkalinity or DIC using CO2SYS program (D.E. Pierrot et al., 2006). This program is developed on a basis of water chemistry or carbonate equilibria, enabling the calculation of DIC species concentrations, as well as pCO_2 or fugacity of CO_2 (fCO_2). The fCO_2 is about 0.3% lower than the pCO_2 under pressures of \sim 1 atm due to the non-ideality of CO_2 . Carbonate equilibria are also constrained by the solubility coefficient of $CO_2(K_0)$,

first (K_1) and second dissociation (K_2) constants of carbonic acid in natural waters as follows:

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$$
fCO_2 = \frac{DIC}{K_0} \times \frac{[H^+]^2}{[H^+]^2 + K_1 \times [H^+] + K_1 \times K_2} \tag{1}
$$

To better understand the potential $CO₂$ dynamics within the subtropical lakes, we computed the areal CO² efflux at distinct spatial and temporal scales. We estimated normalized gas transfer velocity $(k_{600}, \text{m d}^{-1})$ of CO₂ by relating it to wind speed, using previously reported empirical models (Li et al., 2016). It was converted into site-specific gas transfer velocity $(k, m d^{-1})$ with water temperatures and Schmidt numbers. Henry's constant of $CO_2(K_h, \text{mmol m}^{-3} \mu \text{atm}^{-1})$ was corrected for *in-situ* temperature and pressure. The disparity between in-lake (pCO_{2water} , µatm) and atmospheric pCO_2 (pCO_{2air} , μ atm) drives the transfer of CO_2 . We noted relatively high CO₂ concentrations (434–496 ppm, with a mean of 463 ± 20 ppm) and thus pCO_{2air} in each lake (Fig. S2), potentially upscaling the baseline for assessing lake $CO₂$ outgassing. By measuring and compiling these parameters across sampling locations, we calculated water-air areal CO_2 efflux (mmol m⁻² d⁻¹) using thin boundary layer (TBL) model as follows:

$$
CO_2 \text{ efflux} = k \times K_{\text{h}} \times (p \text{CO}_{2 \text{water}} - p \text{CO}_{2 \text{air}}) \tag{2}
$$

For detailed descriptions of estimating areal $CO₂$ efflux in the study subtropical lakes see Supplementary Information S1.

Chromophoric DOM was characterized by the non-normalized absorption coefficients of wavelengths at 254 (a_{254} , a proxy for aromatic abundance) (Ni et al., l

2020a), 350 (a_{350} , a proxy for lignin abundance) (Derrien et al., 2019) and 420 (a_{420} , an indication for photochemical mineralization of lake DOC) (Koehler et al., 2016) in Naperian units. Although these absorption coefficients are highly correlated, they have specific implications for DOM characterization, and the use of multiple coefficients is more conducive to accurate prediction of DOC concentration (Fichot and Benner, 2011). SUVA254, a proxy for DOM aromaticity (D'Andrilli et al., 2022), was further calculated based on a₂₅₄ and DOC concentration. Spectral slope coefficient (S275-295), calculated from exponential fitting to the absorption spectrum of 275–295 nm, was used as a proxy for DOM microbial degradation, photobleaching and molecular weight (Helms et al., 2008). Spectral slope ratio (S_R) , the ratio of S_{275} . 295 to S₃₅₀₋₄₀₀ (computed similarly to S₂₇₅₋₂₉₅), had a negative correlation with DOM molecular weight. Fluorescence DOM was modelled using fluorescence regional integration across five EEM regions (Chen et al., 2003): region I (tyrosine-like), II (tryptophan-like) and IV (microbial exudates) represent biogenic DOM; region III (soil fulvic acid-like) links to terrestrial DOM; and region V expresses humic-like DOM. Fluorescence index (FI) and humification index (HIX) can indicate lake DOM origin and fate (Wilson and Xenopoulos, 2009). Specifically, FI was calculated from emission intensities of 450 to 500 nm at an excitation of 370 nm; while HIX was determined from total emission intensities of 435–480 nm to 300–345 nm at an excitation of 254 nm. We identified the main fluorophores with maximum fluorescence intensity (F_{max}) in the 20 subtropical lakes using parallel factor analysis (PARAFAC). The component models were assessed and validated using residual

analysis and split half analysis. For detailed descriptions of computing chromophoric and fluorescence DOM see Table S1.

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Molecular composition of DOM was analysed using a molecular formula calculator based on criteria with elemental combinations of $C_{0-\infty}H_{0-\infty}O_{0-\infty}N_{0-1}S_{0-1}$, and assigned to the peaks with $S/N > 4$. For FT-ICR MS data, the modified aromaticity index (AI-mod), double bond equivalent (DBE) and nominal oxidation state of carbon (NOSC) were calculated for each assigned molecular as follows:

$$
AI_mod = \frac{1 + C - \frac{1}{2}O - S - \frac{1}{2}(N + H)}{C - \frac{1}{2}O - N - S}
$$
(3)

$$
DBE = 1 + \frac{1}{2}(2C - H + N)
$$
(4)

$$
NOSC = \frac{4C + H - 3N - 2O - 2S}{C}
$$
(5)

Van Krevelen analysis was used to decipher molecular composition from 7 distinct regions based on the elemental ratios of H/C and O/C (Wang et al., 2023): carbohydrates (1.5 \times H/C $<$ 2.4; 0.71 $<$ O/C $<$ 1.2); amino-sugars (1.5 $<$ H/C $<$ 2.2; $0.52 < O/C < 0.71$; N > 0); saturated compounds $(1.5 < H/C < 2.2$; 0 < O/C < 0.52; N > 0); tannin $(0.5 \leq H/C \leq 1.5$; $0.67 \leq O/C \leq 1.2$); lignin $(0.7 \leq H/C \leq 1.5$; $0.1 \leq O/C$ < 0.67); unsaturated hydrocarbons ($0.7 < H/C < 1.5$; $0 < O/C < 0.1$); condensed aromatic structures $(0.2 < H/C < 0.7; 0 < O/C < 0.67)$. We included a visual map to aid in understanding the classification in Fig. S3. Furthermore, molecules were categorized into five compound groups based on AI-mod, H/C and O/C cutoffs (Hu et al., 2023; Linkhorst et al., 2017): combustion-derived polycyclic aromatic structures $(CA, AI-mod > 0.66)$, soil-derived polyphenols and PCAs with aliphatic chains

 $(SP&PCAs, 0.66>AI-mod>0.50)$, soil-derived humic and highly unsaturated compounds (SH&UNs, AI-mod \leq 0.50, H/C < 1.5), unsaturated aliphatic compounds (UAs, $2.0 \geq H/C \geq 1.5$), and saturated fatty and sulfonic acids, carbohydrates (SF&SAC, H/C \geq 2.0 or O/C \geq 0.9).

2.4. Statistical analysis and data uncertainty

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We tested normality and homogeneity of variance using Kolmogorov-Smirnov and Levene's test, respectively. Variables were log-transformed when it was statistically necessary to assume normality. Mann-Whitney *U* test was used to examine temporal variability of in-lake $CO₂$ and DOM datasets. One-way analysis of variance (ANOVA, with Turkey HSD post hoc) allowed us to assess statistical differences in the abundance or proportion of DOM molecular compositions. Linear regression and correlation analysis were assigned to indicate the homology of distinct DOM compositions, and show possible associations of $pCO₂$ with DOM variables. Given these linear relationships were likely impacted by mutual interactions between variables, we conducted an in-depth identification of how lake DOM signals CO₂ sources and forecasts $CO₂$ levels using machine learning. Specifically, random forest model was performed to identify relative importance of $CO₂$ drivers and predict $CO₂$ magnitude in lakes. It constructs multiple decision trees using random subsets of the training data, and synthesizes all predictions from each tree or node. This approach, handles high-dimensional and complex datasets of aqueous gas emissions (Rocher-Ros et al., 2023), demonstrating the ability to manage irrelevant variables without

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causing overfitting. We defined the weights of relative importance (%) as the increased errors caused by permuting variables. The minimum leaf node size and number of trees were set to 2 and 500, respectively. The training and testing datasets have a ratio of approximately 2:1, containing 162 and 85 data points, respectively. The optimal results with the highest R^2 were selected to represent the regressions between observed and modelled *pCO*₂ across multiple modeling. In this study, mathematical statistics were performed using OriginLab OriginPro 2024 and SPSS statistical package (19.0), while random forest model was developed using MATLAB (2021b) algorithm.

Dissolved CO₂ concentrations calculated from water chemistry or carbonate equilibria may involve systematic errors caused by potential issues with pH and alkalinity measurements (Liu et al., 2020). For pH calibration, the breakpoint is often pH 5.4 in data handling: this includes measurements with $pH > 5.4$ to filter out implausible readings and instrument errors (Hotchkiss et al., 2015). Also, a low pH may indicate significant interference from organic acids to DIC species computing. In this study, we demonstrate 95% of pH measurements > 7.0 , suggesting that $CO₂$ calculation errors resulting from pH could be relatively minimal (Fig. S4). For total alkalinity calibration, we realized that non-carbonate alkalinity sourced from organic and inorganic acids (e.g., borate, phosphate, silicate and hydrogen sulfate) can result in significant errors, particularly for low-concentration total alkalinity $\left($ <1000 µmol L⁻ $¹$) (Liu et al., 2020). In this context, we quantified an average uncertainty of 19% for</sup>

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total alkalinity using ancillary data e.g., DOC, dissolved phosphate and nitrogen. In fact, carbonate alkalinity and DIC in freshwaters are conceptually interchangeable terms in water chemistry calculations and thus CO2SYS program (D.E. Pierrot et al., 2006). Therefore, we constrained lake $pCO₂$ uncertainties using total alkalinity-based and directly measured DIC. We show that consistent with the evaluation from ancillary data, DIC and $pCO₂$ were overestimated by 19% and 23% from total alkalinity-based determination, respectively (Fig. S5). Therefore, we used measured DIC instead of total alkalinity in estimating lake $pCO₂$ in this study, in order to minimize the uncertainties. This converted the systematic errors into instrument errors of DIC (and DOC) measurements (< 2% in triplicate). UV absorptions, spectral slopes and S_R were analyzed in duplicates for 10% data, suggesting an uncertainty of $\leq 2\%$. The inner filter effects of EEM data could be minimal (Ohno, 2002), with over 99% of the recorded absorbance at 254 nm being less than 0.3. This left us with one sample that was diluted with Milli-Q water to address the interference issue. Raman and Rayleigh scatter was collected for fluorescence data using interpolation (Bahram et al., 2006). Fluorescence intensities were normalized to Raman Unit (R.U.) using water Raman peak area (Lee et al., 2018).

3. Results and discussion

3.1. Subtropical lake CO² levels and dynamics

Our investigations encompassed mountainous lakes from the Sichuan Basin to the Yunnan-Guizhou Plateau in subtropical China (Fig. 1). Lake pCO_2 spanned four

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orders of magnitude, ranging from <10 to 9175 μatm (17–3385 μatm, 95% confidence intervals) with a mean of 736 ± 1289 µatm. The diverse magnitudes indicate substantial fluctuations in $CO₂$ drivers within these lakes. Particularly, $pCO₂$ in the wet period had a broad range (Fig. 1a), averaging much higher at 1063 ± 1691 uatm than the dry period of 413 ± 534 µatm (Mann-Whitney *U* test, p < 0.001). This is likely modulated by temporal rainfall and soil flushing, therefore influencing lateral $CO₂$ inputs (Vachon et al., 2017). We found large spatial shifts in $pCO₂$ levels in the study subtropical lakes, especially high averages > 2000 μatm for locations with human disturbances e.g., Dianchi, Xihu and Qinglong (Fig. S6). It should be noted that \sim 70% of the samples were undersaturated with CO₂, exhibiting a high potential for C sequestration in subtropical lakes. Lake Chenghai, for instance, retained DIC concentrations of 5413–9228 μ mol L⁻¹ but yielded an unusually low pCO_2 of only 163 $± 76$ μatm.

The modelled *k* values varied from 0.23 to 6.6 m d⁻¹ (0.44–1.3 m d⁻¹, 95%) confidence intervals) and averaged at 0.76 ± 0.53 m d⁻¹ (Fig. S7), which is consistent with the global estimates (0.74 m d^{-1}) from similar wind speed models (Raymond et al., 2013). Combining paired measurements of $pCO₂$ and k values taken at each sampling location, we further calculated areal $CO₂$ efflux ranging from -66 to 231 mmol m⁻² d⁻¹ (-15–64 mmol m⁻² d⁻¹, 95% confidence intervals) for these subtropical lakes (Fig. S8). The mean value, 5 ± 33 mmol m⁻² d⁻¹, was significantly different from the median value of -6 mmol m⁻² d^{-1} , for the CO₂ fluxes. This raises a conflicting

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observation as these subtropical lakes overall represent a net $CO₂$ efflux because of some uniquely high values, despite most sampling locations functioning as $CO₂$ sinks. Site-specific extreme events, for example particular human disturbance (Luo et al., 2022; Raymond et al., 2008), may be responsible for this. Our areal $CO₂$ efflux is lower than the previously reported mean value $(27 \pm 66 \text{ mmol m}^{-2} \text{ d}^{-1})$ for lakes and reservoirs across China (Ran et al., 2021), yet aligns with a recent real-time estimation $($ < 10 mmol m⁻² d⁻¹) for lakes in Yunnan-Guizhou Plateau (Sun et al., 2023). It is important to note that we excluded non-carbonate alkalinity contributions by using detected DIC, and employed a relatively high baseline for atmospheric $CO₂$ based on *in-situ* measurements (Fig. S2). These calibrations avoided potential overestimations and resulted in relatively low magnitudes of lake $CO₂$ evasion estimated. Our findings suggest that the study lakes emitted CO_2 in the wet period (14 \pm 42 mmol m⁻² d⁻¹), but switched to uptake CO_2 in the dry period (-3 \pm 15 mmol m⁻² d⁻¹, Fig. S8). This implies that C turnover, along with $CO₂$ drivers, may differ substantially at timescales in these lake ecosystems.

3.2. Subtropical lake DOM characterizing

Lake DOC concentrations varied from 1.0 to 50.4 mg L⁻¹ (1.9–23.0 mg L⁻¹, 95% confidence intervals), averaging ~2-fold higher in the wet period (11.4 \pm 7.63 mg L⁻¹) than the dry season $(5.9 \pm 4.88 \text{ mg L}^{-1}, p < 0.001)$ (Fig. 2a). Aromatic and lignin abundance, as well as photo-mineralization potentials of lake DOM, were greater in the wet period, as evidenced by absorption coefficients a_{254} , a_{350} and a_{420} (p < 0.001).

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DOM aromaticity (SUVA₂₅₄), however, was higher in the dry period ($p < 0.001$, Fig. 2a). This demonstrates concentrating effect of aromatic compounds during periods of drought stress. Alternatively, biological degradation of DOM can also modulate its aromaticity over time (Ni and Li, 2020). We suggest that photobleaching and microbial degradation drove DOM decay, resulting in large accumulation of photoand bio-metabolites, as well as alterations in DOM relative molecular weight in the dry period, as indicated by spectral slope coefficient $S_{275-295}$ (Fig. 2a). As a result, seasonality can alter DOM photochemical and metabolic pathways in lakes, governing complete (mineralization) or incomplete (biolysis and photobleaching) degradation of organic C. Shifting sunlight and DOM composition over seasons may straightforwardly account for these effects (Cory et al., 2014; Gonsior et al., 2013). It further combined with $S_{350-400}$ to compute S_R , demonstrating that DOM relative molecular weight was statistically similar across time periods ($p > 0.05$, Fig. 2a).

The PARAFAC modeling identified four fluorescent components in the subtropical lakes (Fig. 2b), which were closely associated with terrestrial (C1) and biogenic humic-like (C2), as well as tryptophan-like DOM (C3 and C4). The accumulation of C1 and C3 in particular, sourced from soil leaching and algal metabolism, was intensified by the drought conditions (Fig. S9). Specifically, the coexisting C3 and C4 during the wet period, two alternative tryptophan-like components, represented a changeable relationship in F_{max} (Fig. S10), which likely resulted from a transition state caused by biological degradation of DOM (Ni and Li,

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2023). We realized that PARAFAC modeling may omit individualized components, and thus it employed fluorescence regional integrations for a comprehensive understanding of fluorescence DOM (Fig. 2c). Our results suggest that humic-like DOM (region V) had the highest proportion of 13%–81% across the five EEM regions (ANOVA, $p < 0.001$). Biogenic (region $I + II + IV$) and terrigenous DOM (region III) showed no significant temporal variations (Mann-Whitney U test, p) 0.05), contributing 12%–85% and 2%–17% to the EEM regions, respectively. This demonstrates a robust buffer capacity within the subtropical lakes, able to withstand seasonal fluctuations in DOM evolution. FI varied between 1.5–2.1 with a mean of 1.8 ± 0.10 (Fig. 2d), indicating that autochthonous and allochthonous sources can be equally significant for these fluorophores (implications of FI see Table S1 for more details). We found that soil fulvic acid-like component largely contributed to DOM humification in the lakes, as evidenced by the associations between region III and HIX (Fig. S11).

The DOM molecular formulae were highly consistent across the lakes, as shown by H/C and O/C distributions in van Krevelen Diagram (Fig. S12). Terrestrially derived lignin compounds (or carboxy-rich acyclic molecules) were predominated, contributing 77%–86% of molecular DOM (Fig. 3a). Amino-sugars and saturated compounds, in particular, also exhibited the significant proportions of 2%–4% and 8%-18%, respectively. These DOM molecules, can be preferentially biodegraded in aerobic environments with high H/C >1.5 (Sleighter et al., 2014;

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Spencer et al., 2015). We thus suggest that terrestrial inputs and aquatic metabolism, as well as their potential interactions (Hotchkiss et al., 2015), may act as substantial pathways for organic C evolution in the lakes. We further classified DOM into five compound groups, demonstrating that SH&UNs were highly prevalent, with proportions of 71%–83% (Fig. 3b). This aligns with terrigenous signals from fluorescence DOM characterizing (Fig. 2). By contrast, DOM hydrogenation increased relative abundance of UAs (13%–26%), along with low O/C (< 0.5, Fig. S12) and NOSC $(< 0$, Fig. 3c), resulting in the accumulation of photo-labile compounds (McDonough et al., 2022) within the subtropical lakes. It is especially the case for lake Qinglong, of which DOM had higher AI-mod and DBE (Fig. 3c). Here, photo-produced aromatics may interact with DOM biodegradation, as proposed by a recent study (Hu et al., 2023). Therefore, it is anticipated that biotic processes also concatenate photochemical pathways of DOM, constraining organic C turnover in the study lakes.

3.3. Lake DOM signaling CO² drivers and magnitudes

Organic C shares common pathways with $CO₂$, establishing their causal links in aquatic environments. This supports previously documented associations between DOM and dissolved CO₂ (Li et al., 2024; Luo and Li, 2021). However, these relationships are typically particular at timescales. By linking $pCO₂$ to DOM variables, for instance, we found that a combination of biogeochemical drivers i.e., terrestrial inputs, photo-mineralization and biological metabolism, governed lake $CO₂$

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during wet period. In contrast, photochemical processes likely modulated aquatic $CO₂$ during dry period (see Fig. S13). This was in agreement with temporal shifts of $pCO₂$ levels (Fig. 1), given that photo-mineralization of organic C corresponds only a small fraction of $CO₂$ evasion (Allesson et al., 2021) relative to more diverse sources. It should be noted that from the perspective of DOM molecules, aromaticity or aromatic structures were strongly associated with $pCO₂$, serving as an important indicator for lake $CO₂$ levels (Fig. S14).

Nevertheless, this effort to elucidate lake $CO₂$ cannot rank the drivers and predict $CO₂$ levels. Here, we performed machine learning to identify and allocate in-lake $CO₂$ (Fig. 4). The reliable results from linear regression analysis (Fig. S13) were complemented by random forest modeling, further showing that microbial degradation of DOM (with the relative high importance of biogenic signals i.e., region II , IV and FI) dominantly explained lake pCO_2 among multifaceted CO_2 drivers in the wet period (Fig. 4a). Moreover, the modeling additionally identified significant importance of the terrigenous (a_{350}) and photochemical signals (a_{420} and an ancillary signal $S₂₇₅₋₂₉₅$) in the dry period (Fig. 4b), which reveals that terrestrial inputs, together with photolytic degradation of DOM, constrained lake $pCO₂$. This supports our hypothesis, suggesting the biogeochemical co-trajectories involved in lake $CO₂$ and DOM evolution can greatly differ over time. We highlight that these $CO₂$ mechanisms function not only independently but also interact within organic C processing. By examining the interactions of DOM variables, we can determine that

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aquatic metabolism of terrestrial DOM and photo-mineralization of activated aromatic compounds emerged as two potential pathways for $CO₂$ production in the subtropical lakes (see Fig. S15).

On the basis of these biogeochemical co-trajectories, we clearly show robust correlations between observed and modelled pCO_2 across time periods (Fig. 4c). This indicates there is a generic way to retrieve $CO₂$ levels from DOM signaling, despite temporal variations in $CO₂$ mechanisms. Particularly, the modelled mean $pCO₂$ levels showed good agreement with the measured values, with overestimations of only 0.4% and 2.1% in the wet and dry periods, respectively (Fig. 4d). However, regarding high $pCO₂$ (> 2500 μ atm), the modelled values had significant uncertainties. This may arise from the contributions not fully gauged by the "explainable DOM variables" (Please refer to Fig. 5 for the definition of this term). For instance, our study identified a shared pathway for the uptake of atmospheric polycyclic aromatic compounds and CO² by lakes from a molecular perspective (Fig. S14), which could not be simply omitted particularly in subtropical lakes. Photosynthetic consumption of pCO_2 also complicated predictions of lake $CO₂$ levels (Fig. S16). Particulate organic C typically accounts for 7%–38% of the total organic C pool (Ostapenia et al., 2009; Rouillard et al., 2011) and transforms into DOC through fragmentation (Kiuru et al., 2018), which potentially contributes $51-280$ μatm to lake $CO₂$ levels on average. We note that China's subtropical lakes are susceptible to eutrophication and algal blooms, depending on temporal temperature changes (Yindong et al., 2021) and spatial

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locations e.g., eutrophication-induced algal blooms often emerge along the shoreline rather than in the relative central areas (Stadig et al., 2020; Wang et al., 2018). This potentially shapes spatiotemporal patterns of $CO₂$ levels in these subtropical lake ecosystems (Fig. S17). Mounting evidence that human activities and groundwater can regulate aquatic DOM and CO₂ (Connolly et al., 2020; Duvert et al., 2018; Kang et al., 2023; McDonough et al., 2022). Yet, there is still a lack of the specific explainable DOM variables to correspond with these inputs.

In this study, we propose a DOM-based path for understanding $CO₂$ drivers and magnitudes in lakes (Fig. 5). The mechanistic basis relies on the ubiquitous biogeochemical co-evolution of aquatic $CO₂$ with DOM. It develops a comprehensive conceptual framework to unveil coupling relationships between diverse DOM indicators and aquatic $CO₂$, showing clearly how to differentiate and identify $CO₂$ sources. Furthermore, we suggest that machine learning can unravel the intrinsic connections between various drivers despite potential interferences, and thus establish a universal determination for retrieving $CO₂$ in most natural waters. However, it should be noted that the specific $CO₂$ drivers and associated DOM-based models may vary between distinct aquatic environments e.g., arid regions and saline lakes. Regardless of predictable levels and identifiable sources of lake $CO₂$ and organic C, this determination can provide significant implications for C turnover in aquatic ecosystems. We realize that biological degradation of organic C apparently governs lake $CO₂$ magnitude. For example, microbial metabolism acting as a determinant

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drove much higher $CO₂$ levels during wet period rather than terrestrial inputs during dry period (Fig. 4). This understanding deviates from the prevailing view (Drake et al., 2018; Einarsdottir et al., 2017; Lapierre et al., 2013), yet aligns with the observations in shallow lakes (Bogard et al., 2019), suggesting that terrestrial C inputs may not be the fundamental hotspots of $CO₂$ emissions in subtropical lakes. Subtropical lakes sequester a significant amount of dissolved C but emit less $CO₂$ into the atmosphere (Raymond et al., 2013) due to biogenic constraints, demonstrating the significance in C capture and subsequent C neutrality. Our efforts to signal aqueous CO² through DOM, are thus anticipated to provide new insights into C cycling in subtropical lakes amidst the ongoing impacts of global climate change.

4. Conclusion

Natural DOM shares specific biogeochemical pathways with $CO₂$ in aquatic environments, establishing mechanistic underpinnings of their causal links. Based on these associations, we present a comprehensive conceptual framework to identify and allocate $CO₂$ through DOM signals in China's subtropical lakes. We demonstrate that lake CO₂ was constrained by terrestrial inputs, photo-mineralization and biological metabolism during the wet period, while photochemical processes largely fuelled $CO₂$ during the dry period. The interacting mechanisms involving aquatic metabolism of terrestrial DOM and photo-mineralization of activated aromatic compounds were specifically identified as key drivers of lake $CO₂$. Our findings highlight the crucial role of aquatic biology in regulating $CO₂$ dynamics relative to terrestrial C inputs

within subtropical lakes. Using machine learning, aqueous $pCO₂$ levels were well modelled through DOM signaling, achieving average overestimations of 0.4% and 2.1% in the wet and dry periods, respectively. This study proposes a generalizable approach for unraveling aquatic $CO₂$, which may provide important insights into C cycling from inland waters amidst ongoing climate change.

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Declaration of Competing Interest

All authors agree this submission and the authors declare that there is no conflict of interests regarding the publication of this article

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manuscript. All authors reviewed the manuscript. Journal R

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Figure captions:

Fig. 1. Spatiotemporal distributions of aqueous pCO_2 across the study lakes in subtropical China. Dots show the mean pCO_2 in each lake during the wet period (a) and dry period (b). Blue lines describe drainage networks across the study area. Detailed information of sampling locations is provided in Fig. S1.

Fig. 2. Spectroscopic characteristics of DOM in the subtropical lakes. (a) Temporal pattern of DOC and chromophoric DOM in the lakes. The boxes with whiskers represent 25%–75% range within 1.5IQR. White and black lines show median and mean, respectively. Dots correspond to all data of UV parameters. (b) Threedimensional view of primary fluorophores identified by PARAFAC modeling. (c) Fluorescence regional integration across five excitation-emission matrix regions. The waves display Kernel Smooth distributions of proportions for each region. (d) Distributions of FI and HIX across the time periods. The red line and grey area indicate linear regression and 95% confidence band, respectively. Dots correspond to all data of fluorescent parameters. The waves show Kernel Smooth distributions of the data.

Fig. 3. Molecular information of DOM across the subtropical lakes. (a) Proportions of DOM molecular compounds in each lake, (b) Proportions of compound groups include combustion-derived polycyclic aromatic structures (CA), soil-derived polyphenols and PCAs with aliphatic chains (SP&PCAs), soil-derived humic and highly unsaturated compounds (SH&UNs), unsaturated aliphatic compounds (UAs), and saturated fatty and sulfonic acids, carbohydrates (SF&SAC). (c) The modified

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Fig. 4. Lake DOM signaling CO₂ drivers and magnitudes using machine learning. (a) Random forest modeling assigning the relative importance of DOM variable in evaluating lake $CO₂$. The x axis with whisker shows the mean importance with standard deviation (s.d.). (b) The visual flows of relative importance from DOM variable to temporal pCO_2 in the lakes. The red and blue flows represent the most variables with weights > 0.2. (c) The relationships between observed and modelled *p*CO² across the time periods. The waves show Kernel Smooth distributions of the data. (d) Comparison of observed and modelled $pCO₂$ in the lakes. The boxes with whiskers represent 25th–75th percentiles with s.d. White and black lines show median and mean, respectively. Dots correspond to all data of observed and modelled $pCO₂$. **Fig.** 5 A comprehensive conceptual framework for signaling in-lake $CO₂$ through DOM. We suggest that lake organic C or DOM follows the shared biogeochemical pathways with lake $CO₂$. It includes the common drivers 1) e.g., photochemical drivers, aquatic metabolism, terrestrial inputs and atmospheric uptake that are directly signalled by explainable DOM variables; and 2) e.g., groundwater inputs, human activities and photosynthesis that co-vary with lake $CO₂$ although explainable variables signaling these relationships are lacking. The term "explainable DOM variables" refers to DOM variables that can specifically indicate certain sources or pathways regarding biogenic, terrigenous, and photochemical processes. By contrast, ancillary signals may interact with various pathways but are not specifically indicative

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of any single explainable process. The causal relationships can be visualized through linear regression analysis, correlation analysis or machine learning. This helps to identify $CO₂$ drivers, rank contributions to lake $CO₂$, and particularly discover interactive trajectories in modulating lake $CO₂$ (e.g., aquatic metabolism of terrestrial DOM, and photo-mineralization of activated aromatic compounds). By compiling and analysing these common pathways, it allows us to predict lake $CO₂$ levels using machine learning. We highlight that this conceptual framework could be generic and transferable for other natural waters.

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Graphical Abstract

Declaration of Interests

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All authors agree this submission and the authors declare that there is no conflict of interests regarding the publication of this article

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