



- 1 Spatiotemporal transformation of dissolved organic matter along an alpine stream
- 2 flowpath on the Qinghai-Tibetan Plateau: importance of source and permafrost
- 3 degradation
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- 5 Yinghui Wang ^{a,b}, Robert G.M. Spencer ^c, David Podgorski ^d, Anne Kellerman ^c,
- 6 Harunur Rashid^a, Phoebe Zito^d, Wenjie Xiao^b, Dandan Wei^a, Yuanhe Yang^e, Yunping
- 7 Xu ^{a*}
- 8 ^a Shanghai Engineering Research Center of Hadal Science and Technology, College of Marine
- 9 Sciences, Shanghai Ocean University, Shanghai 201306, China.
- 10 ^b Key Laboratory for Earth Surface Processes of the Ministry of Education, College of Urban and
- 11 Environmental Sciences, Peking University, Beijing 100871, China.
- 12 ^c National High Magnetic Field Laboratory Geochemistry Group and Department of Earth, Ocean,
- 13 and Atmospheric Science, Florida State University, Tallahassee, FL 32306, USA
- ¹⁴ ^d Pontchartrain Institute for Environmental Sciences, Department of Chemistry, University of New
- 15 Orleans, New Orleans, LA, 70148, USA
- 16 ^e State Key Laboratory of Vegetation and Environmental Change, Institute of Botany, Chinese
- 17 Academy of Sciences, Beijing 100093, China
- 18 *Corresponding author. *E-mail*: ypxu@shou.edu.cn (Y. Xu)
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Abstract The Qinghai-Tibetan Plateau (QTP) accounts for approximately 70% of 20 global alpine permafrost and is an area sensitive to climate change. The thawing and 21 mobilization of ice and organic carbon-rich permafrost impact hydrologic conditions 22 23 and biogeochemical processes on the QTP. Despite numerous studies of Arctic permafrost, there are no reports to date for the molecular-level in-stream processing of 24 25 permafrost-derived dissolved organic matter (DOM) on the QTP. In this study, we examine temporal and spatial changes of chemical composition of DOM and ¹⁴C age 26 of dissolved organic carbon (DOC) along an alpine stream (3850-3207 m above sea 27 28 level) by Fourier transform ion cyclotron resonance mass spectrometry (FT-ICR MS), accelerator mass spectrometry (AMS) and UV-visible spectroscopy. Compared to 29 downstream sites, the DOM at the headstream exhibited older radiocarbon (¹⁴C-DOC) 30 age, higher mean molecular weight, higher aromaticity and fewer polyunsaturated 31 components. At the molecular level, 6409 and 1345 formulas were identified as unique 32 to the active layer (AL) leachate and permafrost layer (PL) leachate, respectively. 33 Comparing permafrost leachates to the downstream site, 59% of AL-specific formulas 34 and 90% of PL-specific formulas were degraded, likely a result of rapid instream 35 degradation of permafrost-derived DOM. From peak discharge in the summer to low 36 flow in late autumn, the DOC concentration at the headstream site decreased from 13.9 37 to 10.2 mg/L, while the 14C-DOC age increased from 745 to 1560 years before present 38 (BP), reflecting an increase in relative contribution of deep permafrost carbon due to 39 the effect of changing hydrological conditions over the course of the summer on DOM 40 source (AL vs. PL). Our study thus demonstrates that hydrological conditions impact 41 the mobilization of permafrost carbon in an alpine fluvial network, the signature of 42 which is quickly lost through in-stream metabolism. 43

44 Keywords: dissolved organic matter; permafrost; Qinghai-Tibet Plateau; FT-ICR MS;





- 45 radiocarbon age
- 46

47 1. INTRODUCTION

48 The amount of carbon stored in permafrost is roughly twice as much as that in the atmosphere and represents the largest component of the terrestrial carbon pool (Zimov 49 50 et al., 2006; Tarnocai et al., 2009). Accelerated climate warming has led to a succession 51 of changes associated with permafrost thaw, where liquid water once frozen in permafrost soils has changed watershed hydrology, topography and ecosystem 52 53 biogeochemistry (Frey and Smith, 2005; Abbott et al., 2015; Vonk et al., 2015). When permafrost-derived carbon enters aquatic systems, it can be rapidly bio- and photo-54 degraded (Cory et al., 2014; Drake et al., 2015; Vonk et al., 2015). Therefore, the 55 mobilization of carbon from permafrost soils where it has been relatively stable for 56 thousands of years into dissolved carbon could increase greenhouse gas emissions 57 (Cory et al., 2013; Vonk et al., 2013; Mann et al., 2015; Ward and Cory, 2016; Selvam 58 et al., 2017) and exacerbate climate warming via a positive feedback loop (Koven et al., 59 2011; Schuur et al., 2015). 60

The seasonal thawing-freezing cycle of permafrost soils could change hydrologic 61 inputs and restrict source water contributions to river flow, leading to variability in the 62 flux and the chemical composition of dissolved organic matter (DOM) in permafrost-63 impacted watersheds (Petrone et al., 2006; Laudon et al., 2011). DOM in the Yukon 64 River exhibits seasonal changes in aromaticity, molecular weight, ¹⁴C age and 65 biodegradability (Striegl et al., 2007; Spencer et al., 2008; Wickland et al., 2012; 66 O'Donnell et al., 2014). Since the persistence of DOM in aquatic systems is related to 67 chemical composition (Kellerman et al., 2015; Kellerman et al., 2018), substituting 68 69 space for time to trace changes in DOM composition along a hydrologic flowpath may





illustrate the environmental behavior and fate of seasonally exported permafrost carbon. 70 The Qinghai-Tibet Plateau (QTP), the world's largest and highest plateau, plays a 71 critical role in the evolution of the Asian Monsoon (Sato and Kimura, 2007; Wu et al., 72 73 2007) and supplies water to several large rivers such as the Yangtze River, Yellow River and Yarlung Tsangpo (Yao et al., 2007; Kang et al., 2010). As a climate sensitive region, 74 75 the QTP has experienced significant warming since the 1950s (Qiu, 2008) with the mean annual air temperature rising at a rate of 0.36 °C per decade from 1961 to 2007 76 (Wang et al., 2008). Consequently, the permafrost soils on the OTP have begun to thaw 77 78 and collapse, causing abundant carbon loss from degradation, leaching and lateral displacement (Mu et al., 2016). However, compared with an abundance of studies on 79 Arctic permafrost, biogeochemical studies on QTP permafrost are scant (Mu et al., 80 81 2016). This results in a limited understanding of the permafrost carbon cycle as a whole since the QTP represents nearly 10% of the global permafrost, what's more, the QTP 82 differs from the Arctic in altitude, climate, and hydrology (Bockheim and Munroe, 83 2014). 84

Here, we conducted a study on the spatial and temporal change of permafrost-derived 85 DOM on the northeastern QTP. We applied multiple analytical techniques including 86 Fourier transform ion cyclotron resonance mass spectrometry (FT-ICR MS), AMS 87 radiocarbon (¹⁴C), and UV-visible optical spectroscopy. Our objective is two-fold: 1) 88 determine the dominant sources of alpine stream DOM on the QTP (active layer (AL) 89 vs. permafrost layer (PL)), and 2) trace the persistence and degradation of permafrost-90 derived DOM in an alpine fluvial network. This work represents the first step in 91 characterizing in-stream removal and transformation processes of permafrost carbon at 92 the molecular level on the QTP. 93





95 2. MATERIALS AND METHODS

96 2.1. Study area and sampling

97 Our study area is located in Gangcha County, north of Qinghai Lake. The climate is 98 typical plateau continental climate, characterized by extensive sunshine duration (~3000 hours per year), long cold and dry winters and short cool and humid summers 99 100 (Peng et al., 2015). During 2013-2016, January had the lowest average monthly 101 temperature (-11.82 °C), while December had the lowest average monthly precipitation (0.3 mm). Meanwhile, the highest average monthly temperature and precipitation 102 103 occurred in July (11.66 °C) and August (124.67 mm), respectively. These climate data are available at http://data.cma.cn. The permafrost soil was developed in the late 104 105 Quaternary, and accumulated > 2 m thick in mountainous areas of the Gangcha County. 106 Due to the rapid climate warming on the QTP, the ice-rich permafrost began to thaw, 107 and several thermo-erosion gullies formed a decade ago. In this study, we focused on a continuous system that starts with a thermo-erosion gully (> 200 m long), forms a 108 stream which flows into Qinghai Lake, the largest lake in China with a surface area of 109 ca. 4500 km². Thawed permafrost slumping exposed soil profiles at the gullies' head 110 (ca. 3850 meters above sea level; masl). The top 60 cm is an active layer (AL) that 111 comprises abundant grass litter and roots, underlain by a dark permafrost layer (PL) 112 without visible plant debris. The thaw depth reached 78 cm in August 2015. Seasonal 113 thaw of the entire AL and the upper PL allows for both vertical and lateral percolation 114 of rainwater, which mobilizes large amounts of particulate and dissolved organic matter. 115 The water in the gully flows southward across the hillslope before draining into Qinghai 116 Lake (3196 masl; Fig. 1). 117

Our fieldwork was conducted in the summer and autumn of 2015 and 2016. In 2015,
a time-series sampling campaign was conducted at the headstream (Q-1) from August





1st when the warm and humid climate caused the largest export of leachates, to October 120 15th when the leaching ceased due to little precipitation and low temperature. The AL 121 and PL leachate samples were collected at a depth of 60 cm and 220 cm, respectively, 122 123 at the gullies' head. For each leachate sample, >15 L water was gathered over 2 days using a 20 L pre-cleaned HDPE carboy. Besides soil leachates, water samples (20 L 124 125 each) were collected from twenty sites along the stream (Fig. 1). Sampling sites Q-1 to 126 Q-10 are located in a first-order stream that originates in the largest thermo-erosion gully, whereas sites Q-11 and Q-12 are located in another first-order stream nearby. 127 128 These two streams merge together to form the main stream, along which sampling sites Q-13 to Q-20 were located. Surface water samples were collected using pre-cleaned 129 HDPE carboys and kept on ice and in the dark until filtering through Whatman GF/F 130 filters (0.7 μ m) within 6 hours after sampling. To obtain enough carbon for ¹⁴C analyses, 131 aliquots of the 0.7 µm filtrate were concentrated over a cross-flow ultrafiltration system 132 with 1 kDa cut off (Millipore®, Pellicon 2 system). The retentates and the remaining 133 filtrate were all stored at -20 °C until further analysis. All glassware and GF/F filters 134 were combusted at 450 °C for at least 4 hours prior to use. 135

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137 2.2. Hydrological condition, DOC concentration and spectral absorbance in alpine 138 streams

On Aug. 1st 2015, stream water temperature, pH and conductivity were measured with a portable Horiba W-23XD Water Quality Monitoring System. The water flux was calculated according to flow rate and cross-sectional area of the stream. The DOC concentration of each water sample was determined by 3-5 injections on a Shimadzu TOC-V_{CPH} analyzer using high temperature combustion, and the coefficient of variance across measurements was < 2%.





The optical properties of the water samples were determined using a Shimadzu UV-145 1800 spectrophotometer. The scan range was between 200 and 800 nm and Milli-Q 146 water (18.2 M Ω cm⁻¹) was used as the blank. A quartz cell with 1.0 cm path length was 147 148 used. The spectral slope of the 275-295 nm region (S275-295), an indicator for the molecular weight of DOM (Helms et al., 2008), was determined by applying log linear 149 150 fits across the wavelengths 275-295 nm. Specific UV absorbance (SUVA254), an indicator for relative aromatic C content, was calculated by dividing the decadic UV 151 absorbance at 254 nm with DOC concentration (Weishaar et al., 2003). 152

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154 2.3. Electrospray ionization Fourier transform ion cyclotron resonance mass 155 spectrometry (ESI FT-ICR MS)

156 Selected water samples collected in 2016 from headstream (Q-1), mid-stream (Q-9), and downstream (Q-17), as well as leachate samples collected from the AL and PL, 157 were prepared for FT-ICR MS analyses. They were solid phase extracted (SPE) using 158 the Bond Elut PPL (Agilent Technologies) following the procedures of Dittmar et al. 159 (2008). The aliquot volume of SPE DOM was adjusted for a target final eluate 160 concentration of 40 µg C/ml (in methanol) to aid ionization in negative mode 161 electrospray ionization (ESI). The methanol extracts were analyzed on a 9.4 Tesla 162 custom-built FT-ICR MS at the National High Magnetic Field Laboratory (NHMFL; 163 Tallahassee, FL; Kaiser et al., 2011). The injection speed was 0.7 µL/min. A total of 100 164 broadband scans was accumulated for each mass spectra. Other instrumental parameters 165 can be found in Hodgkins et al. (2016). After internal calibration in MIDAS Predator 166 Analysis (NHMFL), formulas were assigned based on published rules to peaks with 167 intensities > 60 baseline noise (Stubbins et al., 2010) using EnviroOrg®TM software and 168 169 categorized by compound class based on the elemental composition of molecular





- $\label{eq:spectral} \mbox{formulas} \ (\mbox{Spencer et al.}, 2014; \mbox{Corilo}, 2015). \ \mbox{A modified aromaticity index} \ (\mbox{AI}_{mod}) \ \mbox{was}$
- 171 calculated according to the definition of Koch and Dittmar (2006): $AI_{mod} =$
- 172 $\frac{1+C-0.5O-S-0.5H}{C-0.5O-S-N}$, and if AI_{mod} is negative, then AI=0. The groups referenced in this study
- are: 1) aliphatics (Ali.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, N = 0; 2) peptides (Pep.): H/C 1.5 2.0, O/C < 0.9, O/C < 0.9, O/C < 0.9, O/C < 0.9
- 174 2.0, O/C < 0.9, N > 0; 3) highly unsaturated compounds (Uns.): $AI_{mod} < 0.5$, H/C < 1.5;
- 4) polyphenols (Pol.): $0.5 < AI_{mod} < 0.67$; 5) condensed aromatics (CA): $AI_{mod} \ge 0.67$.
- The relative abundance of the defined compound class was weighted by signalmagnitude in each spectrum.
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179 2.4. Radiocarbon analyses

Freeze-dried retentates were fumigated with concentrated hydrochloric acid (12 M) in order to remove inorganic carbon. After that, the samples were analyzed on the Keck Carbon Cycle Accelerator Mass Spectrometry (AMS) Facility at the University of California, Irvine, USA. Processing blank and sample preparation backgrounds were subtracted. Radiocarbon concentrations are given as conventional ¹⁴C age following Stuiver and Reimer (1993).

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187 **3. RESULTS**

188 3.1. Hydrology and DOC concentration from headstream to downstream water

Discharge increased along the stream reach, from 0.15 m³/min at the headstream (Q-1) on August 1st 2015 to 24.14 m³ /min (Q-19) (Fig. 2). pH increased from 7.4 at Q-1 to 8.2 at Q-4 and remained elevated in the middle and lower stream (7.9 to 8.4). Conductivity was relatively constant from Q-1 to Q-6 (35 to 38 μ s/cm), then increased at Q-7 and remained elevated downstream (48 to 60 μ s/cm). The DOC concentration was high in headstream waters (e.g., 11.7 \pm 0.9 mg/L at Q-1 and 10.2 \pm 1.5 mg/L at Q-





- 195 2; mean \pm SD, same hereafter) and decreased downstream (2.5 to 5.8 mg/L from Q-5
- to Q-20). The mean DOC concentration of the AL leachates (11.6 \pm 1.1 mg/L) was an
- order of magnitude lower than that of the PL leachates ($126.4 \pm 20.9 \text{ mg/L}$).
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199 **3.2.** Optical properties of DOM in leachates and stream waters

The mean $S_{275-295}$ was $(14.5 \pm 0.48) \times 10^{-3}$ nm⁻¹ for the AL leachates and $(18.3 \pm$ 200 1.3) × 10⁻³ nm⁻¹ for the PL leachates. In the stream waters, the S₂₇₅₋₂₉₅ ranged from 15.8× 201 10^{-3} to 22.5×10^{-3} nm⁻¹, increasing in downstream reaches. SUVA₂₅₄ was 3.52 ± 0.24 L 202 mg C⁻¹ m⁻¹ for the AL leachates and 0.95 ± 0.14 L mg C⁻¹ m⁻¹ for the PL leachates, and 203 decreased in the stream from Q-1 to Q-10 (3.06 to 1.27 L mg C⁻¹ m⁻¹), and then remained 204 low (Fig. 3). A strong negative correlation was found between SUVA254 and S275-295 for 205 water samples from both years ($R^2 = 0.77$, P < 0.01). Neither stream waters nor 206 permafrost leachates show an interannual variation of optical properties (Fig. 3). 207

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3.3. Spatiotemporal change of ¹⁴C-DOC age through fluvial networks

¹⁴C-DOC age of the PL leachate was 4145 years BP, which was much older than that of the AL leachate (535 years BP; Fig. 4a). The ¹⁴C-DOC age decreased along the stream from 745 years BP for the headstream water (Q-1) to 160 years BP at Q-19, a site close to Qinghai Lake. Besides apparent spatial variability, the ¹⁴C-DOC age also changed temporally. In 2015, the ¹⁴C-DOC age of the headstream water (Q-1) increased from 745 years BP on August 1st, to 1015 years BP on August 11th and 1560 years BP on September 5th (Fig. 4b).

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218 3.4. FT-ICR MS characterization of SPE-DOM

219 Compared with the PL leachate, the AL leachate was characterized by higher





220	molecular richness (14709 vs. 9645 assigned formulae), higher mean molecular weight
221	(498.81 vs. 452.73 Da) and higher AI _{mod} (0.47 vs. 0.30). Elemental composition
222	revealed that compounds containing both N and S were only detected in the AL
223	leachates and headstream waters. The AL leachate contained 54.28% highly unsaturated
224	compounds, 27.10% polyphenols and 17.23% condensed aromatic compounds,
225	whereas the proportion of aliphatics and peptides was minor (ca. 1.30%). Compared
226	with the AL leachate, the PL leachate comprised a higher proportion of polyunsaturated
227	compounds (74.23%) and aliphatics + peptides (10.04%), but a lower proportion of
228	polyphenols (11.33%) and condensed aromatics (4.32%).

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Along the stream (Q-1, Q-9, and Q-17), the molecular richness, mean molecular weight and modified aromaticity index of SPE-DOM decreased by 26% (14924 to 11074), 4.7% (510.1 to 486.5 Da), and 16.3% (0.43 to 0.36), respectively (Table 1). The relative abundance of aromatics (condensed aromatics and polyphenols) decreased by 48% (35.7% at Q-1 vs. 18.4% at Q-17), whereas that of highly unsaturated compounds increased by 28% (62.8% at Q-1 vs. 80.3% at Q-17). Aliphatics and peptides were minor components of stream DOM (<1.3%) and did not exhibit a downstream trend.

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238 4. DISSCUSSION

239 4.1. AL leachates as a major source of stream DOM

The UV-visible optical parameters and molecular formulas resolved by FT-ICR MS show that the AL and PL leachates have different chemical compositions (Table 1 and 2). Since chemical composition impacts the reactivity of DOM (Kellerman et al., 2015), the differing chemical composition between the AL and PL leachates that enter the stream may influence bioavailability (Vonk et al., 2013) and photolability (Stubbins et al., 2017). Thus, distinguishing DOM source is crucial for understanding in-stream





biogeochemical processes in permafrost-impacted systems. DOM may originate from 246 a variety of sources including permafrost soil (AL and PL) leaching, in-situ microbial 247 production, and wet deposition from snow and rain. At the headstream site (Q-1), 248 249 however, the dominant source of DOM is permafrost soil leaching, as short residence times at the gully head restrict in-stream production and wet deposition is likely 250 251 negligible due to low DOC concentrations in Tibetan glaciers (0.2-3.3 µg/ml; Spencer 252 et al., 2014). Assuming that headstream DOM is derived only from permafrost soil leaching, we are able to estimate the relative contributions of DOM from the AL and 253 254 PL.

The DOC concentration of the AL leachate (ca. $11.6 \pm 1.1 \text{ mg/L}$; mean \pm SD based 255 on samples from 2015 and 2016, n = 2; same hereafter) is similar to that of the 256 headstream (Q-1; ca. 11.7 \pm 0.9 mg/L), but substantially lower than that of the PL 257 leachates (ca. 126.4 ± 20.9 mg/L), supporting a predominance of AL-leachate DOM in 258 stream waters. In addition, the SUVA₂₅₄ is 3.51 ± 0.24 L mg C⁻¹ m⁻¹ for AL leachates 259 (and 0.95 \pm 0.14 L mg C⁻¹ m⁻¹ for PL leachates, whereas the S₂₇₅₋₂₉₅ is (14.5 \pm 0.48) \times 260 10^{-3} nm⁻¹ for AL leachates and $(18.0 \pm 1.33) \times 10^{-3}$ nm⁻¹ for PL leachates. Similar optical 261 properties and DOC concentrations between AL-leachates and the headstream water 262 $(16.5 \pm 0.40 \times 10^{-3} \text{ nm}^{-1} \text{ for } S_{275\text{-}295} \text{ and } 2.92 \pm 0.19 \text{ L mg C}^{-1} \text{ m}^{-1} \text{ for SUVA}_{254})$ support 263 DOM that leaches from the AL dominates stream DOM. Furthermore, the stream water 264 at Q-1 has a ¹⁴C-DOC age of 745 years BP, close to that of the AL leachate (535 years 265 BP), and much younger than that of the PL leachate (4145 years BP). We estimate the 266 portion of AL and PL-derived C by using a binary mixing model based on Δ^{14} C values 267 of bulk DOC (Criss, 1999): 268

269
$$\Delta^{14}C_{DOM} = f_{AL} \times \Delta^{14}C_{AL} + f_{PL} \times \Delta^{14}C_{PL}$$

270
$$1.0 = f_{AL} + f_{PL}$$





According to this model, ca. 94% of DOC collected from stream site Q-1 on Aug. 271 1, 2015 originated from the AL (Fig. 6a). Headstream ¹⁴C-DOC age increased from 272 summer to fall (Fig. 4b), reflecting an enhanced contribution of old carbon from the 273 274 deeper soils (i.e., PL), however, the AL still accounted for \geq 72% of total DOC exported (Fig. 6a). This binary mixing model may overestimate the contribution of AL to stream 275 276 DOC since PL-derived DOC may be degraded faster than AL-derived DOC, due to the 277 high biolability of ancient permafrost carbon as shown in Arctic ecosystems (Vonk et 278 al., 2013). Nonetheless, the AL appears as a major contributor to stream DOC in the 279 QTP.

Seasonal variation of ¹⁴C-DOC (Fig. 4b) has been previously observed in high 280 latitude permafrost areas in Alaska (Aiken et al., 2014; O'Donnell et al., 2014), with the 281 most enriched ¹⁴C values observed in the spring and becoming more depleted through 282 summer-fall and/or during winter. The mean monthly air temperature of Gangcha 283 County, after reaching the maximum in July (ca. 10.5 °C), decreases to 2.1 °C in 284 September (data from http://data.cma.cn). As air temperature drops, surface soils freeze 285 earlier than deeper soils, leading to an increase in the relative contribution of deep soil 286 carbon (i.e. PL) to stream DOM, although the DOC concentration in Q-1 decreased 287 from 13.9 mg/L to 10.2 mg/L (Fig. 6b). 288

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290 4.2. Selective removal of DOM along the alpine stream on the QTP

The DOC concentration decreased (12.3 to 4.0 mg/L) from upper to mid-stream (Q-1 to Q-5), which could be attributed to a dilution effect and/or in-stream degradation of DOM. Dilution from groundwater is likely since groundwater discharge sustains baseflow of rivers and streams in the QTP (Ge et al., 2008). Downstream groundwater inputs were further supported by the order of magnitude increase in discharge (1.49 to





24.14 m³/min) and increase in conductivity (37 to 60 µs/cm). Moreover, downstream 296 DOC concentrations remained about 3.0-4.0 mg/L (Q-15 to Q-20), indicative of the low 297 DOC concentrations of groundwater. Conversely, a tributary that originated from 298 299 another thermo-erosion gully merged into the study stream, however, the different tributaries exhibited similar DOC concentrations (e.g., Q-9 and Q-10 vs. Q-11 and Q-300 301 12; Fig. 2d). The similarities in DOC concentrations were attributed to homogeneity in 302 dominant vegetation, soil type and climate, and thus, homogeneity in DOM inputs to the different tributaries in our study area. Therefore, additional tributaries could not 303 304 explain the spatial pattern of DOC concentration.

Despite evident dilution, DOC attenuation could be partly due to in-stream 305 degradation given several lines of evidence from optical properties, radiocarbon age 306 307 and molecular composition. The UV-visible optical parameters, S275-295 and SUVA254, have been widely used to reveal mean molecular weight and aromaticity of DOM, 308 respectively (Weishaar et al., 2003; Helms et al., 2008; Spencer et al., 2009; Mann et 309 al., 2012). In our study, the $S_{275-295}$ of stream waters varied from 15.8×10^{-3} to $22.5 \times$ 310 10^{-3} nm⁻¹ (Fig. 3a), comparable to typical riverine DOM values (13.19×10^{-3} to 22.96 311 $\times 10^{-3}$ nm⁻¹), but much lower than that of DOM from continental shelf and slope (29.7) 312 $\times 10^{-3}$ to 48.5×10^{-3} nm⁻¹) (Fichot and Benner, 2012), suggesting a moderate degradation 313 of stream DOM on the QTP. A downstream increase for S275-295 regardless of sampling 314 time (Fig. 3a) reflects selected degradation of high molecular weight compounds, 315 leading to the enrichment of low molecular weight DOM. This spatial trend is in 316 accordance with the size-reactivity continuum model (Amon and Benner, 1996) that the 317 bioreactivity of DOM decreases along a continuum of size (from large to small). In 318 addition to S₂₇₅₋₂₉₅, SUVA₂₅₄ varied from 1.27 to 3.06 L mg C⁻¹ m⁻¹, showing a general 319 320 decrease downstream (Fig. 3b). Lignin, an aromatic biopolymer specific for vascular





plants (Hedges et al., 1997), is relatively resistant to biodegradation (Hedges et al., 321 1985), but highly photo-labile (Lanzalunga and Bietti, 2000). Cory et al. (2014) found 322 323 that sunlight accounts for 70% to 95% of water column carbon processing in Arctic 324 rivers and lakes. Given strong solar radiation and long sunshine duration (~3000 hours per year) on the QTP (Peng et al., 2015), photo-degradation could be an important 325 pathway for carbon removal in QTP streams. A strong negative correlation between 326 $S_{275-295}$ and SUVA₂₅₄ (R² = 0.73, p < 0.01) indicates that photodegradation of high 327 328 molecular weight aromatic compounds (like lignin) may play a role in the decrease of 329 mean molecular weight of DOM along the stream.

Similar to SUVA254 and S275-295, the data from FT-ICR MS also show a downstream 330 331 decrease in aromaticity (AI_{mod}: 0.43 to 0.36) and mean molecular weight of stream DOM (510.0 to 486.5 Da; Table 1). Compared with headstream DOM at Q-1, DOM at 332 Q-9 and Q-17 was characterized by a lower proportion of condensed aromatics and 333 polyphenols and enriched in highly unsaturated compounds (Table 1). The decrease in 334 relative abundance of aromatic compounds is consistent with the reports for the 335 photolability of aromatic formulas within permafrost, river and ocean DOM (Stubbins 336 and Dittmar, 2015; Stubbins et al., 2017). 337

As discussed in section 4.1, AL is the principal contributor to stream DOM. Thus, 338 tracing AL-derived DOM is paramount in estimating biogeochemical processes of 339 carbon in the stream. FT-ICR MS identified 6409 molecular formulas specific to AL-340 leachates (i.e. not observed in the PL, Table 2). Through various stream processes, some 341 AL specific formulas were removed from the DOM pool (from 17% by Q-1 up to 59% 342 by Q-17), which accounted for 66% of the aromatic compounds and 51% of the highly 343 unsaturated compounds (Table 2). Molecular formulas containing N and/or S were more 344 345 labile in the fluvial networks than CHO formulas, with 84% of S-containing formulas





and 100% of S and N-containing formulas lost (Table 2). Furthermore, the removal of 346 DOM formulas (ca. 83% of AL-specific formulas, and >95% of AL-specific formulas) 347 occurred in upper and mid-stream (leachates to Q-9). Concurrent with the rapid loss of 348 349 AL-specific formulas, some new molecular formulas were detected by FT-ICR MS, which was mainly attributed to in-situ production by stream algae/microbes, although 350 351 an import form groundwater could not be excluded. The addition of those new molecular formulas was also reflected by the ¹⁴C enrichment in middle and lower-352 stream (Fig. 3b). 353

354 Overall, our multiple analyses demonstrate a rapid and selective degradation of stream DOM on the QTP. The attenuation of aromatic compounds and enrichment of 355 highly unsaturated compounds could change the environmental photo- and bio-lability 356 357 of DOM, increasing relative importance of photodegradation in upper stream and biodegradation in lower stream. The continuous change in chemical properties of DOM 358 along the alpine stream flowpath has a potential to shift the aquatic microbial 359 community since DOM serves as an important energy and nutrient source(Wild et al., 360 2014). 361

362

363 4.3. Prediction of in stream carbon dynamic under continued warming

The DOC concentrations, UV-visible optical parameters and FT-ICR MS all suggest that currently, PL is a minor source to stream DOM (see 4.1). However, the QTP is a sensitive area to global warming, with a rate of air temperature rise that is approximately three times the global warming rate (Qiu, 2008). According to climate model predictions, spatial average temperatures of the QTP will increase by 0.68–0.98 °C for the period of 2015–2050 (Zhu et al., 2013), and in 2050, the mean AL thickness on the QTP permafrost will increase by approximately 0.3-0.8 m more than that in 2010





(Zhang and Wu, 2012). With the deepening of the AL, carbon that is currently stable in 371 the PL will be thawed and mobilized into the downslope aquatic environments, which 372 373 inevitably changes the proportion of AL vs. PL contributions to stream DOM. Thus, it 374 is worth to trace chemical change of PL leachates along the stream. The PL leachate contained only 1345 formulas unique to the PL leachate in comparison to the AL, 375 376 accounting for 14% of total assigned formulas (Table 2). Most PL-specific formulas 377 were more biolabile components, e.g. aliphatics and peptides (73%), followed by highly unsaturated formulas (23.6%) and aromatics (1.9%). At the downstream site (i.e., Q-378 379 17), 90% of these PL-specific molecular formulas were lost, substantially higher than that of AL-specific formulas (59%). Furthermore, the vast majority of PL-specific 380 formulas were lost within < 1 km (Q-1: 83%) whereas only 17% of AL-specific 381 382 formulas were lost by Q-1 (Table 2). Therefore, the FT-ICR MS data demonstrate that permafrost thaw can trigger a rapid degradation of old carbon that was frozen in soils 383 for thousands of years (Fig. 3a). This is consistent with findings in Arctic fluvial 384 networks that show the utilization of ancient permafrost carbon in headstream waters 385 was rapid (Mann et al., 2015; Frey et al., 2016). Therefore, we hypothesize that with 386 enhanced leaching of deep soil C under continued warming on the QTP, DOM in alpine 387 streams will be more enriched in biolabile aliphatics/peptides and depleted in 388 photolabile aromatics 389

Finally, despite substantial in-stream degradation, some old permafrost-derived carbon (i.e., polyphenols and highly unsaturated compounds) could persist downstream. In addition, CO₂ produced by respiration of old DOC could be utilized by stream algae to biosynthesize new DOM with an old carbon age. These effects resulted in a sustained deviation from modern ¹⁴C-DOC age in the alpine stream (e.g., 160 years BP at Q-19), and were even detected in large rivers on the QTP (e.g., Yangtze River and Yellow River;





- Qu et al., 2017). Thus, under continued warming, a greater quantity of older C may be
 transported into large watersheds on the QTP, and thereby exert an important role in
 biogeochemical cycles there since older carbon has different photo and bio-lability from
 young carbon in AL soils.
- 400

401 5. CONCLUSIONS

Permafrost thaw represents positive feedbacks to climate change, but its carbon alteration and removal mechanism is not well known, particularly for the alpine permafrost such as the QTP. Here we use multiple analytical methods (e.g., FT-ICR MS, radiocarbon and UV-visible spectroscopy) to trace spatial and temporal variability of permafrost DOM along an alpine stream in the northeastern QTP, from which four conclusions have been reached.

1) Presently, the AL is the major source to stream DOM with relatively high aromaticity. This character, combined with strong solar radiation on the QTP, suggests sunlight may be an important driver for DOM removal in alpine fluvial networks, which was corroborated by an almost 60% loss of AL specific formulas from the thermoerosion gully head to downstream waters.

2) From summer to fall (seasonal permafrost thawing to freezing cycle), the
concentrations and chemical composition of stream DOM varied significantly at the
thermo-erosion gully head. Even though the total amount of the leached DOC decreased,
the contribution of deep permafrost carbon increased, reflected by an increase of ¹⁴CDOC age and a decrease in aromaticity of DOM.

Although both the AL and PL leachate DOM underwent rapid degradation in
the alpine stream, some components with old ¹⁴C-DOC age (mainly highly unsaturated)
were recalcitrant to degradation and could be transported downstream, causing ¹⁴C-





- 421 DOC values that were more depleted than modern radiocarbon age downstream in our
- 422 study, and even in large watersheds as observed in Qu et al. (2017).
- 423 4) With deepening of the AL under continued climate warming on the QTP, 424 currently stable PL soils will thaw and release greater amounts of old, aliphatic/peptide-425 rich DOM to downstream waters. This change in source and chemical composition will 426 make microbial degradation more important for carbon removal and may shift 427 downstream microbial communities, even in large watershed systems. All these factors 428 should be taken into account when interpreting alpine permafrost carbon dynamics 429 under the amplified climate warming trend on the QTP.

430

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617 Figure and table captions

Fig. 1. Location of the QTP and sampling sites Q1 to Q20. Sites marked by a star were
selected for FT-ICR MS and ¹⁴C-DOC analyses. The AL and PL denote the sampling
locations of the active and permafrost layers. The blue line and the red line represent
the first order and second order stream, respectively, and the blue dashed line denotes

- 622 stream without GPS data.
- **Fig. 2.** (a) Stream water discharge, (b) pH, and (c) conductivity at the sampling sites in 2015; and (d) DOC concentration in stream water and PL/AL leachates collected in
- 625 2015 (filled circles) and 2016 (open circles).

Fig. 3. UV-visible optical indices of the stream water and PL/AL leachate samples collected in 2015 (filled circles) and 2016 (open circles) on the QTP: $S_{275-295}$ (a) and SUVA₂₅₄ (b).

Fig. 4. Variations of ¹⁴C-DOC age across the alpine stream spatially (a), and at
 headstream Q-1 temporally (b).

Fig. 5. van Krevelen diagrams of AL leachate DOM (a), PL leachate DOM (b),
headstream DOM Q-1 (c), downstream DOM Q-17 (d), the relative abundance of
defined compound class in different samples (e). CA = condensed aromatics, Pol. =
polyphenols, Uns. = highly unsaturated compounds, Ali. = aliphatics, Pep. = peptides;
and Sug. = Sugar.

Fig. 6. (a) Relative contribution of AL leachate DOM to headstream DOM (Q-1); and (b) temporal variation of the DOC concentration at headstream Q-1.

Table 1 The number of molecular formulas assigned, modified aromaticity index
(AI_{mod}), mean molecular weight (mean MW) and relative abundance of defined
compound classes detected by FT-ICR MS for DOM samples from the QTP, including
soil leachates (AL and PL) and stream waters (Q-1, Q-9 and Q-17). CA = condensed
aromatics, Pol. = polyphenols, Uns. = highly unsaturated compounds, Ali. = aliphatic,
Pep. = peptides.

Table 2 The number of specific molecules identified in the AL leachate DOM and the
PL leachate DOM within the fluvial network, and the percentage of peaks totally
degraded during the transportation.





Fig. 1





































Table 1. The number of molecular formulas assigned, modified aromaticity index
(AI_{mod}), mean molecular weight (mean MW) and relative abundance of defined
compound classes detected by FT-ICR MS for DOM samples from the QTP, including
soil leachates (AL and PL) and stream waters (Q-1, Q-9 and Q-17). CA = condensed
aromatics, Pol. = polyphenols, Uns. = highly unsaturated compounds, Ali. = aliphatic,
Pep. = peptides.

Sample	Formulas assigned	Mean MW	$\mathrm{AI}_{\mathrm{mod}}$	CA (%)	Pol. (%)	Uns. (%)	Ali. (%)	Pep. (%)
AL	14709	498.81	0.47	17.23	27.10	54.28	1.16	0.14
PL	9645	452.73	0.30	4.32	11.33	74.23	7.92	2.12
Q-1	14924	510.07	0.43	12.05	23.69	62.85	1.14	0.14
Q-9	11724	500.19	0.38	6.86	15.82	76.32	0.86	0.06
Q-17	11074	486.50	0.36	5.53	12.91	80.31	1.11	0.08





Samples		All peaks	Only CHO	Contains N	Contains S	Contains N& S	Condensed aromatics	Polyhoenols	Unsatuated	Aliphatics	Peptides
	AL	6409	1793	3370	424	822	1620	1720	2970	38	23
AL	Q-1	5311 (17%)	1653 (8%)	2791 (17%)	349 (18%)	517 (37%)	1278 (21%)	1416 (18%)	2549 (14%)	20 (47%)	14 (39%)
specific	6-9	3365 (47%)	1294 (28%)	1917 (43%)	153 (64%)	0(100%)	748 (54%)	838 (51%)	1759 (41%)	6 (84%)	1 (96%)
	Q-17	2623 (59%)	985 (45%)	1570 (53%)	67 (84%)	0~(100%)	550 (66%)	602 (65%)	1453 (51%)	5 (87%)	0 (100%)
	PL	1345	515	551	278	0	2	23	318	597	385
PL	Q-1	222 (83%)	90 (83%)	102 (81%)	30 (89%)	0	0 (100%)	11 (52%)	126 (60%)	46 (92%)	36 (91%)
specific	Q-9	117 (91%)	44 (91%)	46 (92%)	27 (90%)	0	2 (0%)	14 (39%)	96 (70%)	1(100%)	4 (99%)
	Q-17	130 (90%)	47 (91%)	55 (90%)	28 (90%)	0	2 (0%)	13 (43%)	104 (67%)	(%66) 9	5 (99%)

Table 2: The number of specific molecules identified in the AL leachate DOM and the PL leachate DOM within the fluvial network, and the