Editor Comments

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Author Response: We sincerely thank the Editor for her additional comments/edits. We have added statements in **blue** below that detail our response to each comment.

Although you might have not found studies on DOM precipitation (or flocculation might be a better term) specifically related to systems with thaw slumps, I think you will on this process in other systems (try boreal, if not temperate, if not any humic aquatic system on the planet). There is no reason for excluding this loss process. See for example this study on lakes: Wachenfeldt, E. von, D. Bastviken, and L. J. Tranvika (2009), Microbially induced flocculation of allochthonous dissolved organic carbon in lakes. Limnol. Oceanogr. 54: 1811–1818, doi:10.4319/lo.2009.54.5.1811.

This is an excellent suggestion and reference and we have added the following to this section: "(ii) flocculation of terrestrial DOM resulting in the settling of particulate organic matter (Wachenfeldt et al., 2009)"

You wrote: "to the point where A350 at a 1 cm path length was ≤ 0.02 " but I think it needs to be replaced by "to the point where A350 was ≤ 0.02 for a 1 cm path length" This change has been made.

You wrote: "In other words, the more abundant "virgin" bioavailable molecules upstream are replaced downstream by photobleached smaller molecules (originating from aromatic compounds), resulting in the fraction of DOC used relatively constant without any clear pattern overall. If this (or something similar) were not the case, we would expect to see a declining fraction of bioavailable DOC along the flow-path continuum."

I was using the term "virgin" quoted loosely as it's jargon, but you probably need to find a more accurate expression: never exposed, fresh, or something like that. Moreover, I think you need to tone down the sentence, as you do not have direct evidence of this replacement; your results suggest this (use words like may, could, possibly, suggest, etc.). And I don't think "(or something similar)" is necessary. This is an excellent suggestion to improve this section. We have replaced these sentences to now read the following: "In other words, our results suggest that more abundant newly exposed bioavailable molecules upstream are replaced downstream by photobleached smaller molecules (originating from aromatic compounds), resulting in the fraction of DOC used relatively constant without any clear pattern overall. If this were not the case, we would expect to see a declining fraction of bioavailable DOC along the flow-path continuum."

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6	Optical properties and bioavailability of dissolved organic matter along a flow-path
7	continuum from soil pore waters to the Kolyma River mainstem, East Siberia
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25	Keywords: East Siberia, Kolyma River, permatrost, DOC, CDOM, biolability
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30 Abstract

The Kolyma River in Northeast Siberia is among the six largest arctic rivers and drains a region 31 underlain by vast deposits of Holocene-aged peat and Pleistocene-aged loess known as yedoma, 32 most of which is currently stored in ice-rich permafrost throughout the region. These peat and 33 yedoma deposits are important sources of dissolved organic matter (DOM) to inland waters that 34 35 in turn play a significant role in the transport and ultimate remineralization of organic carbon to CO_2 and CH_4 along the terrestrial flow-path continuum. The turnover and fate of terrigenous 36 37 DOM during offshore transport largely depends upon the composition and amount of carbon 38 released to inland and coastal waters. Here, we measured the ultraviolet-visible optical properties of chromophoric DOM (CDOM) from a geographically extensive collection of waters 39 spanning soil pore waters, streams, rivers, and the Kolyma River mainstem throughout a ~250 40 km transect of the northern Kolyma River basin. During the period of study, CDOM absorption 41 42 coefficients were found to be robust proxies for the concentration of DOM, whereas additional 43 CDOM parameters such as spectral slopes (S) were found to be useful indicators of DOM quality along the flow-path. In particular, the spectral slope ratio (S_R) of CDOM demonstrated 44 statistically significant differences between all four water types and tracked changes in the 45 46 concentration of bioavailable DOC, suggesting that this parameter may be suitable for clearly discriminating shifts in organic matter characteristics among water types along the full flow-path 47 48 continuum across this landscape. However, despite our observations of downstream shifts in 49 DOM composition, we found a relatively constant proportion of DOC that was bioavailable (~3-6% of total DOC) regardless of relative water residence time along the flow-path. This may be a 50 51 consequence of two potential scenarios allowing for continual processing of organic material 52 within the system, namely: (a) aquatic microorganisms are acclimating to a downstream shift in

DOM composition; and/or (b) photodegradation is continually generating labile DOM for 53 continued microbial processing of DOM along the flow-path continuum. Without such 54 processes, we would otherwise expect to see a declining fraction of bioavailable DOC 55 downstream with increasing residence time of water in the system. With ongoing and future 56 permafrost degradation, peat and yedoma deposits throughout the Northeast Siberian region will 57 58 become more hydrologically active, providing greater amounts of DOM to fluvial networks and 59 ultimately to the Arctic Ocean. The ability to rapidly and comprehensively monitor shifts in the quantity and quality of DOM across the landscape is therefore critical for understanding potential 60 61 future feedbacks within the arctic carbon cycle.

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63 **1. Introduction**

There is increasing evidence that inland freshwater ecosystems play a significant role in 64 the global carbon cycle owing to the metabolism of terrestrially-derived organic matter as it 65 66 moves through fluvial networks from land to ocean (Cole et al., 2007; Battin et al., 2009a, b). Recent research suggests that arctic watersheds may increasingly augment the role of freshwater 67 ecosystems in the global flux of terrestrial carbon to the atmosphere (Walter et al., 2007; Denfeld 68 69 et al., 2013; Vonk et al., 2013; Hayes et al., 2014; Spencer et al., 2015) and ocean (Frey and Smith, 2005; Frey and McClelland, 2009; Schreiner et al., 2014; Tesi et al., 2014) as a result of 70 climate warming and changing regional hydrology. Terrestrial sources of organic matter 71 72 generally dominate the energy and carbon fluxes through stream, riverine, and estuarine ecosystems (Mulholland, 1997; Holmes et al., 2008), but the lability and composition of this 73 74 carbon remain poorly characterized. Headwater and intermediate streams dominate overall 75 channel length in large dendritic drainage basins (e.g., Denfeld et al., 2013), thus the functional

role of streams and intermediate rivers is magnified when assessing landscape controls on carbonand nutrient fluxes to the atmosphere and Arctic Ocean.

Following the publication of the "river continuum concept" (Vannote et al., 1980), there 78 has been much research focused on the delivery and processing of terrestrially-derived organic 79 matter within temperate stream ecosystems. Through these studies, it has been shown that 80 81 biological processes within streams alter the transport of organic matter to downstream ecosystems (e.g., Webster and Meyer, 1997), but the fate of terrestrial organic matter in arctic 82 83 streams and rivers has only more recently been explored (e.g., Frey and Smith, 2005; Neff et al., 84 2006; Holmes et al., 2008; Denfeld et al., 2013; Spencer et al., 2015). Furthermore, a variety of conceptual and pragmatic issues complicate the study of arctic rivers, including: (i) large 85 seasonal variations in discharge accompanied by large seasonal variations in nutrient and organic 86 matter inputs from rivers to the coastal ocean (e.g., McClelland et al., 2012); (ii) the 87 heterogeneity of vegetation, permafrost extent, topography, and soil attributes within arctic 88 89 watersheds (e.g., Frey and McClelland, 2009); and (iii) spatial and temporal inaccessibility hindering comprehensive sampling; among others. 90

Hydrologic flow-paths and organic matter transport in arctic regions dominated by 91 92 permafrost are markedly different than temperate regions with well-drained soils. In particular, 93 permafrost-dominated watersheds lack deep groundwater flow-paths owing to the permafrost 94 boundary in soil that prevents deep groundwater movement (Judd and Kling, 2002; Frey et al., 2007). As a result, the delivery of terrestrial-permafrost organic matter to aquatic ecosystems 95 may in fact lack significant terrestrial or groundwater processing. Once dissolved organic matter 96 97 (DOM) enters aquatic ecosystems, multiple processes remove DOM from the water column: (i) 98 photochemical reactions, where DOM is degraded to CO_2 or to compounds bioavailable for

99	bacterial uptake (Moran and Zepp, 1997; Laurion and Mladenov, 2013; Cory et al., 2014); (ii)
100	flocculation of terrestrial DOM resulting in the settling of particulate organic matter
101	(Wachenfeldt et al., 2009); (iii) loss via aggregation of DOM owing to changes in ionic strength
102	when freshwater mixes with sea water (Sholkovitz, 1976); (iiiiv) DOM sorption to particles and
103	sedimentation (Chin et al., 1998); and/or ($\frac{1}{2}v$) bacterial uptake and utilization of the bioavailable
104	fraction (Bronk, 2002; Karl and Björkman, 2002; Mann et al., 2014; Spencer et al., 2015).
105	Measurements of waters along a hydrologic flow-path may indeed give insight into the
106	characteristics of DOM as it is modified through these various processes along the soil-stream-
107	river continuum.
108	Recent work on the Kolyma River in Northeast Siberia has identified marked variation in
109	annual discharge that is associated with large pulses of organic matter flux to the Arctic Ocean
110	during spring freshet, providing detailed temporal characterization of DOM in the Kolyma River
111	mainstem across the annual hydrograph (e.g., Mann et al., 2012). Furthermore, selective
112	processing and loss of permafrost-derived DOM has been shown to occur via microbial
113	metabolism throughout the Kolyma River basin, as waters move downstream through the fluvial
114	network (Mann et al., 2014; Mann et al., 2015; Spencer et al., 2015). Here, we complement
115	these previous studies by providing extensive spatial characterization of DOM along a flow-path
116	continuum from soil pore waters to the Kolyma River mainstem during mid-summer (July)
117	baseflow. The heterogeneity of environmental characteristics and extensive continuous
118	permafrost of the Kolyma River basin combine to make this a critical region to investigate and
119	monitor. In particular, we measured the ultraviolet-visible absorption spectra (200-800 nm) of
120	chromophoric DOM (CDOM) from a geographically extensive collection of waters throughout a
121	~250 km transect of the northern Kolyma River basin, including samples of soil pore waters,

122 streams, rivers, and the Kolyma River mainstem. CDOM absorption and spectral slopes (calculated within log-transformed absorption spectra) were used to investigate contrasting water 123 types and were found to be useful indicators of both the concentration and reactivity of DOM. 124 With ongoing permafrost degradation and subsequent release of a long-term storehouse of 125 organic material into the contemporary carbon cycle, the ability to easily and comprehensively 126 127 monitor the quantity and quality of DOM across the landscape through investigation of its optical properties is becoming critical for understanding the global significance of the arctic carbon 128 129 cycle. Here, we explore a full suite of CDOM parameters as well as concentrations of dissolved 130 organic carbon (DOC) and bioavailable DOC as they vary across a full flow-path continuum in the Kolyma River basin in Northeast Siberia. 131

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133 **2. Data and Methods**

The Kolyma River in Northeast Siberia is among the six largest arctic rivers and drains a 134 ~650,000 km² region underlain by vast deposits of Holocene-aged peat and Pleistocene-aged 135 loess known as yedoma, much of which is currently stored in ice-rich permafrost throughout the 136 region (Holmes et al., 2012; Holmes et al., 2013). These peats and yedoma deposits are 137 138 important sources of DOM to terrestrial waters that in turn play a significant role in the transport and ultimate remineralization of organic carbon to atmospheric CO₂ and CH₄ (e.g., Walter et al., 139 2006; Mann et al., 2012; Denfeld et al., 2013; Spencer et al., 2015). The Kolyma River basin 140 141 and its subwatersheds exhibit extreme hydrologic seasonality, with ice breakup and peak river discharge typically occurring in late May or early June. In this study, sampling took place along 142 143 the most northern ~250 km of the Kolyma River in the vicinity of Cherskiy, Sakha Republic, 144 Russia (68.767°N, 161.333°E) during the mid-summer period of July 2009 (Figure 1). Samples

145 were collected over a narrow temporal window from July 11-25, 2009 in order to capture a "snapshot" of observations during the mid-summer period. In total, 47 water samples were 146 collected, including soil pore waters in shallow wetlands (n=9), small streams with watersheds 147 $<100 \text{ km}^2$ (n=15), major river tributaries with watersheds 900–120,000 km² (n=14), and Kolvma 148 mainstem locations with watersheds >400,000 km^2 (n=9). Although we did not determine 149 residence times directly for our sampled sites, Vonk et al. (2013) estimated that in higher relief 150 areas near Duvannyi Yar (adjacent to the Kolyma River mainstem), the transport time from 151 152 permafrost thaw to entry into the Kolyma River may be less than one hour. Furthermore, with 153 respect to the mainstem, it has been estimated that water residence times in the Kolyma River from Duvannyi Yar to the river mouth may be ~3–7 days, assuming average mainstem velocities 154 of 0.5–1.5 m/s (Holmes et al., 2012; Vonk et al., 2013). As such, permafrost-derived C may not 155 156 be easily detectable at the river mouth, as this time is likely comparable to the rapid removal rates of highly labile permafrost C determined through incubation experiments (e.g., Holmes et 157 al., 2012; Vonk et al., 2013). 158

Samples were collected by hand using a 1 L acid-washed high density polyethylene 159 (HDPE) bottle as a collection vessel, where sample waters were used to rinse the bottle several 160 161 times before filling. Soil pore waters were collected by depressing the soil surface within the wetlands and allowing the water to slowly seep into the collection vessel. In shallow streams, 162 less than 0.5 m in depth, samples were collected approximately midway below the surface and 163 164 the bottom. In larger tributaries and rivers, samples were collected at a depth of ~0.5 m. Water 165 samples were then filtered through precombusted (450°C for 6 hours) Whatman 0.7 µm GF/F 166 filters in the field and stored in acid-washed HDPE bottles without headspace to minimize degassing and algal growth. Upon returning to the laboratory (typically within ~1 day), DOC 167

samples were acidified with concentrated HCl to a pH of ≤ 2 and stored refrigerated and in the dark until analysis via high-temperature combustion using a Shimadzu TOC-VCPH Analyzer (within one month of collection). DOC was calculated as the mean of 3 to 5 injections with a coefficient of variance less than 2%.

We additionally conducted a series of organic matter bioavailability assays to assess the 172 173 total and relative amounts of bioavailable DOC in soil, stream, and river environments. These assays relied upon 5-day biological oxygen demand (BOD) incubations, with methods similar to 174 175 those in Mann et al. (2014). Water samples were collected in triplicate glass 300 mL BOD 176 bottles and filtered as DOC (above). The samples were initially allowed to equilibrate via filtering in a controlled laboratory environment at 15°C, after which t=0 was the start time of the 177 incubations. The Winkler titration method was used to measure dissolved oxygen (DO) 178 concentrations initially (t=0) (i.e., in situ DO) as well as after 5-day incubations at 15° C, where 179 bottles were kept in the dark in between measurements. At t=0, DO measurements were at 180 concentrations expected at equilibrium with the 15°C laboratory temperature (~8.5–9.0 mg/L). 181 This temperature was only slightly warmer than environmental sampling conditions (i.e., the 182 Kolyma River mainstem samples ranged from 11.40–13.90°C, river samples ranged from 10.70– 183 184 14.20°C, and stream samples ranged from 4.40–13.80°C). However, we maintained samples at 15°C as is standard in the BOD method, which allowed samples to be treated identically in the 185 186 controlled experiment (in situ temperatures varied depending not only upon location but also 187 date and time of day). Furthermore, bottles were wrapped tightly with paraffin such that physical degassing should have been minimal during the incubations. BOD was then calculated 188 189 as the difference between DO concentrations at t = 0 and following the 5-day incubations. We 190 assumed 100% of DO consumed was converted to CO_2 via aerobic respiration and that the

191 carbon source respired was DOM, where resulting BOD measurements were used an analog for bioavailable DOC. The Winkler method we used here has been used extensively and is attractive 192 for a variety of reasons, including: (i) enabling DO to be measured with precision of 0.01 mg/L, 193 thus low respiration rates can be accurately measured; (ii) allowing for convenient replication of 194 assays within habitats; (iii) permitting experimental manipulation of standard bioassays (e.g., N 195 196 and P amendments, photolysis experiments, alteration of initial microbial consortia, and temperature manipulation; (iv) helping to segregate the relative roles of water column and 197 198 sediment processes (through comparisons with sediment analyses); and (v) helping to inform 199 more realistic ecosystem-level experiments that are much more laborious and time intensive. In order to investigate the optical characteristics of the DOM in these samples, we 200 201 additionally measured the ultraviolet-visible absorption spectra of CDOM from this broad collection of waters. CDOM absorbance was measured on filtered (precombusted Whatman 0.7 202 203 µm GF/F), unacidified waters stored in acid-washed HDPE bottles immediately after collection (within ~1 day) at the Northeast Science Station in Cherskiy using a Thermo Scientific 204 GENESYS 10 UV/Vis Spectrophotometer across wavelengths 800-200 nm (1 nm interval) with 205 206 a 1 cm quartz cuvette. All sample spectra were blank corrected using Milli-Q water (18 Ω). Measurements were made after samples had equilibrated to the laboratory temperature in order 207 to minimize temperature effects. Null-point adjustments were performed on all spectra, such that 208 209 CDOM absorbance was assumed to be zero across wavelengths greater than 750 nm and the 210 average absorbance between 750 nm and 800 nm was subtracted from each spectrum to correct for offsets owing to instrument baseline drift, temperature, scattering, and refractive effects 211 (Green and Blough, 1994; Helms et al, 2008). CDOM absorption coefficients were calculated 212 213 as:

$$a(\lambda) = 2.303A(\lambda)/l \tag{1}$$

where a is the Napierian absorption coefficient (m⁻¹) at a specified wavelength (λ , in nm), $A(\lambda)$ is 215 the absorbance at the wavelength, and *l* is the cell path length in meters (Green and Blough, 216 217 1994). To avoid inner-filtering effects, several highly absorbing samples (primarily the soil pore 218 waters) were diluted with Milli-Q water before analysis (to the point where A_{350} was ≤ 0.02 at for a 1 cm path length was ≤ 0.02) to avoid saturation of the spectra at short wavelengths, where the 219 220 final CDOM absorbance and therefore absorption coefficients were corrected for these 221 procedures. CDOM spectral slopes (S, nm^{-1}) between 290–350 nm $(S_{290-350})$, 275–295 nm $(S_{275-295})$, 222 and 350–400 nm ($S_{350-400}$), calculated within log-transformed absorption spectra, were also 223 224 utilized to investigate DOM characteristics of contrasting water types, and were calculated as: $a(\lambda) = a(_{\lambda ref}) e^{-S(\lambda - \lambda ref)}$ (2)225 where $a(\lambda)$ is the absorption coefficient at a specified wavelength, λ_{ref} is a reference wavelength, 226 and S is the slope fitting parameter (Hernes et al., 2008; Helms et al., 2008; Spencer et al., 227 228 2009a). All slopes are reported here as positive values, such that higher (i.e., steeper) slopes indicate a greater decrease in absorption with increasing wavelength. Additional CDOM 229 parameters investigated here include the spectral slope ratio (S_R) , calculated as the ratio between 230 $S_{275-295}$ and $S_{350-400}$; the ratio between CDOM absorption coefficients (a) at 250 nm and 365 nm 231 $(a_{250}:a_{365})$; and specific UV absorbance (SUVA₂₅₄), determined by dividing UV absorbance (A) 232

at 254 nm by the sample DOC concentration and reported in units of L mg C^{-1} m⁻¹ (Weishhar et

al., 2003). These six CDOM parameters ($S_{290-350}$, $S_{275-295}$, $S_{350-400}$, a_{250} : a_{365} , SUVA₂₅₄, and S_R)

have been shown to provide insights for various DOM characteristics such as molecular weight,

source waters, composition, age, and aromatic content for a variety of geographic regions (e.g.,

Weishaar 2003; Neff et al., 2006; Helms et al., 2008; Spencer et al., 2008; Spencer et al., 2009a;
Spencer et al., 2009b; Mann et al., 2012). As such, we chose our method for spectral slope
calculations to be consistent with previous studies to foster intercomparisons between datasets,
however future studies may derive further insight utilizing methods that calculate a continuous
spectral slope curve over the full 200–800 nm span (e.g., Loiselle et al., 2009) rather than only
specific wavelength intervals as presented here.

243

244 **3. Results**

245 Total DOC concentrations (and the variance among values within each water type) decreased markedly downstream along the flow-path continuum from soil pore waters to the 246 Kolyma River mainstem (Figure 2a). Mean (± 1 standard deviation) DOC values were 43.3 \pm 247 22.8 mg L⁻¹ (soil pore waters), 11.6 ± 3.0 mg L⁻¹ (streams), 4.9 ± 1.6 mg L⁻¹ (rivers), and $3.6 \pm$ 248 0.4 mg L^{-1} (mainstem waters). Soil pore waters, in particular, showed highly variable DOC 249 concentrations (ranging from 13.2 to 64.7 mg L^{-1}) demonstrating the heterogeneous supply of 250 DOM from terrestrial systems to streams. By contrast, DOC concentrations in the Kolyma 251 mainstem along the ~250 km stretch sampled were remarkably similar (ranging from 3.0 to 4.4 252 $mg L^{-1}$) during this mid-summer July period (Figure 2a). Furthermore, DOC concentrations of 253 the four water types sampled were found to be significantly different from one another (one-way 254 ANOVA, *p*<0.05). 255

256 Concentrations of bioavailable DOC showed similar patterns to DOC, declining 257 downstream along the flow-path continuum with increasing water residence time in the system 258 (Figure 2b). Bioavailable DOC concentrations averaged 0.9 ± 0.2 mg L⁻¹ (soil pore waters), 0.3 259 ± 0.1 mg L⁻¹ (streams), 0.3 ± 0.2 mg L⁻¹ (rivers), and 0.2 ± 0.2 mg L⁻¹ (mainstem waters), and

260	showed relative greater variability than DOC within the stream, river and mainstem water types.			
261	Concentrations of bioavailable DOC in soil pore waters were statistically different from the othe			
262	three water types (one-way ANOVA, $p < 0.05$), although by contrast, streams, rivers, and			
263	mainstem waters were not statistically different from one another (p >0.05). Importantly, the			
264	percentage of bioavailable DOC (i.e., calculated as the amount of bioavailable DOC divided by			
265	total DOC) did not significantly decrease downstream (one-way ANOVA, $p>0.05$) and showed			
266	relatively similar values among the four water sample types along the flow-path continuum			
267	(Figure 2c), where percentages averaged $3.9 \pm 3.8\%$ (soil pore waters), $3.2 \pm 1.9\%$ (streams), 6.2			
268	\pm 4.3% (rivers), and 4.5 \pm 4.5% (mainstem waters).			
269	CDOM absorption spectra (200-800 nm) showed clear separation between soil pore			
270	waters, streams, rivers, and the Kolyma mainstem, where soil pore waters exhibited values			
271	markedly higher than the other three water sample types (Figure 3a). CDOM absorption also			
272	clearly declined downstream from streams, rivers, to mainstem waters when assessing those			
273	waters only (Figure 3b). Furthermore, we investigated the potential for utilizing CDOM			
274	absorption as a proxy for DOC concentrations in these waters. Our data revealed that			
275	independent of water type along the stream-river-mainstem flow-path, CDOM absorption was			
276	strongly linearly correlated to DOC concentrations at 254, 350, and 440 nm (Figure 4). In			
277	particular, CDOM absorption at 254 nm had the highest predictive capability of DOC ($r^2 =$			
278	0.958, <i>p</i> <0.01), with CDOM absorption at 350 nm ($r^2 = 0.855$, <i>p</i> <0.01) and 440 nm ($r^2=0.667$,			
279	p < 0.01) less strongly predictive (Figure 4).			
280	We additionally investigated the quantitative distribution of the six derived CDOM			

281 parameters ($S_{290-350}$, $S_{275-295}$, $S_{350-400}$, a_{250} : a_{365} , SUVA₂₅₄, and S_R) across the four water types

(Figure 5; Table 1). In general, four parameters ($S_{290-350}$, $S_{275-295}$, a_{250} : a_{365} , and S_R) showed an 282

283	increasing pattern along the flow-path continuum, whereas two parameters ($S_{350-400}$ and
284	SUVA ₂₅₄) showed a decreasing pattern. In terms of whether the values of the six parameters
285	were statistically significantly different among water sample types, one-way ANOVA tests (at
286	the 0.05 level) revealed inconsistent results. Most commonly, soil pore waters were statistically
287	different from all other water types for four of the parameters ($S_{290-350}$, $S_{275-295}$, a_{250} : a_{365} , and S_R),
288	but no consistent pattern was observed in significant differences across other water types.
289	However, the spectral slope ratio (S_R) was the only CDOM parameter of the six investigated that
290	showed statistically significant differences between all four water types ($p < 0.05$).
291	Lastly, we examined the relationships between CDOM optical properties and DOM
292	bioavailability. To this end, we performed linear regressions between all six of our derived
293	CDOM parameters and bioavailable DOC concentrations to determine the strength of their
294	ability to predict bioavailable DOC. Our results indicated that five of the CDOM parameters
295	$(S_{290-350}, S_{275-295}, a_{250}:a_{365}, SUVA_{254}, and S_R)$ were statistically significant predictors at the 0.05
296	level (Table 2). In particular, S_R showed the strongest relationship with bioavailable DOC
297	concentrations (r^2 value = 0.45, $p < 0.01$). The relationship between bioavailable DOC
298	concentrations and S_R (Figure 6) showed a distinct negative trend (bioavailable DOC mg L ⁻¹ = -
299	2.204(S_R) + 2.518), with the highest bioavailable DOC concentrations and lowest S_R values for
300	soil pore waters, and lowest bioavailable DOC concentrations and highest S_R values for Kolyma
301	River mainstem waters. We found a clear gradation in the relationship between S_R and
302	bioavailable DOC down the flow-path continuum, as one would also expect by examining these
303	parameters individually (e.g., Figures 2b, 5f). In summary, not only was S_R the only CDOM
304	parameter that showed statistically significant separation between all four water types examined,

but it also had the strongest relationship when compared with concentrations of bioavailableDOC.

307

308 4. Discussion and Conclusions

In this study, we present a full suite of DOC, bioavailable DOC, and CDOM parameters 309 310 throughout the permafrost-dominated Kolyma River basin in Northeast Siberia with the purpose of helping to elucidate the processing of DOM along a full flow-path continuum from soil pore 311 312 waters to the mainstem. Our findings show that average concentrations of DOC and bioavailable 313 DOC generally decrease as waters travel downstream from soil pore waters, streams, rivers, and ultimately to the Kolyma River mainstem. This pattern suggests the occurrence of rapid in-314 315 stream processing of DOM and potential remineralization of DOC to atmospheric CO_2 during this July baseflow period well before these waters reach the Arctic Ocean (e.g., Denfeld et al., 316 2013; Mann et al., 2015; Spencer et al., 2015). The amount of total DOC putatively lost to 317 318 remineralization is a relatively small fraction ($\sim 3-6\%$ depending upon water type), but on par with similar studies across the Arctic for this time of year (e.g., Holmes et al., 2008). Although 319 this may be a relatively small proportion, it is likely the permafrost-derived, ancient DOC found 320 321 in headwaters that is contributing to permafrost carbon feedbacks to climate warming (Mann et 322 al., 2015). Moving downstream, the river continuum concept predicts that relative diversity of organic molecules decreases from the headwaters to the river mouth (Vannote et al., 1980). As 323 324 energetically favorable compounds are converted to living tissue or respired as CO₂, bulk DOM in the Kolyma basin has indeed been shown in previous studies to become less diverse moving 325 326 from headwaters to mainstem waters before exported to the Arctic Ocean (Spencer et al., 2015).

327 CDOM parameters presented in this study give further insight into characteristics of DOM along the full flow-path continuum throughout the Kolyma River basin. For instance, the 328 specific ultraviolet absorbance (SUVA₂₅₄) has been shown to be correlated with DOM 329 composition, where SUVA₂₅₄ values are positively correlated with percent aromaticity and 330 molecular size of DOM (and for a given river have been shown to be greatest during spring 331 332 flood) (e.g., Weishaar et al., 2003; Spencer et al., 2009a; Mann et al., 2012). In this study, we generally found progressively decreasing $SUVA_{254}$ values along the flow-path from soil pore 333 334 waters towards mainstem waters, suggesting that soil pore waters contain higher molecular 335 weight, aromatic terrestrial DOM that generally becomes lower in molecular weight and aromaticity along the flow-path continuum towards the Kolyma River mainstem. In addition, the 336 a_{250} : a_{365} ratio has been shown to be negatively correlated to aromaticity and molecular size of 337 DOM (Peuravuori and Pihlaja, 1997). In fact (similar to samples from the Yukon River, Alaska 338 (Spencer et al., 2009a)), our data showed that the a_{250} : a_{365} ratio is significantly negatively 339 correlated with SUVA₂₅₄ (a_{250} : a_{365} = -0.947 (SUVA₂₅₄) – 0.947; r²=0.49, p<0.01). As such, the 340 a_{250} : a_{365} ratio may potentially be utilized as a first-order proxy for SUVA₂₅₄ when DOC 341 concentrations cannot be easily determined. 342

However, despite our observations of downstream shifts in DOM composition, we find a relatively constant proportion of DOC that was bioavailable (~3–6% of total DOC) regardless of relative water residence time along the flow-path. This suggests that continual microbial processing of organic matter is able to occur with similar rates during transit from headwaters throughout the Kolyma River drainage network to the Arctic Ocean concurrent with ongoing downstream CDOM compositional changes. Microbial demand in headwater streams of the Kolyma River basin is subsidized by significant quantities of DOC specifically derived from

permafrost and aged soils, yet the proportion of permafrost supporting DOC mineralization
declines as waters move downstream through the fluvial network (Mann et al., 2015). Thus, our
results importantly show that microbial metabolism continues at similar rates independent of
dominant DOM source and radiocarbon age.

There may be several reasons for why microbial metabolism maintains this consistent 354 355 rate along the flow-path, including the possibility that aquatic microorganisms are acclimating to 356 a downstream shift in DOM composition. The higher overall amounts of bioavailable DOC we 357 measured in soil pore waters may reflect a highly bioreactive permafrost or aged surface soil 358 derived fraction of the bulk DOC pool (e.g., Vonk et al., 2013; Mann et al., 2014). Further downstream in larger tributary and Kolyma mainstem waters, it has been shown that lower total 359 amounts of bioavailable DOC is supported almost entirely from predominantly modern 360 radiocarbon aged surface soils and vegetation sources (Mann et al., 2015). Aquatic 361 microorganisms may therefore be readily acclimating to significant shifts in DOM composition 362 caused by selective losses of unique DOM fractions (e.g., Kaplan and Bott, 1983; Spencer et al., 363 2015) alongside high-internal demand for labile DOM by stream communities in lower order 364 streams, which would otherwise generally be expected to result in decreased DOM lability with 365 366 increasing water residence time (Stepanauskas et al., 1999a,b; Wikner et al., 1999; Langenheder et al., 2003; Sondergaard et al., 2003; Fellman, 2010; Fellman et al., 2014). 367

Additional mechanisms such as increasing photodegradation downstream may also account for our observed patterns in downstream DOM. Previous studies have indicated that CDOM spectral slopes (particularly $S_{290-350}$ and $S_{275-295}$) can serve as indicators of DOM source and composition, where a steeper spectral slope typically suggests lower molecular weight material with decreasing aromatic content and a shallower (i.e., lower) slope typically suggests

373 higher molecular weight material with increasing aromatic content (Green and Blough, 1994; 374 Blough and Del Vecchio, 2002; Helms et al., 2008; Spencer et al., 2008; Spencer et al., 2009a). Furthermore, $S_{275-295}$ has been identified as a reliable proxy for dissolved lignin and therefore 375 terrigenous DOM supply across Arctic Ocean coastal waters, as well as photobleaching history 376 (Helms et al., 2008; Fichot et al., 2013). We found a general increase in $S_{290-350}$ and $S_{275-295}$ 377 378 moving downstream through the network, indicative of progressive photodegradation of DOM alongside likely reductions in average DOM molecular weight and aromaticity. We found 379 380 spectral slopes over longer wavelength regions $(S_{350-400})$ decreased through the network, also 381 suggesting constant photochemical degradation of DOM as waters flowed downstream (e.g., 382 Helms et al., 2008). The slope ratio (S_R) has also been shown to be a proxy for DOM molecular weight and source, where low ratios typically correspond to more allochthonous, higher 383 molecular weight DOM (Helms et al., 2008; Spencer et al., 2009b; Mann et al., 2012). The 384 advantage of S_R ratios over individual S values is apparent when each spectral slope responds to 385 a process in an opposing manner, emphasizing the response in calculated $S_{\rm R}$ values. The clear 386 increases in S_R we observed moving downstream in the fluvial network (from a minimum of 0.74 387 in soil pore waters to a maximum of 1.24 in the mainstem) indicate that during July summer 388 389 conditions, soil pore waters contain higher molecular weight, aromatic terrestrial DOM that 390 generally becomes lower in average molecular weight and aromaticity along the flow-path continuum towards the Kolyma River mainstem. The maximum S_R value of 1.24 we report in 391 392 the Kolyma River mainstem is markedly higher than the range of $S_{\rm R}$ (0.82–0.92) reported in Stedmon et al. (2011) for the Kolyma from 2004 and 2005, demonstrating the heterogeneity of 393 394 DOM properties even in mainstem waters and the necessity for greater temporal resolution in 395 monitoring. Similar to spectral slopes, S_R values may also be indicative of photobleaching

history (e.g., Helms et al., 2008) and our we observed increase in S_R downstream through the network suggests evidence of on-going photochemical degradation of surface water DOM during transit.

Photodegradation may indeed play an important and direct role in our observed consistent 399 fraction of bioavailable DOC along the flow-path. Previous studies in the Arctic underscore the 400 401 importance of residence times as well as a significant combined role for photo- and biological degradation along the flow-path in Arctic watersheds (Cory et al., 2007; Merck et al., 2012; Cory 402 et al., 2013; Laurion and Mladenov, 2013). These previous results show that the photochemical 403 404 "pretreatment" of stream DOM that occurs during export into lakes and coastal zones may impact the ability of microorganisms to mineralize DOM. Therefore, the residence times and 405 flow-paths of waters should greatly influence the ultimate fate of DOM (e.g., DOM vs. CO₂) 406 exported to the adjacent ocean. In our case, we find that our increasing S_R values downstream 407 suggest important photodegradation processes are occurring along the flow-path continuum, 408 409 where this photodegradation may potentially release significant quantities of labile DOM for continued microbial processing of DOM further downstream in these stream networks. In other 410 words, our results suggest that the more abundant "virgin" newly exposed bioavailable molecules 411 412 upstream are replaced downstream by photobleached smaller molecules (originating from aromatic compounds), resulting in the fraction of DOC used relatively constant without any clear 413 pattern overall. If this (or something similar) were not the case, we would expect to see a 414 415 declining fraction of bioavailable DOC along the flow-path continuum. In this study, we have provided new and important findings with regards to the spatial 416 417 distribution of DOM concentration, bioavailability, and optical properties during mid-summer 418 hydrologic conditions throughout the Kolyma River basin in Northeast Siberia. Freshwater DOC

419	measurements across the network were strongly positively correlated to CDOM absorption at
420	254 nm ($r^2 = 0.958$, $p < 0.01$), confirming the utility of simple CDOM optical measurements for
421	estimating carbon concentrations in arctic freshwaters (Spencer et al., 2008, 2009a; Stedmon et
422	al., 2011) and across water types within the Kolyma River basin in particular. Furthermore, the
423	optical parameter S_R proved to be the only CDOM compositional measure that showed
424	statistically significant separation between all four water types examined during the study period,
425	suggesting that this parameter may be useful for easily distinguishing characteristics and
426	processes occurring in organic matter among water types along the full flow-path continuum.
427	The significant increase in S_R values we observed downstream through the network suggests
428	evidence of on-going photochemical degradation of surface water DOM during transit.
429	Additionally, of all the CDOM parameters, S_R values were most closely related to concentrations
430	of bioavailable DOC ($r^2 = 0.454$, $p < 0.01$), suggesting that this value may be correlated with a
431	decline in bioavailable DOC through the network. However, biological degradation has
432	previously been shown to typically slightly decrease S_R values (Helms et al., 2008), which
433	indicates that the opposite relationship observed here may instead be a consequence of co-
434	variance with photodegradation of DOM, or demonstrate that S_R values may reflect a broader,
435	more complex range of physical and biological processes than previously recognized. Garnering
436	further insight from our measurements, the relatively constant proportion of DOC that was
437	bioavailable regardless of relative water residence time along the flow-path may be a
438	consequence of two potential scenarios allowing for continual processing of organic material
439	within the system, namely: (a) aquatic microorganisms are acclimating to a downstream shift in
440	DOM composition; and/or (b) photodegradation is continually generating labile DOM for
441	continued microbial processing of DOM along the flow-path continuum. Without such

442 processes, we would otherwise expect to see a declining fraction of bioavailable DOC443 downstream with increasing residence time of water in the system.

Unlike many previous studies that focus on only mainstem rivers in the Arctic, we focus 444 here on a variety of waters along a full flow-path continuum, showing that CDOM metrics (in 445 particular, $S_{\rm R}$) reflect important compositional differences in DOM of waters along the transit 446 447 from headwaters to the Arctic Ocean. The range in DOM properties of waters travelling downstream through the Kolyma Basin often spanned wider ranges than DOM compositional 448 449 differences reported annually among the six major arctic rivers. For example, $S_{\rm R}$ values across 450 the major arctic rivers over the years 2004 and 2005 spanned a minimum of 0.79 in the Yenisey River, to a maximum value of 1.11 in the Mackenzie River (Stedmon et al., 2011), compared to 451 452 the range of 0.74–1.24 for waters in our study within a single basin. It is therefore essential that changes taking place in the quality of CDOM exported by these rivers be examined throughout 453 entire river basins in order to adequately assess climate driven shifts in terrigenous carbon supply 454 455 and reactivity. Future work that includes both photo- and microbial degradation experiments may further elucidate the ability for $S_{\rm R}$ to serve as a direct proxy for these processes along a 456 flow-path gradient. Our overall results thus far demonstrate promise for utilizing ultraviolet-457 458 visible absorption characteristics to easily, inexpensively, and comprehensively monitor the 459 quantity and quality of DOM (over broad ranges) across permafrost landscapes in the Arctic. 460 This is particularly critical for remote arctic landscapes such as those in Northeast Siberia, where 461 the future fate of organic carbon currently frozen in permafrost soils (and whether it ultimately is released as CO₂ and CH₄) is tightly linked to the lability of this material. 462

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- **Table 1.** Mean spectral slope and other CDOM parameters for soil pore waters, streams, rivers,
- 722 and the Kolyma River mainstem.

	$S_{290-350}$ (× 10 ⁻³ nm ⁻¹)	$S_{275-295}$ (× 10 ⁻³ nm ⁻¹)	$S_{350-400}$ (× 10 ⁻³ nm ⁻¹)	a_{250} : a_{365}	$\begin{array}{c} SUVA_{254} \\ (L\ mg\ C^{-1}\ m^{-1}) \end{array}$	S _R
Soil pore waters	15.35	15.27	18.65	5.47	3.52	0.82
Streams	17.08	17.39	18.89	6.44	2.94	0.92
Rivers	17.17	17.79	18.19	6.27	2.77	0.98
Kolyma Mainstem	18.10	18.57	17.50	6.53	2.56	1.06

Table 2. Relationships between bioavailable DOC and each of the six CDOM metrics

investigated. $S_{\rm R}$ shows the highest r-squared value, with a *p*-value of 0.00002.

	r^2	<i>p</i> -value
S ₂₉₀₋₃₅₀	0.3560	0.00025
S ₂₇₅₋₂₉₅	0.4497	0.00002
S ₃₅₀₋₄₀₀	0.0443	0.23987
<i>a</i> 250: <i>a</i> 365	0.2645	0.00220
SUVA ₂₅₄	0.1980	0.01376
$S_{\mathbf{R}}$	0.4540	0.00002

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Figure 1. The northern reaches of the Kolyma River in East Siberia and the locations of the 47 734 water samples collected throughout the region in this study (including soil pore waters, streams, 735 736 rivers, and the Kolyma River mainstem). 737 Figure 2. Concentrations of (a) dissolved organic carbon (DOC), (b) bioavailable DOC, and (c) 738 739 percentage of total DOC that is bioavailable for the four water sample types. The mean (hollow squares), median (horizontal lines), ± 1 standard deviation (gray boxes), and total range 740 741 (whiskers) for each sample population are shown. 742 Figure 3. Chromophoric dissolved organic carbon (CDOM) absorption spectra from 200–800 743 744 nm for (a) all samples; and (b) streams, rivers, and the Kolyma River mainstem only. 745 Figure 4. Relationships between DOC and CDOM absorption at 254, 350, and 440 nm for 746 747 streams, rivers, and the Kolyma River mainstem. 748 749 **Figure 5.** The six presented CDOM metrics, (a) $S_{290-350}$, (b) $S_{275-295}$, (c) $S_{350-400}$, (d) a_{250} : a_{365} , (e) SUVA₂₅₄, and (f) S_R, show the separation between soil pore, stream, river, and Kolyma mainstem 750 751 waters. The mean (hollow squares), median (horizontal lines), ± 1 standard deviation (gray 752 boxes), and total range (whiskers) for each sample population are shown.

- **Figure 6.** The CDOM metric S_R shows a relatively strong relationship with concentrations of
- bioavailable DOC present in the sampled waters, with an r-squared value of 0.4540 and *p*-value

756 <0.01.



Figure 1.



Figure 2.



Figure 3.



Figure 4.



Figure 5.



Figure 6.