



# Consequences of Increased Variation in Peatland Hydrology for Carbon Storage: Legacy Effects of Drought and Flood in a Boreal Fen Ecosystem

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Kane ES, Dieleman CM, Rupp D, Wyatt KH, Rober AR and Turetsky MR (2021) Consequences of Increased Variation in Peatland Hydrology for Carbon Storage: Legacy Effects of Drought and Flood in a Boreal Fen Ecosystem. Front. Earth Sci. 8:577746. doi: 10.3389/feart.2020.577746 Globally important carbon (C) stores in boreal peatlands are vulnerable to altered hydrology through changes in precipitation and runoff patterns, groundwater inputs, and a changing cryosphere. These changes can affect the extent of boreal wetlands and their ability to sequester and transform C and other nutrients. Variation in precipitation patterns has also been increasing, with greater occurrences of both flooding and drought periods. Recent work has pointed to the increasing role of algal production in regulating C cycling during flooded periods in fen peatlands, but exactly how this affects the C sink-strength of these ecosystems is poorly understood. We evaluated temporal trends in algal biomass, ecosystem C uptake and respiration (using static and floating chamber techniques), and spectroscopic indicators of DOM quality (absorbance and fluorescence) in a boreal rich-fen peatland in which water table position had been experimentally manipulated for 13 years. Superimposed on the water table treatments were natural variations in hydrology, including periods of flooding, which allowed us to examine the legacy effects of flooding on C cycling dynamics. We had a particular focus on understanding the role of algae in regulating C cycling, as the relative contribution of algal production was observed to significantly increase with flooding. Ecosystem measures of gross primary production (GPP) increased with algal biomass, with higher algal biomass and GPP measured in the lowered water table treatment two years after persistent flooding. Prior to flooding the lowered treatment was the weakest C sink (as CO<sub>2</sub>), but this treatment became the strongest sink after flooding. The lower degree of humification (lower humification index, HIX) and yet lower bioavailability (higher spectral slope ratio, Sr) of DOM observed in the raised treatment prior to flooding persisted after two years of flooding. An index of free or bound proteins (tyrosine index, TI) increased with algal biomass across all plots during flooding, and was lowest in the raised treatment. As such, antecedent drainage conditions determined the sink-strength of this rich fen - which was also reflected in DOM characteristics. These findings indicate that monitoring flooding history and its effects on algal production could become important to estimates of C balance in northern wetlands.

Keywords: boreal, peat, climate change, carbon dioxide, dissolved organic matter, DOM, spectroscopy, algae

1

# INTRODUCTION

Northern peatlands are a globally important reservoir of carbon (C), representing a large magnitude of C vulnerable to loss in a changing climate (Goldstein et al., 2020). Rich fens are one of the most common boreal peatland types in western North America (Vitt et al., 2000), and open peatlands and mineral wetlands comprise about 85% of wetland area in Alaska (Kolka et al., 2018) - an area of land nearly twice that of all wetland area in the contiguous United States (Ford and Bedford, 1987). Moreover, the extent of fens in interior Alaska has been increasing owing to permafrost degradation and changing ground water and runoff patterns in recent decades (Jorgenson et al., 2001; Lara et al., 2016). Observed regional climate trends and modelling efforts also suggest increased precipitation with overall higher variation (anomalies) in total amounts in Alaska's boreal region (Stewart et al., 2013; Euskirchen et al., 2016). These patterns suggest prolonged or more variable periods of inundation in the future, with yet uncertain consequences for the C storage capacity of Alaska's vast fen ecosystem complexes.

The carbon balance of peatlands has shown varied responses to changes in the water table position occurring with periods of flooding or drought. While high water tables are generally associated with higher fluxes of methane (e.g., Turetsky et al., 2014), the balance between ecosystem respiration (ER) and gross primary production (GPP) can vary with changes in water table position (Belyea, 2009; Strack et al., 2009). For example, a decade of research in an interior Alaskan fen has shown that drought conditions generally reduced GPP, but had muted effects on ER, which switched from increased autotrophic respiration in wet years to increased heterotrophic respiration in dry years (Olefeldt et al., 2017). However with flooding, variation in respiration in this fen complex changed with the timing and sources of inundation, with persistent flooding late into the growing season resulting in lower GPP relative to ER (Euskirchen et al., 2020). Exactly how constituent autotrophic and heterotrophic respiration components are likely to change with different regimes of flooding are not yet well understood.

A key difference in the active mode of primary production during flooded conditions, which affords a photic zone within the water column, is the relative activity of algae (Goldsborough and Robinson, 1996). While little is still known about algal production in boreal fens, recent work highlights the importance of algae as a component of the photosynthetic community during flooded conditions in interior Alaskan peatlands (Wyatt and Turetsky, 2015; DeColibus et al., 2017). Moreover, antecedent drainage conditions prior to flooding have been shown to exert strong controls over algal production. Experimental drainage in an Alaskan fen peatland resulted in increased pore water nutrient concentrations which fostered algal production upon flooding due to high rainfall (Wyatt et al., 2012). However, exactly how long this "ecosystem memory" of antecedent dry conditions can affect algal production, and how this in turn affects net C balance with persistent flooding is not clear (DeColibus et al., 2017). For example, observed increases in the exudates (dissolved organic matter, DOM) concomitant with increased algal production are very bioavailable (Bertilsson and Jones, 2003; Wyatt et al., 2012;

Wyatt and Rober 2020) and could increase ER with persistent flooding. As such, component fluxes of ecosystem respiration along with algal production need to be measured together during periods of persistent flooding to ascertain the net consequences of flooding history on C balance in fens.

We investigated trace gas production, algal biomass, and DOM quality in an experimental manipulation of water table positions that simulate both drought (lowered water table treatment) and flooding (raised water table treatment) conditions relative to a control treatment without manipulation in an Alaskan boreal rich fen. We conducted our measurements prior to flooding across all experimental treatments and after two years of consecutive flooding. Our over-arching hypothesis was that hydrologic variation affects the balance between GPP and ER (Figure 1). With flooding, we expected algal production to offset carbon losses through ER, increasing the C sink-strength of the ecosystem. As such, we hypothesized the lowered water table treatment would be a stronger sink of C when reflooded owing to greater algal biomass associated with elevated nutrient availability, whereas the previously raised water table treatment would be a weaker sink of C when flooded, owing to reduced GPP from vascular plants. On the other hand, increased water-soluble organic carbon, algae-derived DOM and nutrients from the lowered water table treatment could increase heterotrophic respiration, and hence drive higher ER relative to GPP in the flooded environment. However, exactly how antecedent changes in DOM quality persist with prolonged flooding periods are not well understood. Therefore, we specifically ask: 1) How does the legacy of flooding history influence peatland C balance? 2) What is the role of DOM quality in driving ecosystem C fluxes and how does algal biomass relate to DOM quality?

# **MATERIALS AND METHODS**

# Study Site and Experimental Design

The rich fen (surface pH ~6) is in the boreal peatland lowlands of the Tanana Flats of interior Alaska (cf. Hopkins et al., 1955) approximately 30 km southeast of Fairbanks, Alaska United States. The vegetation and climate characteristics of this site have been previously described in detail (Churchill et al., 2015; Olefeldt et al., 2017; Euskirchen et al., 2020), but briefly, emergent vascular species (Equisetum, Carex, and Comarum (Potentilla) genera) and some brown mosses and Sphagnum dominate the site. The fen complex lacks permafrost but is within the region of discontinuous permafrost in Alaska's interior. There is little microtopography at the rich fen, which has accumulated peat to a depth of approximately 1–1.5 m. The site is associated with the Bonanza Creek Long Term Ecological Research Program (Iter. uaf.edu).

The Alaska Peatland Experiment (APEX) was established in 2005 to investigate the effects of altered hydrologic regimes on C cycling processes in a controlled experiment. Three plots (approximately  $20 \times 20$  m) were assigned each to one of three water table (WT) treatments (raised, lowered, and a control

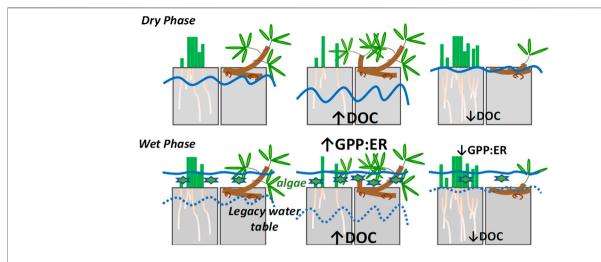


FIGURE 1 | Conceptual model of how our experimental water table treatments (control, lowered and raised) have affected the pool size of nutrients, dissolved organic carbon [DOC] and its quality, and how these factors have affected algal production during flooding (wet phase). We hypothesize that these factors affect the balance between gross primary production and ecosystem respiration (GPP:ER). The solid blue lines indicate water table position by treatment, and the dashed lines indicate legacy water table position during flooding. The green stars indicate the relative abundance of algae. The plants indicate relative abundances of sedges and shrubs (see Rupp et al., 2019).

reference plot, each approximately 25 m apart), as previously described in detail (see Turetsky et al., 2008 and Chivers et al., 2009 for experiment details). Briefly, in the winter of 2005 channels were dug to facilitate water drainage from the lowered water table plot. This water in turn was pumped into the raised water table treatment with solar-powered bilge pumps, throughout the growing season (approximately June-September). There were no significant differences in water table position or vegetation composition among plots prior to treatment initiation. The goal of the experiment was to maintain lowered (drained) and raised water table positions relative to the control while allowing for the natural seasonal variability in water table position at this site. However, it was impossible to experimentally manipulate the water table position during periods of natural flooding, including years when flooding (water table > 40 cm above peat surface) persisted for the entire growing season (2014, 2017 and 2018; see Euskirchen et al., 2020 for hydrograph data). In this current study, we measured trace gas flux, algal biomass, and DOM quality within the APEX plots prior to a natural flooding event in 2016, which persisted through 2017 and 2018. We remeasured the experimental plots after continuous flooding during the growing season of 2018 to test our hypotheses regarding persistent flooding effects on C cycle dynamics.

#### Gas Flux Measurements

We used the static chamber technique for gas flux measurements during non-flooded years as previously described in detail (Chivers et al., 2009). Briefly, six collars ( $60 \times 60 \,\mathrm{cm}$ ) were inserted to a depth of 10 cm in 2005 within each WT treatment. A clear lexan chamber (0.227 m³) with closed-cell foam sealed the chamber with the collar. Two computer CPU fans mixed the air within the chamber during measurements. The change in chamber  $CO_2$  concentrations was measured for approximately 4 min using a portable infrared gas analyzer

(IRGA; PP Systems EGM 4, Amesbury, Massachusetts, United States). ER was measured by shading chambers with an opaque shroud and net ecosystem exchange (NEE) was measured without a shroud. Temperature, relative humidity, and photosynthetically active radiation (PAR) were logged continuously within the chamber during each flux measurement with a PP Systems TRP-1 sensor. Measurements occurred approximately bi-weekly.

A floating chamber was employed for gas flux measurements in flooded conditions during the growing season of 2018. This static chamber was constructed from a clear polycarbonate plastic bucket (18.9 L) with Styrofoam floatation around the base and one CPU fan inside for air circulation. This design allows for the measure of CO<sub>2</sub> flux in the airspace above the water. The floating chamber was equipped with an airlock to equilibrate pressure. The clear chamber was covered with an opaque shroud during ER measurements and NEE was measured without a shroud. Temperature, relative humidity, and PAR were logged continuously with a PP Systems TRP-1 sensor mounted within the clear floating chamber. The change in chamber CO<sub>2</sub> concentrations was measured with an IRGA as previously described. Floating chamber measurements were conducted within water table manipulation plots (within plot n = 3) from a small inflatable boat and by wading along the established boardwalk network.

# Algal Biomass

Algal material was collected biweekly during the 2018 growing season for estimates of algal biomass measured as chlorophyll a (mg cm<sup>-2</sup>). Owing to shallow water with a well-lit photic zone, algae within this fen complex grow primarily affixed to submersed substrata (DeColibus et al., 2017). Attached algal material was collected from four senesced *Carex utriculata* stems (10 cm length) cut below the water surface within each

of three randomly selected 1 m<sup>2</sup> areas per water table treatment using methods modified from DeColibus et al. (2017). There was no floating or loosely attached algae (i.e. metaphyton) observed during the study. Each composite sample was placed into a clean 15 ml polyethylene tube and kept on ice until returning to the lab where samples were frozen at -20°C until subsequent processing. Algal chlorophyll a was extracted with 90% ethanol in the dark for 24 h and the resulting pigment extracts were analyzed with an Agilent Cary 60 UV-Vis spectrophotometer (Agilent Technologies, Santa Clara, CA, United States) at 665 and 750 nm after acidification to correct for phaeophytin (APHA, 1998).

# **Dissolved Organic Matter Quality**

When the water table was below the peat surface in 2016, a stainless steel "sipper" (0.52 cm diameter tubing with 2 cm slotted region at the end) was carefully inserted into the peat to a depth of 20 cm. Pore water was drawn up through the sipper into a syringe that was rinsed first with deionized (DI) water, then with sample prior to collection. When conditions were flooded in 2018, surface waters from the photic zone were collected in addition to pore water. Specifically, a clean, sample rinsed 60 ml syringe was used to sample the water column 10 cm below the surface of the water. After collection, the sample was injected into a 125 ml dinitrogen gas flushed and evacuated Wheaton glass vial capped with a butyl rubber septa via a 0.45 µm Whatman syringe filter and needle and then refrigerated prior to analysis, following Rupp et al. (2019). Samples were collected at three locations, spaced approximately 3 m apart, in each WT treatment throughout the growing seasons of 2016 (prior to flooding) and 2018 (during sustained flooding). Dissolved organic carbon (DOC) and total dissolved nitrogen (TDN) were determined on samples within 10 days on an TOCV Analyzer with TDN module (Shimadzu Scientific Instruments, Columbia, MD, United States).

Pore water was extracted from each of the Wheaton vials and run on a fluorometer (Horiba–Jobin–Yvon Aqualog C; Horiba Co., Edison, NJ) to simultaneously collect UV–Vis absorbance and florescence spectra. Run parameters were excitation: 240–600 nm in 3 nm increments; emission: 212–608 nm by 3 nm bandpass; integration time = 0.25 s. Samples with absorbance greater than 0.6 at  $\lambda$  = 254 were diluted to satisfy the assumption of detector linearity required by modeling (Stedmon and Bro 2008; Lawaetz and Stedmon 2009). Data post-processing and correction for inner filter effects were as described in detail by Veverica et al. (2016), following Stedmon and Bro (2008) and Lawaetz and Stedmon (2009).

Fluorescence and absorbance spectroscopic indices were calculated to characterize dissolved organic matter in all samples as previously described (Veverica et al., 2016; Rupp et al., 2019). Fluorometric indices considered in this study included an index of DOM humification (humification index; HIX; Zsolnay et al., 1999; Ohno, 2002; cf. Zsolnay, 2002). Higher HIX values can generally be considered as more biologically recalcitrant or aromatic in boreal systems (Chen et al., 2011; Frey et al., 2016; Herzsprung et al., 2017). We calculated an index of free or bound proteins (tyrosine or tryptophan index; TI; Fellman et al., 2010; cf. Chen et al.,

2003). Higher TI values are considered relatively labile in comparison to the humified products associated with HIX (cf. Cory and Kaplan, 2012). We also calculated the Spectral slope ratio (Sr), an absorbance index of photodegradation, molecular weight (Helms et al., 2008) and source (Chen et al., 2011). The Sr is also an inverse indicator of relatively labile carbon compounds, with lower values associated with more bioavailable DOM (Frey et al., 2016).

# **Statistics and Analyses**

We examined WT treatment effects, flooding, and their interaction (WT treatment x flooded or not flooded state) on CO<sub>2</sub> flux data (ER, GPP, NEE), algal biomass, and DOM composition variables (HIX, Sr, TI). We collected these data throughout the growing seasons of 2016 (pre-flood; four campaigns) and 2018 (after continuous flooding; five campaigns). We used a generalized linear mixed model approach to examine differences in CO2 flux and DOM composition data prior to flooding and in the flooded state; this approach enables statistical models to be fit to data where the response is not necessarily normally distributed (PROC GLIMMIX; SAS version 9.4, SAS Institute, Cary, North Carolina, United States). Sampling campaign was treated as a random effect to account for repeated measures. The distributions of response variables were evaluated in Kolmogorov-Smirnov tests using the UNIVARIATE procedure, and were assigned as gaussian or lognormal as appropriate. Differences in DOC and TDN concentrations were explored using a mixed effects model via the "nlme" package in R statistical software version 3.4.2 (R Core Development Team, 2017). The WT treatment was used as a fixed effect, while sampling date was a random effect to account for repeated measures. The TDN values were transformed using the power-based Tukey Ladder of Powers transformation via the "rcompanion" package (Mangiafico, 2018; TDN 10 cm below water surface lambda = 2.2; TDN 20 cm below peat surface lambda = 1.5). We explored relationships with algal biomass (chlorophyll a) and ER, Sr, and TI with linear regression. Type-III tests of fixed effects and post hoc comparisons of least squared means tests across treatments were considered significant at  $\alpha$  = 0.05. All data used in these analyses are available in the public domain via the Bonanza Creek Data Catalog (https://www.lter. uaf.edu/data/data-catalog).

#### **RESULTS**

# **Change in Carbon Balance**

Across all treatments, there were changes in ER and GPP in response to flooding that resulted in interactions between drainage history and flooding for net C balance (**Table 1**). Prior to flooding, the raised water table (WT) treatment had marginally lower ER (2.79  $\pm$  0.14 CO $_2$  µmol m $^{-2}$  s $^{-1}$ ) than the lowered (3.25  $\pm$  0.30 CO $_2$  µmol m $^{-2}$  s $^{-1}$ ) or control (4.10  $\pm$  0.54 CO $_2$  µmol m $^{-2}$  s $^{-1}$ ) treatments (F = 3.31, p = 0.08), though there were no differences in ER among treatments after flooding (mean of 3.28  $\pm$  0.63 CO $_2$  µmol m $^{-2}$  s $^{-1}$ ; F = 0.75, p = 0.60; cf. **Figure 2**).

**TABLE 1** | Results of mixed effects models testing the effects of experimental water table treatment, flooding, and their interaction on ecosystem respiration (ER), gross primary production (GPP) and net ecosystem exchange (NEE). Model numerator (Num.) and Denominator (Den.) degrees of freedom (DF) are provided.

Model	Effect	Num. DF	Den. DF	F	p
ER	Treatment plot	2	14	5.330	0.019
	Flooded state	1	18	9.090	0.008
	Plot x flooded state	2	14	1.080	0.365
GPP	Treatment plot	2	10	1.900	0.200
	Flooded state	1	5	6.310	0.054
	Plot x flooded state	2	10	4.280	0.045
NEE	Treatment plot	2	10	0.250	0.786
	Flooded state	1	5	9.340	0.028
	Plot x flooded state	2	10	4.090	0.050

In contrast, GPP was reduced (less uptake, less negative) following flooding in both the control ( $-8.22 \pm 0.65$  vs.  $-5.33 \pm 0.99 \text{ CO}_2 \, \mu\text{mol m}^{-2} \, \text{s}^{-1}$ ) and raised (-7.19  $\pm 0.33 \, \text{vs.}$   $-4.18 \pm 0.99 \, \text{CO}_2 \, \mu\text{mol m}^{-2} \, \text{s}^{-1}$ ) treatments, whereas in the lowered treatment GPP did not decrease with flooding  $(-6.36 \pm 0.82 \text{ vs. } -6.07 \pm 0.34 \text{ CO}_2 \, \mu\text{mol m}^{-2} \, \text{s}^{-1})$ , resulting in an interaction between WT treatment and flooding (Table 1; F = 4.28, p = 0.045). As a result of these processes the site was a stronger sink of C prior to flooding (mean NEE  $-3.88 \pm 0.67$  $CO_2 \, \mu mol \, m^{-2} \, s^{-1}$ ) than after flooding (-1.18 ± 0.87)  $CO_2 \mu mol m^{-2} s^{-1}$ ), with the lowered treatment becoming a stronger sink, and the raised treatment being a source of C on one measurement day with flooded conditions (Figure 2B). The change in the C sink-strength with flooding in the raised treatment contributed to a weak interaction between water table treatment and flooding in explaining variation in NEE (**Table 1**; F = 4.09, p = 0.05).

In 2018, mean [DOC] obtained from 10 cm beneath the surface of the water table was significantly lower in the raised WT treatment (32  $\pm$  1.4 mg  $L^{-1}$ ) than in both the lowered (36  $\pm$ 

0.4 mg L<sup>-1</sup>) or control (39  $\pm$  0.5 mg L<sup>-1</sup>) treatments (F = 16.53, p < 0.001). Mean [DOC] obtained from 20 cm beneath the surface of the peat was also lower in the raised WT treatment (56  $\pm$  3 mg L<sup>-1</sup>) than in both the lowered (93  $\pm$  7 mg L<sup>-1</sup>) or control (80  $\pm$  2 mg L<sup>-1</sup>) treatments (F = 29.88, p < 0.001). There were no WT treatment effects on mean [TDN] obtained from 10 cm beneath the surface of the water table (0.5  $\pm$  0.1 mg L<sup>-1</sup>; F = 0.73, p = 0.49). Mean [TDN] obtained from 20 cm beneath the surface of the peat was lower in the raised WT treatment (1.1  $\pm$  0.1 mg L<sup>-1</sup>) than in both the lowered (2.8  $\pm$  0.2 mg L<sup>-1</sup>) or control (2.1  $\pm$  0.2 mg L<sup>-1</sup>) treatments (F = 20.61, p < 0.001).

# **Algal Biomass and Carbon Quality**

Algal biomass (chlorophyll a) was significantly higher in the control (mean  $\pm$  standard error; 2.61 mg m<sup>-2</sup>  $\pm$  0.46) and lowered (2.57 mg m<sup>-2</sup>  $\pm$  0.38) treatment plots compared with the raised (1.55 mg m<sup>-2</sup>  $\pm$  0.22) treatment (F = 4.24, p = 0.04). Ecosystem respiration and the index of free or bound-in proteins (TI) increased with algal biomass, while the degree of DOM photodegradation or molecular weight (Sr, or inversely related to humification, or HIX) declined with algal biomass (**Figure 3**). After accounting for changes in chlorophyll a content in mixed models (Type III tests of fixed effects), WT treatment also had significant effects on Sr (F = 6.50, p = 0.03), and marginally significant treatment effects on ER and TI (F = 3.12, p = 0.09 and F = 3.50, p = 0.08, respectively).

# **DOM Quality Before and After Flooding**

Indices of DOM quality changed as a function of WT treatment and flooding. Prior to flooding (2016), DOM humification (HIX) at 20 cm in the peat was greater in the lowered water table treatment than in the raised WT treatment (F = 8.09, p = 0.004; **Figure 4A**). Following flooding, humification was much more variable at the lowered WT treatment, and was only significantly different (lower value) in the raised treatment

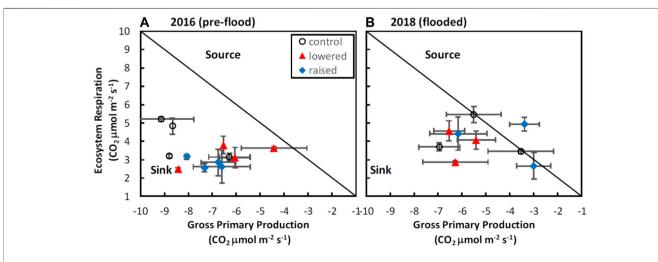


FIGURE 2 | Changes in ecosystem carbon balance prior to flooding, and after two years of continuous flooding. Depicted are means for a given campaign by water table treatment plot. Error bars represent standard errors of the mean values.

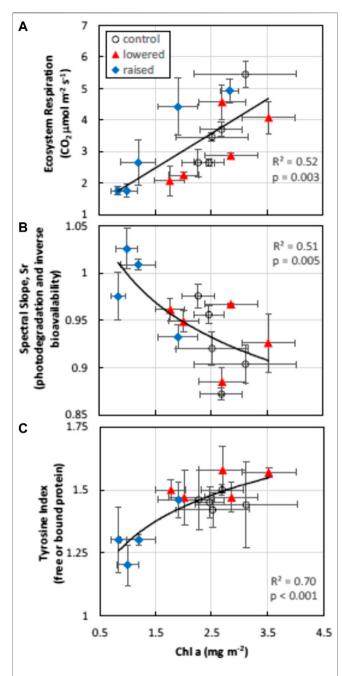


FIGURE 3 | Algal biomass (chlorophyll a content) was strongly related to changes in ecosystem respiration (top pane) measured with floating chambers during flooding, the bioavailability of DOM (inversely related to spectral slope ratio, Sr, middle pane) and the content of free or bound protein (TI, bottom pane). Depicted are daily means by water table treatment plot in the flooded year (2018). Error bars represent standard errors of the mean values.

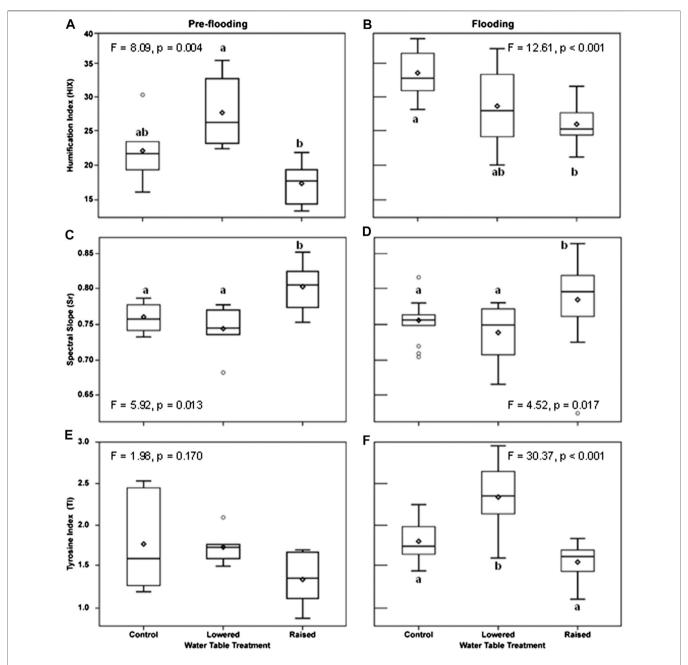
(**Figure 4B**; p < 0.001). Overall, the degree of humification was higher, in the flooded year, in the control and lowered plots (year x WT treatment interaction, F = 9.84, p < 0.001). In contrast, Sr, an indicator of photodegradation and an inverse indicator of molecular weight and bioavailability, was higher in the raised WT treatment than in the other treatments (**Figures 4C**, **D**), and there was no difference between the flooded and non-flooded years (p =

0.70). The index of free or bound-in proteins (TI) did not vary by WT treatment prior to flooding (Figure 4E; p = 0.17), but was significantly greater in the lowered WT treatment, and less in the raised treatment following flooding (Figure 4F; p < 0.001). Moreover, TI was greater overall in the flooded year than prior to flooding (F = 17.67, p < 0.001) and there was a marginal year x WT treatment interaction (F = 2.92, p =0.06). These changes in DOM quality measured at 20 cm within the peat pore water were maintained in DOM collected within the water column two years after persistent flooding (10 cm from water table surface), with the raised plot being different from the other treatments (Figure 5). With flooding, DOM in the water column of the raised WT treatment exhibited less bioavailability or increased photodegradation (Sr; F = 6.17, p = 0.005), a lower degree of humification (HIX; F = 11.29, p <0.001) and less proteinaceous compounds (TI; F = 8.12, p =0.001).

# DISCUSSION

# **Carbon Balance With Drainage History and Flooding**

In conditions of variable hydrology, lowering the water table generally promotes an increase in both ER and GPP, which has made it difficult to ascribe single mechanisms for the sign of net C balance with hydrologic change (Weltzin et al., 2003; Strack and Waddington, 2007; Ballantyne et al., 2014). A major difference between previous studies and this current work, and fens in general, was the prolonged period of saturation where water table remained above the peat surface. Prior work at this site had shown no significant WT treatment effects on ER (Chivers et al., 2009; Olefeldt et al., 2017), but was not able to investigate WT treatment effects on C cycling when the site was flooded. More recent work has shown that the site as a whole exhibited higher ER in years when flooding occurred from the spring season onward (Euskirchen et al., 2020). This work showed that with persistent flooding, the previously raised WT treatment exhibited higher ER relative to GPP, with the most parsimonious explanation being limitations to the production of algal biomass (cf. Figure 3). While prior work had also shown the ability of algae to capitalize on the nutrients released from increased decomposition in the lowered WT treatments when flooded (DeColibus et al., 2017), this current work demonstrated that algae also affected ecosystem C balance. Interestingly, the increase in labile C supply in response to algal production (e.g., higher TI) did not result in a significant increase in the relative amount of heterotrophic respiration (cf. Wyatt et al., 2019), which certainly warrants further investigation as to the limiting factors of ER when the ecosystem is flooded. For example, the protein-like fluorescence in this study may not be reflecting the pool of bioavailable DOM (Maie et al., 2007). These findings refine our prior hypothesis (Olefeldt et al., 2017) that the lack of a response in ER to WT treatments reflects a shift in the component flux contributions to ER, from autotrophic respiration in wet years to heterotrophic respiration in dry



**FIGURE 4** Indexes of peat pore water (20 cm into the peat) character before flooding in 2016 **(A,C,E)** and after continuous flooding, measured in 2018 **(B,D,F)**. Depicted are seasonal means by water table treatment plot. An index of DOM humification (HIX) was higher in the flooded year (b, F = 44.09, p < 0.001), and there was a year × treatment interaction (F = 8.84, p < 0.001). An inverse index of DOM bioavailability and molecular size (Spectral Slope Ratio, Sr) did not vary by year (F = 0.15, p = 0.70), but was higher in the raised treatment **(C,D)**. An index of peat pore water free or bound protein (TI) was higher in the flooded year (F = 17.67, p < 0.001), and the lowered treatment was higher than the raised treatment **(F)**. Letters indicate significant differences in post-hoc comparisons of means within a treatment year (flood state).

years, and demonstrate that an algal-mediated increase in GPP in the lowered WT treatments can attenuate net C losses during flooding.

# **Insights Gained From DOM Composition**

The DOM absorbance and fluorescence spectroscopy provided insights into C cycle processes within the peat prior to and after two years of flooding. Prior to flooding, pore water was more

humified in the lowered WT treatment, which is consistent with the accumulation of water-soluble products of increased decomposition in a broader, more aerobic region above the water table, or acrotelm (Strack et al., 2008; Tfaily et al., 2018). HIX remained higher (and yet more variable) in the lowered WT treatment upon being flooded, which is consistent with prior work demonstrating that a sustained HIX upon rewetting drained peatland soil represents an increased amount of oxygen-rich and relatively unsaturated (O/C >

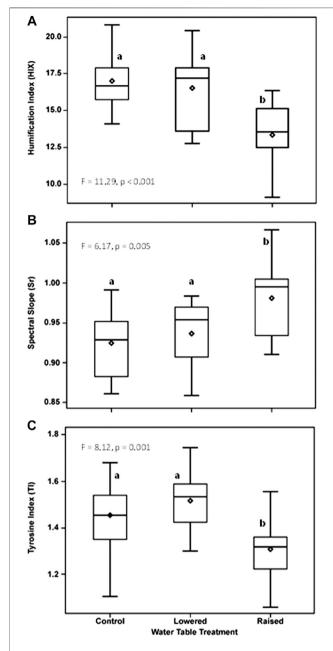


FIGURE 5 | Water table treatment differences in an index of humification (HIX, (A), an index of photodegradation or an inverse index of bioavailability (Sr, (B), and an index of free or bound protein (TI, (C) of water taken 10 cm below the surface of the water during flooding throughout the 2018 growing season. Depicted are seasonal means by water table treatment plot. Letters indicate significant differences in post-hoc comparisons of means.

0.5; H/C < 1.0) DOM (Herzsprung et al., 2017). The lower HIX values in the raised WT treatment, regardless of flooding history, suggest less accumulation of the products of peat mineralization. Lower HIX is consistent with the increased Sr (Chen et al., 2011) observed at the raised WT treatment, and could also be reflecting increased photodegradation of the DOM as opposed to biological degradation of the DOM (Helms et al., 2008; Wilske et al., 2020). Increased photodegradation would be facilitated by the water table

being at or slightly above the peat surface in the raised WT treatment. The potential for DOM photodegradation to CO2 could be contributing to the lower C sink strength in the raised WT treatment. In turn, DOM with relatively high Sr exhibits low bioavailability (Frey et al., 2016). In contrast, TI—an index of free or bound proteins, is generally thought to be highly bioavailable (Cory and Kaplan, 2012). There were no WT treatment effects on soil TI values prior to flooding, but significant increases in the lowered WT treatment and decreases in the raised WT treatments were measured when flooded (Figure 4E, F; Figure 5). Taken together, these indices indicate that the raised WT treatment had less bioavailable DOM within the peat (higher Sr, low TI) than the lowered WT treatment, even though the lowered WT treatment also had a fraction of more humified DOM (higher HIX). A parsimonious explanation for increased HIX and decreased Sr with increasing TI is the increase in algal production and algal exudation in response to the increased nutrients released in the lowered WT treatment relative to the raised WT treatment (Wyatt et al., 2012), while also having more water-soluble products from increased decomposition in the lowered WT treatment prior to flooding (Kane et al., 2010). Moreover, it could be that the proteinlike fluorescence in this case reflects higher molecular weight lignin polyphenols (Herns et al., 2009; Stubbins et al., 2014), and more detailed analysis of the presence of aromatic amino acids in this system is certainly warranted.

These results are consistent with prior work demonstrating the ability of algae to capitalize on the "legacy effect" of prior drainage and nutrient release (Wyatt et al., 2012; DeColibus et al., 2017), and further demonstrate that the elevated nutrient legacy effects on algal production can persist after two consecutive years of flooding. While vascular plants are a significant component of NEE (Rupp et al., 2019), photosynthetic capacity (NDVI) has been shown to be lower in the lowered WT treatment during flooded conditions (McPartland et al., 2019), which further suggests a different mode of primary production to be driving C uptake when flooded. The duration of flooding is an important consideration for C cycling and algal production. For example, relatively short-term (<40 days) wet events did not generate lag effects in C cycling, and the lowered WT treatment in wet years exhibited similar GPP and NEE to that of the raised WT treatment in dry years (Olefeldt et al., 2017). After persistent flooding, however, the lowered WT treatment became the strongest C sink, whereas the raised WT treatment became a source (e.g., Table 1; Figure 2). However, it is not clear how long the "legacy effect of drought" can continue to fuel increased algal production. For example, nutrient co-limitations or competition with vascular plants may override any nutrient legacy effect on algal production over the span of decades (Lougheed et al., 2011; Lougheed et al., 2015). Flooding events at this rich fen site have been increasing over time, commensurate with the study region becoming a greater source of C to the atmosphere (Euskirchen et al., 2020). If periods of inundation increase in duration (i.e., become more lake-like), as has been observed in other regions of interior Alaska (Jorgenson et al., 2001), these data suggest rich fens could become a weaker sink for C, with

reduced algal production likely being a contributing factor. These findings indicate that incorporating an understanding of flooding history and its effects on algal production are important in assessments of C storage in boreal fen peatlands.

#### DATA AVAILABILITY STATEMENT

The datasets presented in this study can be found in online repositories. The names of the repository/repositories and accession number can be found below: All data used in these analyses are available in the public domain via the Bonanza Creek Data Catalog (https://www.lter.uaf.edu/data/data-catalog).

# **AUTHOR CONTRIBUTIONS**

All authors contributed to the conception of this study. EK lead the analyses and writing, with input from all authors. CD lead field activities during flooding (2018), with input from all authors. KW and AR lead algal biomass methods and analyses.

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