

**Nutrient enrichment effects are conditional on upstream nutrient concentrations:
Implications for bioassessment in multi-use catchments**

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Abstract: Human impacts on stream ecosystems are expected to intensify with population growth and climate change. Decisive information on how stream communities respond to cumulative human impacts is therefore integral for protecting streams draining multi-use catchments. To determine cumulative influences of nutrient enrichment and assess more nuanced approaches for the evaluation of human impacts, we present results from one factorial and two gradient assessment designs applied to benthic algae and macroinvertebrate data from 14 mid-order streams in southern Ontario, Canada with pre-existing human impacts (i.e., sewage effluent and agriculture). We found that among stream variability in ecological indicators measured downstream of sewage effluent outfalls confounded our generalized factorial assessment and provided inconclusive information on a known human impact. Despite our gradient assessment also not having strong statistical support, accounting for the extent of nutrient enrichment associated with differences in sewage effluent and agricultural inputs revealed that larger

longitudinal changes in stream communities were associated with increased nutrient enrichment. However, re-weighting our nutrient enrichment gradient based on upstream nutrient concentrations to account for nonlinearities in the response of stream communities to nutrient enrichment produced more robust assessment results that were consistent with predicted effects of nutrients on stream ecosystems. Thus, while our factorial assessment suggests that the communities are resistant to nutrients from cumulative human impacts, our targeted gradient assessment demonstrates that the effects of nutrient enrichment are highly conditional on upstream ecosystem conditions. Future assessments may need to go beyond traditional approaches (i.e., impact presence/absence) and more explicitly consider the environmental stressors and their associated complexities related to the impact under investigation.

Keywords: diatoms; macroinvertebrates; nitrogen; phosphorus; agriculture; sewage; wastewater

1. Introduction

Streams support and regulate important ecosystem processes (e.g., nutrient cycling, water purification, and waste assimilation) and provide provisional and cultural services (e.g., food, water, and recreation) to sustain societal needs (Lowe & Likens, 2005). However, human impacts on stream ecosystems are diverse (e.g., habitat alteration and pollution) and are expected to increase with continued population growth and climate change (Søndergaard & Jeppesen, 2007). Thus, there is an ongoing need to assess the impacts of human activities to provide managers and policy makers with scientifically defensible information on how to better protect the structure and function of stream ecosystems.

One of the most effective study designs to evaluate the effects of an impact (e.g., impoundment, point source pollution, and land use change) while controlling for natural

variability is the before-after control-impact (BACI) assessment (Green, 1979; Smokorowski & Randall, 2017). BACI assessments compare the temporal change in ecosystem conditions after exposure to an impact to the natural temporal variability in an unexposed stream and can readily identify the potential effect of an impact if its timing and location are known, and pre-exposure monitoring data are collected. However, researchers do not often have sufficient advance notice of when or where an impact will occur to collect pre-exposure data. Thus, suboptimal (e.g., before-after only, control-impact only) or modified (e.g., upstream-downstream control-impact) BACI assessments are often used to evaluate the effect of human impacts, despite producing less robust assessment results (Smokorowski & Randall, 2017). Moreover, BACI assessments are usually limited by the use of spatial replicates to measure ecosystem conditions within individual control-impact streams (Hurlbert, 1984), and while temporal replicates can be employed to eliminate spatial pseudo-replication (Stewart-Oaten et al., 1986), inference on the potential effect of an impact is restricted to the studied location alone (Underwood, 1991; 1994). Generalizability of cause-effect relationships from BACI assessments can be strengthened with the inclusion of multiple control-impact sites as independent replicates (*aka* multiple-BACI; Downs et al., 2002). However, in highly developed regions locating a set of suitable control-impact streams with similar environmental conditions can be near impossible and remains a common challenge for impact assessments (Stoddard et al., 2006; Herlihy et al., 2008).

Past studies have shown that although BACI assessments employ a factorial approach to provide information on how the presence of an impact influences stream ecosystems, quantitative differences in environmental stressors (e.g., chemical and sediment concentrations, and thermal pollution) associated with an impact can govern the size of the perturbation from prior ecosystem conditions (Welch et al., 1992; Chambers et al., 1997; Scrimgeour & Chambers,

2000; Bowman et al., 2005). For example, Quinn and Hickey (1993) observed that variability in the difference between benthic macroinvertebrate assemblages sampled upstream and downstream of eight sewage treatment lagoons was associated with the increase in organic matter concentration from the differential addition of effluent. Thus, the factorial evaluation of an individual impact derived from multiple study streams with differing degrees of human influence may be confounded by increased variability in the response of stream communities to specific environmental stressors. Variability in the response of ecosystem conditions to a common impact can result in decreased confidence in factorial assessment results, particularly in assessments with small sample sizes. Moreover, bioassessment in highly developed regions can be further complicated by the presence of multiple stressors from independent human impacts that may interact (i.e., additively, synergistically, or antagonistically) and accumulate along the stream continuum to contribute to a cumulative extraneous effect on ecosystem conditions (Seitz et al., 2011; Galic et al., 2018). Increased variability from cumulative and multiple stressors can therefore make it difficult to select well-matched control-impact streams that can provide decisive assessment information on the potential effects of an individual human impact.

In recognition that human impacts can have diverse effects on stream ecosystems, many studies have elected to use human activity gradients (e.g., land cover and volume of effluent discharged) to evaluate impacts at the catchment-scale or have taken a gradient approach to isolate the effects of specific stressors (e.g., nutrient and pesticide concentrations; Davies & Jackson, 2006; Yates & Bailey, 2010a; Hausmann et al., 2016). A gradient approach infers that stream ecosystems will respond to future impacts in direct accordance with identified spatial patterns in stressor-response relationships (Jarvie et al., 2013). Gradient studies have provided a more targeted understanding of how quantitative differences in a stressor affect stream

communities and many have reported that stream communities respond nonlinearly to stressor gradients (Allan, 2004; Clements & Rohr, 2009; D’Amario et al., 2019). For example, stream communities often exhibit a threshold response to increased environmental stress where little to no further change in community structure is observed after a critical point has been surpassed (Groffman et al., 2006; Dodds et al., 2010; Hilderbrand & Utz, 2015). However, because human impacts in highly developed regions often occur on streams exposed to existing environmental stressors, these nonlinearities in stressor-response relationships have the potential to mask the cumulative effects of additional human impacts on stream conditions in factorial before-after comparisons (Clements et al., 2016). Indeed, although local stream communities can be resistant to additional degradation, cumulative human impacts may decrease the resilience of stream communities and have farther-reaching consequences throughout the stream network. Moreover, ill-informed management decisions from suboptimal impact assessments may promote environmental degradation across larger spatial scales. Thus, in order to improve and further protect imperilled stream ecosystems, there is a need for a more robust understanding of how cumulative impacts from human activities influence streams draining highly developed regions.

Nutrient (i.e., nitrogen and phosphorus) enrichment is a pervasive chemical stressor that affects many streams draining developed regions (Dodds & Smith, 2016; Wurtsbaugh et al., 2019). Human impacts from point (e.g., sewage effluent) and nonpoint (e.g., agriculture) sources of pollution can cumulatively increase instream nitrogen and phosphorus concentrations and degrade the structure and function of stream ecosystems through accelerated eutrophication (Mainstone & Parr, 2002; Withers & Jarvie, 2008). Eutrophication is characterized by excessive primary production (i.e., sestonic and benthic algae, and macrophytes) and while enriched nutrient concentrations can increase the biomass of primary producers, past studies have often

reported threshold or breakpoint type relationships between stream degradation by eutrophication and nutrient concentrations (Evans-White et al., 2013; Jarvie et al., 2013; Heiskary & Bouchard, 2015). The ecological implications of stream nutrient enrichment have been well established (e.g., Biggs, 2000; Friberg et al., 2010; Woodward et al., 2012) and because point sources can be readily identified, there has been a substantive amount of research on the nutrient impacts of treated sewage effluents (see Hamdhani et al., 2020). For example, many studies have reported an increase in tolerant and decrease in sensitive benthic macroinvertebrate taxa downstream of sewage outfalls compared to upstream reference reaches (e.g., Quinn & Hickey, 1993; Ortiz et al., 2005; Englert et al., 2013). Assemblage-level differences have also been observed in benthic algal communities (e.g., Chambers et al., 1997; Scrimgeour & Chambers, 2000; Bowman et al., 2005). Most of these past studies occurred in streams with either low upstream nutrient concentrations or a pronounced spike in concentration due to input of sewage with little or no treatment. However, in developed regions sewage effluent is often discharged into streams already enriched with nutrients from upstream nonpoint sources, such as agricultural lands, resulting in more pronounced cumulative effects and nonlinear responses. The point source discharge of effluent into agricultural streams thus provides an opportunity to determine if the cumulative effects of nutrient enrichment can be detected with general and more nuanced assessment approaches that account for nonlinearities associated with environmental stressors.

Our goal was to demonstrate the challenges of impact assessments in highly developed regions by evaluating the cumulative effects of nutrient enrichment on instream ecological communities using both a general and more targeted study design. First, a multiple before-after (upstream-downstream) control-impact assessment was used to conduct a factorial evaluation of the generalized impact that treated sewage effluent has on ecological communities (i.e., benthic

diatoms and macroinvertebrates) of agricultural streams. Second, a targeted nutrient enrichment gradient assessment was used to evaluate the association between upstream-downstream differences in stream communities and the extent of nitrogen and phosphorus enrichment from human nutrient sources in stream catchments. Third, we tested if accounting for nonlinear stressor-response relationships when evaluating cumulative effects would provide a more robust understanding of how increased nutrient enrichment influences ecological communities in streams draining highly developed regions.

2. Methods

2.1 Study Area and Design

We assessed the cumulative impact of nutrients from agriculture and sewage effluent on 14 streams in the mixed-wood plains ecozone (Crins et al., 2009) of southern Ontario, Canada (Figure 1). Southern Ontario is characterized by a humid, temperate climate with an average annual temperature between 4.9 and 7.8 °C and total precipitation from 759 to 1087 mm (Crins et al., 2009). The underlying physiography consists of post-glacial surface geologies dominated by mixed agricultural land covers (i.e., crop and livestock operations) with patches of natural (i.e., forest and wetland) and urban areas throughout the landscape (Yates & Bailey, 2010b).

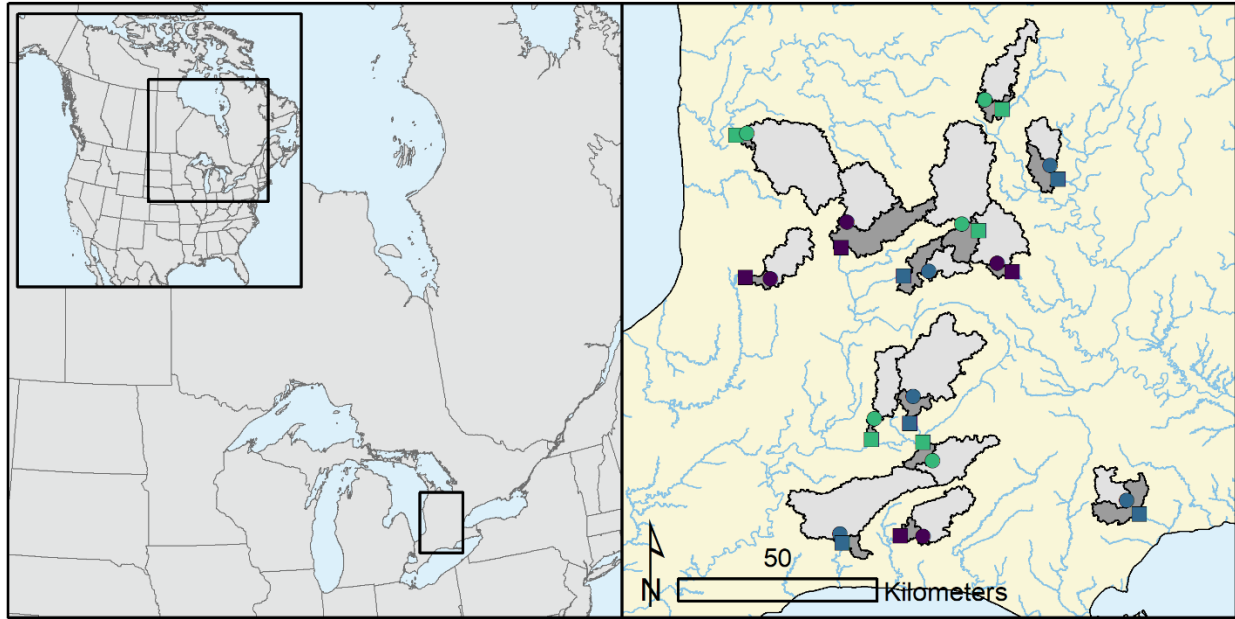


Figure 1: Location of the 14 study streams in southern Ontario, Canada. Upstream sampling locations are symbolized by circles and light grey catchments. Downstream sampling locations are symbolized by squares and dark grey catchments correspond to the additional drainage area captured by the nested stream reach. Green, purple, and blue symbols correspond to watersheds exposed to agriculture, agriculture + passively treated lagoon sewage effluent, and agriculture + actively treated mechanical sewage effluent, respectively.

Study streams were selected to have comparable catchment sizes, land cover composition, and surface geology (glacial till), but differ in their exposure to anthropogenic sources of nutrients (Table 1). All 14 streams were exposed to nutrients from agricultural activities upstream in study catchments with 9 of those streams also being exposed to nutrients from point sources of municipal sewage effluent. Of the 9 effluent receiving streams, 5 streams were exposed to actively treated sewage effluent from mechanical sewage treatment plants and 4 streams were exposed to passively treated sewage effluent from sewage lagoons (a.k.a. wastewater stabilization ponds). Operational differences between active and passive sewage treatment plants are described in Pearce et al. (2020).

To evaluate the potential impact of anthropogenic nutrient enrichment, two paired sampling sites were established on each study stream. Sites were located upstream and downstream of identified point sources or a comparable longitudinal distance apart along

agricultural streams (without point sources). Catchment size was on average (\pm standard deviation) $170 \pm 107 \text{ km}^2$ for upstream sites and $198 \pm 123 \text{ km}^2$ for downstream sites. Average (\pm standard deviation) land cover composition was largely similar between upstream and downstream study catchments (agriculture upstream = $85 \pm 6 \%$ versus downstream = $83 \pm 8 \%$; urban upstream = $2 \pm 2 \%$ versus downstream = $4 \pm 4 \%$; natural upstream = $13 \pm 6 \%$ versus downstream = $13 \pm 5 \%$).

Table 1: Properties of upstream and downstream sampling locations on the 14 study streams in southern Ontario, Canada. Distance (Dist.) between sampling locations reflects longitudinal distance along the stream network and downstream distance of sampling locations from identified point sources is provided in brackets. Agriculture (Agri.), urban (Urb.), and natural (Nat.) land cover was summarized from the Agriculture and Agri-Food Canada Annual Crop Inventory (2013) dataset.

| Stream Name | Dist. | Upstream | | | | Downstream | | | |
|---------------------------|------------------|-------------------|-------------|------------|-------------|-------------------|-------------|------------|-------------|
| | | Area | Agri. | Urb. | Nat. | Area | Agri. | Urb. | Nat. |
| | (km) | (km^2) | (%) | (%) | (%) | (km^2) | (%) | (%) | (%) |
| Agriculture | | | | | | | | | |
| Mallet River | 6.7 | 111.6 | 84.2 | 1.2 | 14.5 | 144.1 | 84.4 | 1.2 | 14.4 |
| Reynolds Creek | 5.9 | 123.4 | 85.9 | 0.7 | 12.9 | 153.0 | 86.0 | 0.8 | 12.8 |
| S. Maitland River | 4.4 | 359.4 | 87.8 | 1.1 | 10.1 | 371.5 | 87.6 | 1.1 | 10.3 |
| Upper Nith River | 6.4 | 311.8 | 89.5 | 1.3 | 9.3 | 377.2 | 88.0 | 1.3 | 10.6 |
| Waubuno Creek | 7.3 | 98.6 | 87.1 | 0.5 | 12.1 | 106.4 | 86.4 | 1.0 | 12.3 |
| <i>Mean</i> | <i>6.11</i> | <i>201.0</i> | <i>86.9</i> | <i>0.9</i> | <i>11.8</i> | <i>230.4</i> | <i>86.5</i> | <i>1.1</i> | <i>12.1</i> |
| <i>Standard Deviation</i> | <i>1.1</i> | <i>124.4</i> | <i>2.0</i> | <i>0.3</i> | <i>2.1</i> | <i>132.5</i> | <i>1.5</i> | <i>0.2</i> | <i>1.7</i> |
| Lagoon | | | | | | | | | |
| Ausable River | 7.1 (1.2) | 100.5 | 88.5 | 0.7 | 10.7 | 115.8 | 86.4 | 3.0 | 10.3 |
| Catfish Creek | 8.7 (2.3) | 119.1 | 86.9 | 2.2 | 10.8 | 141.8 | 84.3 | 5.1 | 10.2 |
| Lower Nith River | 12.2 (1.1) | 217.4 | 79.4 | 2.4 | 18.0 | 546.4 | 84.4 | 2.7 | 12.9 |
| North Thames River | 7.1 (2.6) | 166.6 | 93.7 | 0.8 | 5.5 | 323.6 | 91.4 | 2.1 | 6.4 |
| <i>Mean</i> | <i>8.8 (1.8)</i> | <i>150.9</i> | <i>87.1</i> | <i>1.5</i> | <i>11.2</i> | <i>281.9</i> | <i>86.6</i> | <i>3.2</i> | <i>10.0</i> |
| <i>Standard Deviation</i> | <i>2.4 (0.8)</i> | <i>52.3</i> | <i>5.9</i> | <i>0.9</i> | <i>5.1</i> | <i>199.1</i> | <i>3.3</i> | <i>1.3</i> | <i>2.7</i> |
| Mechanical | | | | | | | | | |
| Avon River | 6.7 (1.1) | 53.6 | 83.6 | 2.7 | 13.3 | 115.6 | 71.2 | 14.8 | 13.5 |
| Canagagigue Creek | 5.7 (1.6) | 65.4 | 85.3 | 0.6 | 13.1 | 115.9 | 81.8 | 6.3 | 11.2 |
| Kettle Creek | 4.0 (1.4) | 329.3 | 80.9 | 6.5 | 12.4 | 354.5 | 78.3 | 9.1 | 12.3 |
| Lynn River | 7.2 (1.6) | 59.1 | 67.2 | 1.8 | 30.6 | 138.0 | 61.7 | 9.0 | 29.1 |
| Middle Thames River | 9.1 (2.4) | 269.7 | 84.4 | 0.6 | 14.9 | 304.9 | 84.0 | 1.5 | 14.5 |
| <i>Mean</i> | <i>6.5 (1.6)</i> | <i>155.4</i> | <i>80.3</i> | <i>2.4</i> | <i>16.9</i> | <i>205.8</i> | <i>75.4</i> | <i>8.1</i> | <i>16.1</i> |
| <i>Standard Deviation</i> | <i>1.9 (0.5)</i> | <i>133.3</i> | <i>7.5</i> | <i>2.5</i> | <i>7.7</i> | <i>114.8</i> | <i>9.1</i> | <i>4.8</i> | <i>7.4</i> |
| Mean | 7.0 (1.7) | 170.4 | 145.4 | 3.3 | 21.1 | 236.3 | 198.2 | 8.5 | 29.0 |
| Standard Deviation | 2.0 (0.6) | 106.9 | 92.4 | 5.5 | 13.0 | 140.1 | 123.0 | 8.4 | 17.3 |

2.2 Sample Collection and Processing

Grab water samples were collected at each sampling site every three weeks from May to September of 2013 (n = 8). Samples were collected in the thalweg of each stream at 60% depth and stored at 4 °C prior to overnight transport to the National Laboratory for Environmental Testing (Environment and Climate Change Canada) for colorimetric determination of total phosphorus (TP) and total nitrogen concentrations (TN). Detection limits were 0.0005 mg P L⁻¹ for TP and 0.015 mg N L⁻¹ for TN.

Benthic diatom and macroinvertebrate assemblages were sampled in September 2013. Diatoms were sampled by collecting biofilm from 10 randomly selected cobbles throughout the defined sampling reach. Composite biofilm samples were preserved in dark bottles with Lugols iodine (~ 1% v/v) and subsampled for taxonomic identification. Biofilm subsamples were digested in 800 µL of 100% (v/v) nitric acid for 48 hours and 200 µL of hydrogen peroxide 30% (v/v) for an additional 48 hours. Digested samples were rinsed to remove nitric acid and were mounted with Naphrax® on prepared microscope slides (refractive index: 1.74; Brunel microscopes Ltd., Wiltshire, UK). Diatoms valves were enumerated (minimum 400) with use of a Reichert-Jung Polyvar microscope equipped with differential interference contrast (magnification 1250x). Diatom frustules were identified to lowest possible taxonomic, usually species, following Lavoie et al. (2008b).

Benthic macroinvertebrates were sampled following the Canadian Aquatic Biomonitoring Network (CABIN) protocols (Reynoldson et al., 2012), which consisted of a 3-minute traveling kick sample (400 micron D-frame net) over a defined sampling reach (six times the bankfull width). Macroinvertebrates were fixed with 10% buffered formalin and preserved in 75% ethanol. Collected macroinvertebrate samples were subsampled at random with the use of a

Marchant box until a minimum of 300 individuals were enumerated. Subsampled macroinvertebrates were identified to genus or family.

2.3 Data Analysis

Consistency in taxonomic resolution was achieved for benthic macroinvertebrate samples where individuals were enumerated at the family and genus levels following Velk et al. (2004). In brief, if greater than 20% of individuals in a taxon were identified to the family level, then all individuals from the lower genus levels would be included in the family level count. However, if less than 20% of individuals were identified to the family level, and all samples had at least one individual enumerated at the genus level for that taxon, then the family level data was removed from analysis. In cases where less than 20% of individuals were identified to the family level, but not all samples had individuals at the genus level, all individuals were adjusted to the family level. Following taxonomic adjustments, any taxon that was present in less than 5% of the samples was declared as rare and was removed for dissimilarity analyses, unless the number of individuals for the taxon within an individual sample was greater than 5% of total individuals counted. Removal of rare taxa was completed for both benthic diatom and macroinvertebrate assemblages. Rare taxa were not removed in the calculation of community composition indices.

2.4 Diatom and Macroinvertebrate Indices

The composition of benthic diatom and macroinvertebrate assemblages were described using common bioassessment indices (see Barbour et al., 1999; Table 2). Diatom assemblages were described by the Eastern Canadian Diatom Index (IDEC; Lavoie et al., 2014), pollution tolerance index (PTI_D; Barbour et al., 1999; Muscio, 2002), percent abundance of high nutrient taxon (%high-NP; Potapova & Charles, 2007), percent abundance of low nutrient taxon (%low-

228 NP; Potapova & Charles, 2007), taxon richness (s_D), and Bray-Curtis dissimilarity
229 ($dissimilarity_D$). Benthic macroinvertebrate assemblages were described using the Hilsenhoff
230 Family Biotic Index (FBI; Hilsenhoff, 1998), pollution tolerance index (PTI_{BMI} ; *aka* enrichment
231 tolerance in Krynak & Yates, 2018), percent Ephemeroptera, Plecoptera, and Tricoptera
232 abundance (%EPT), percent Diptera abundance (%Dipt.), taxon richness (s_{BMI}), and Bray-Curtis
233 dissimilarity ($dissimilarity_{BMI}$). Bray-Curtis dissimilarity matrices were calculated for benthic
234 diatom and macroinvertebrate assemblages from Hellinger transformed taxon abundance data
235 with the *vegan* package in R (Oksanen et al., 2019). Pairwise dissimilarities between sampling
236 locations on each stream were used to evaluate the effect of different human impact categories.

Table 2: Description of benthic diatom and macroinvertebrate indices used to evaluate nutrient enrichment. Predicted direction of association between nutrient enrichment and each ecological indicator is denoted by positive (+) and negative (–) symbols.

| Indicator | Description | Nutrient Enrichment Effect |
|------------------------------|---------------------------------------------------------------------------------------------------------|----------------------------|
| <i>Diatoms</i> | | |
| IDECA | Eastern Canadian Diatom Index used to assess biological integrity associated with stream eutrophication | – |
| PTI _D | Metric summarizing the pollution sensitivity of taxa present in an assemblage | – |
| %high-NP | Proportional abundance of taxon associated with nutrient enriched environments | + |
| %low-NP | Proportional abundance of taxon associated with nutrient poor environments | – |
| S _D | Taxon richness or number of unique taxa at lowest possible taxonomic resolution | – |
| Dissimilarity _D | Pairwise Bray-Curtis dissimilarity between assemblages | + |
| <i>Macroinvertebrates</i> | | |
| FBI | Hilsenhoff Family Biotic Index used to assess biological integrity associated with organic pollution | + |
| PTI _{BMI} | Metric summarizing the pollution sensitivity of taxa present in an assemblage | + |
| %EPT | Proportional abundance of Ephemeroptera, Plecoptera, and Tricoptera taxa | – |
| %Dipt. | Proportional abundance of Diptera taxa | + |
| S _{BMI} | Taxon richness or number of unique taxa at lowest possible taxonomic resolution | – |
| Dissimilarity _{BMI} | Pairwise Bray-Curtis dissimilarity between assemblages | + |

2.5 Statistical Analyses

Two statistical approaches were used to evaluate the cumulative impact of nutrients from agriculture and sewage effluent on instream ecological communities. First, a multiple BACI assessment model was used to evaluate differences ($\alpha = 0.05$) in the mean of each dependent variable (TN, TP, IDECA, PTI_D, %high-NP, %low-NP, and S_D, FBI, PTI_{BMI}, %EPT, %Dipt., S_{BMI},) associated with sampling location (i.e., upstream or before, and downstream or after) and human impact (i.e., agriculture or control, and passive/active effluent or impact). Human impact was evaluated in separate analyses (1) as a three-level factor accounting for the operational

differences of passive and active sewage treatment facilities and (2) as a two-level factor where effluent treatment was amalgamated to increase statistical power. An interaction term (sampling location: human impact) was included in the model to determine if differences in measured variables between sampling locations depended on the presence/type of human impact. Stream ID was included as a random effect term to account for site nestedness. Mixed model analyses of variance for BACI assessments were performed with the *nlme* package in R version 3.6.1 (Pinherio et al., 2020). The sampling location : human impact interaction term represents the BACI contrast and is equal to an analysis of variance between the stream specific differences (downstream – upstream) in each dependent variable and human impact (i.e., agriculture, passive effluent, and active effluent). Because measures of Bray-Curtis dissimilarity correspond to stream specific differences in assemblage composition, a one factor analysis of variance was performed between the pairwise dissimilarity (dissimilarity_D and dissimilarity_{BMI}) measures for each study stream and human impact. For all models, Cohen's d effect size and 90% confidence intervals were calculated from the stream specific differences (downstream – upstream) in each dependent variable to discern differences in the cumulative effects of sewage effluent (i.e., all sewage effluent, passive sewage effluent, active sewage effluent) on agricultural rivers. Analyses were performed with the *stats* and *effsize* packages in R (R Core Team, 2020; Torchiano, 2020).

Second, the potential cumulative effects of increased nitrogen and phosphorus concentrations on the upstream-downstream differences in stream communities were evaluated through a gradient assessment approach with data from streams exposed to agriculture only and sewage effluent. Multiple regression analyses were used to determine the association ($\alpha = 0.05$) between longitudinal differences (downstream – upstream) in TP and TN (independent variables) and longitudinal differences in individual ecological indicators (dependent variables). The

272 downstream cumulative effects of TP and TN were assessed directly based on the absolute
273 difference in concentration. However, stream communities, particularly benthic algae, have been
274 commonly found to exhibit nonlinear (threshold) response to low-level nutrient enrichment (e.g.,
275 Biggs, 2000; Stevenson et al., 2008; Black et al., 2011; Taylor et al., 2014) and thus the effect of
276 increased nutrient enrichment may be conditional on upstream ecosystem conditions
277 (Lacoursière et al., 2011; Clements et al., 2016). Thus, in addition to absolute differences in
278 concentration, a separate assessment was conducted with TP and TN concentrations that were
279 inverse (reciprocal) transformed prior to differencing ($1/\text{downstream} - 1/\text{upstream}$) to increased
280 the numerical weight of nutrient inputs into streams that had lower upstream nutrient
281 concentrations (Figure 2). For example, the absolute difference in concentration would be the
282 same for a stream that increased from 0.01 mg L^{-1} TP to 0.03 mg L^{-1} TP as a stream that
283 increased from 0.05 mg L^{-1} TP to 0.07 mg L^{-1} TP, but the inverse transformed differences would
284 be about 11-fold greater for the stream with the lower upstream TP concentration. Independent
285 variables were normalized prior to analysis such that greater values correspond to increased
286 cumulative effects of nutrients. Statistical analyses were performed in R with the *stats* package
287 (R Core Team, 2020).

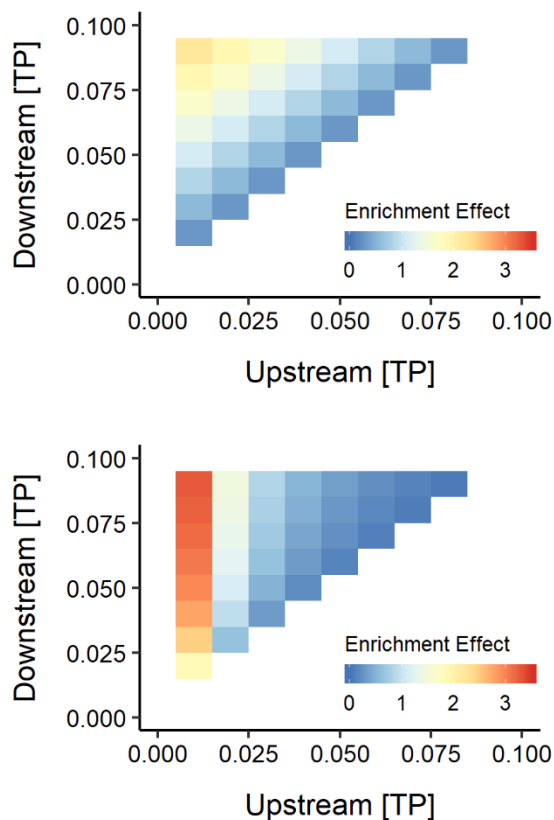


Figure 2: Illustration of the mean normalized enrichment effect of nutrients (colour gradient) based on (a) absolute concentration differencing and (b) inverse concentration differencing of hypothetical total phosphorus (TP) concentrations (mg L^{-1}) between upstream and downstream study sties.

3. Results and Discussion

3.1 Multiple Before-After Control-Impact Assessment

Nutrient concentrations in agricultural streams and at upstream sampling locations in effluent impacted streams were on average (\pm standard deviation) $0.055 \pm 0.034 \text{ mg P L}^{-1}$ for TP and $5.25 \pm 1.27 \text{ mg N L}^{-1}$ for TN (Figure 3). Past studies in southern Ontario have reported similar total nutrient concentrations in streams that drain agricultural catchments (Raney & Eimers, 2014; Thomas et al., 2018; DeBues et al., 2019; Pearce et al., 2020) and compared to regional nutrient criteria ($0.024 \text{ mg P L}^{-1}$ for TP; 1.07 mg N L^{-1} for TN) most control streams in our study had enriched nutrient concentrations (Chambers et al., 2012). Sewage effluent was

301 observed to cumulatively increase stream nutrient concentrations at downstream sampling
302 locations. Mean nutrient concentrations were 0.144 ± 0.136 mg P L⁻¹ for TP and 6.40 ± 1.68 mg
303 N L⁻¹ for TN downstream of lagoon sewage effluent, and 0.104 ± 0.085 mg P L⁻¹ for TP and 6.81
304 ± 2.64 mg N L⁻¹ for TN downstream of mechanically treated sewage effluent (Figure 3). Many
305 studies have shown increased nutrient concentrations in stream reaches that receive effluent
306 compared to upstream reaches (see Carey & Migliaccio, 2009; Hamdhani et al., 2020). However,
307 while we observed an increase in mean nutrient concentration downstream of sewage outfalls, a
308 significant ($p < 0.05$) sampling location : human impact interaction term was not detected in
309 multiple BACI assessment models of TN and TP (Table S1) suggesting that sewage effluent was
310 not impacting nutrient concentrations of agricultural streams.

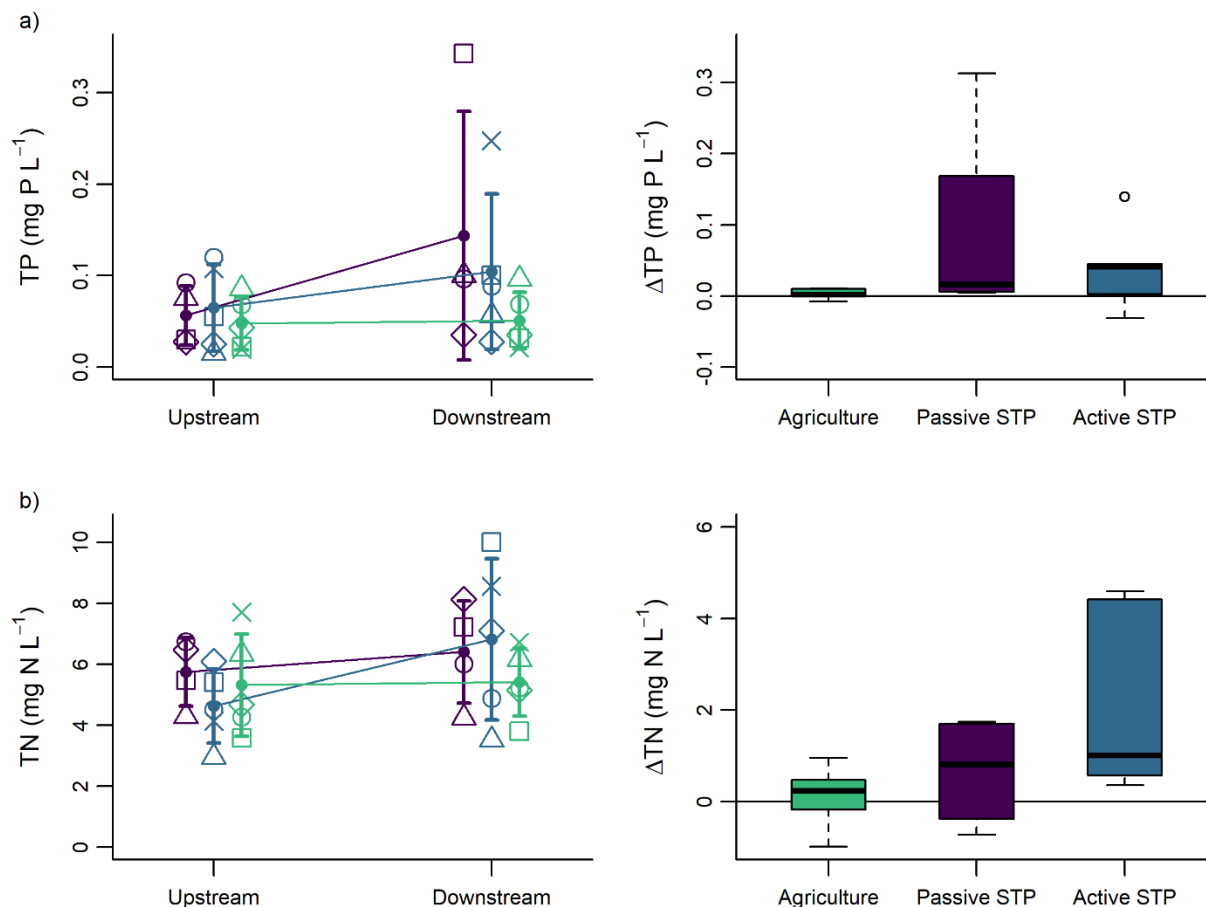


Figure 3: Mixed model analysis of variance for multiple before-after (upstream-downstream) control-impact (agriculture-sewage effluent) assessment of (a) total phosphorus (TP) and (b) total nitrogen (TN) concentrations. Green, purple, and blue colours correspond to streams exposed to agriculture, agriculture + passively treated lagoon sewage effluent, and agriculture + actively treated mechanical sewage effluent, respectively. Open symbols correspond to nutrient concentrations for individual study streams and filled points represent the mean \pm one standard deviation within each human impact category. Longitudinal differences (downstream – upstream) in nutrient concentrations are summarized by boxplots (min, 25th percentile, 50th percentile, 75th percentile, max) for each human impact category.

Longitudinal differences (downstream – upstream) in nutrient concentrations were highly variable among streams that received sewage effluent (-0.03 to 0.31 mg P L^{-1} for TP; -0.73 to 4.60 mg N L^{-1} for TN) compared to agricultural streams (-0.008 to 0.01 mg P L^{-1} for TP; -0.99 to 0.95 mg N L^{-1} for TN) that received no point source inputs of effluent (Figure 3). Variable water quality impacts of sewage effluent have also been reported in past studies (Quinn & Hickey, 1993; Chambers et al., 1997; Scrimgeour & Chambers, 2000; Bowman et al., 2005) and are often

associated with operational differences (e.g., effluent quality, compliance limits, discharge volume, and dilution factor) in sewage treatment facilities (Carey & Migliaccio, 2009). For example, Welch et al., (1992) observed a 1 to 103-fold increase in dissolved reactive phosphorus and a 1 to 176-fold increase in dissolved inorganic nitrogen among seven streams in New Zealand that received effluent from point sources. Although primary operational differences in sewage treatment facilities were accounted for in our three-level study design (i.e., passive versus active treatment), high variability in the sewage nutrient load and its dilution appears to have confounded our ability to detect a generalizable cumulative effect of sewage effluent on the nutrient concentrations of agricultural streams when summarised categorically.

Among the 12 ecological indices evaluated in our study (IDEC, PTI_D, %high-NP, %low-NP, *s_D*, dissimilarity_D, FBI, PTI_{BMI}, %EPT, %Dipt., *s_{BMI}*, and dissimilarity_{BMI}), no significant BACI interaction effects were observed in three-level human impact models (Table S1). Our finding of no difference in any of the ecological indices suggests that the discharge of sewage effluent had no influence on longitudinal patterns in benthic diatom and macroinvertebrate assemblages of agricultural streams. All upstream reaches in our study, as well as downstream reaches in streams without point sources, were affected by agricultural nutrient enrichment and thus may have been comprised of tolerant communities resistant to further changes from the addition of sewage effluent (Lavoie et al., 2008a). Although it is not uncommon to observe community tolerance to cumulative stress (Gücker et al., 2006; Burdon et al., 2016; 2019), like TP and TN concentrations, we observed high variability in the differences of measured ecological indicators between sampling locations within all three human impact categories (Figure 4; Figure S-1). Given the known effects of sewage effluent on instream biological communities (e.g., Quinn & Hickey, 1993; Scrimgeour & Chambers, 2000; Bowman et al., 2005;

Englert et al., 2013), it appears more likely that high variability in measured ecological indicators at the categorical level masked any generalizable impact of exposure to sewage effluent.

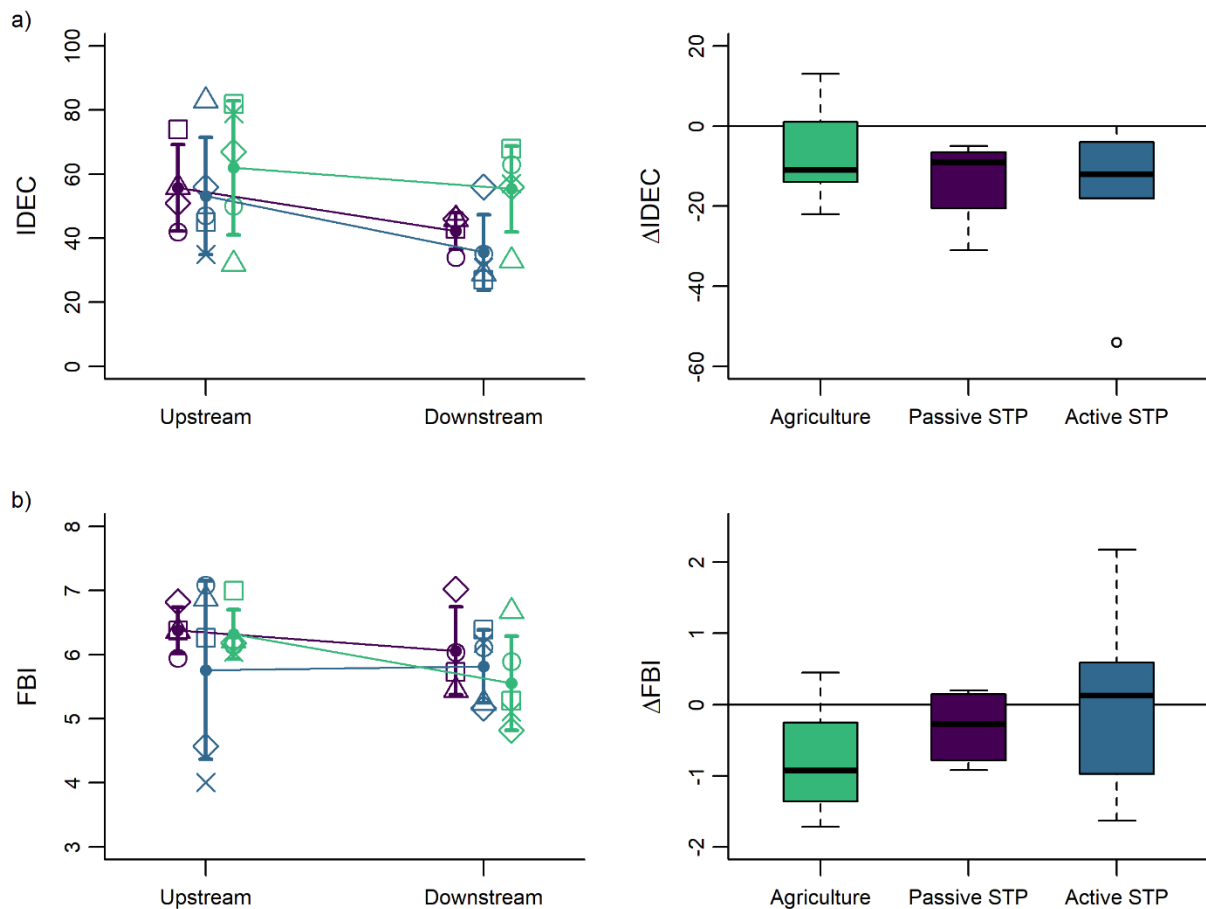


Figure 4: Mixed model analysis of variance for modified before-after (upstream-downstream) control-impact (agriculture-sewage effluent) assessment of (a) the eastern Canadian diatom index (IDEC) and (b) the Hilsenhoff family biotic index (FBI). Green, purple, and blue colours correspond to streams exposed to agriculture, agriculture + passively treated lagoon sewage effluent, and agriculture + actively treated mechanical sewage effluent, respectively. Open symbols correspond to indicator values for individual study streams and filled points represent the mean \pm one standard deviation within each human impact category. Longitudinal differences (downstream – upstream) in ecological indicators are summarized by boxplots (min, 25th percentile, 50th percentile, 75th percentile, max) for each human impact category. Results for other ecological indicators are depicted in the supplementary material (Figure S1).

Our study was limited to a small number of potential sampling locations within each human impact category in order to minimize variation in catchment physiography, land cover composition, and reach characteristics, in addition to other logistical constraints (e.g., site

access). Small sample sizes likely accentuated the variability observed in nutrient concentrations downstream of sewage treatment facilities leading to large variation in ecological indices among sampling locations. Increased variability in stream conditions within human impact categories likely resulted in the lack of significant BACI interaction effects among streams. Although the multiple BACI experimental design is an effective method to establish generalizable conclusions on the effects of an identifiable impact on stream ecosystem conditions (e.g., Keough & Quinn, 2000; Ried & Quinn, 2004; Roberts et al., 2007; Moreno et al., 2007), suboptimal study designs or assessments with high variability have the potential to result in the false conclusion of an effect (i.e., false positive/type I error) or the false conclusion that there is no effect (i.e., false negative/type II error) (Smokorowski & Randall, 2016). Because of the importance of impact assessments in guiding management interventions, both types of error can have severe implications for environmental management (Keough & Mapstone, 1997). For example, falsely concluding that there is no effect of an impact can result in the further degradation of ecological conditions. Post hoc improvements to study designs (i.e., amalgamation of related human impact levels) and increasing the alpha value (i.e., probability of type I error) in hypothesis tests can lower the probability of type II error and increase the statistical power to detect differences in ecological variables. Where study design is limited by sample size, post hoc leniency is common when extra caution regarding an environmental impact is warranted (Hanson, 2011; Murtaugh, 2014). Thus, it is important to critically interpret the results from BACI experiments, especially experiments with suboptimal study designs or high variability (Smokorowski & Randall, 2016).

Evaluating our three-level BACI assessment results at an alpha value of 0.1, we found that two ecological indices had significant BACI interaction terms: $SBMI$ (Sampling Location : Human Impact, $F_{(2,11)} = 3.27$, $p = 0.077$) and dissimilarity_D (Sampling Location : Human Impact,

$F_{(2,11)} = 3.34$, $p = 0.073$). No additional significant BACI interaction terms were identified in two-level models that combined passive and active sewage effluent impact categories (Table S1), but the probability of type I error decreased for both s_{BMI} (Sampling Location : Human Impact, $F_{(1,12)} = 3.90$, $p = 0.072$) and dissimilarity_D (Sampling Location : Human Impact, $F_{(1,12)} = 6.60$, $p = 0.025$). Richness of benthic macroinvertebrate assemblages (s_{BMI}) decreased downstream of point sources with streams that received actively treated sewage effluent having a significantly larger reduction in richness compared to streams exposed to agricultural activities alone (Cohen's d , 1.451 [0.129, 2.773]; Table S2). However, longitudinal differences in benthic macroinvertebrate richness in agricultural streams did not differ in comparison to streams that received passively treated sewage effluent. Likewise, Bray Curtis dissimilarity of diatom assemblages (dissimilarity_D) increased downstream of point sources with streams that received actively treated sewage effluent having assemblages that were significantly more dissimilar than streams exposed to agricultural activities alone (Cohen's d , 1.833 [0.432, 3.235]), but longitudinal differences in agricultural streams again did not differ from streams that received passively treated effluent. Although the response observed in these two ecological indicators corresponds to the predicted effect of increased nutrient enrichment, our findings do not provide strong support for a generalizable conclusion on the cumulative impact of sewage effluent on ecological communities of agricultural streams.

Past studies on the impact of sewage treatment facilities have commonly attributed variation in the response of stream communities to the concentration and bioavailability of effluent downstream of point sources (Welch et al., 1992; Chambers et al., 1997; Scrimgeour & Chambers, 2000; Bowman et al., 2005). For example, Quinn and Hickey (1993) found that the density of common benthic macroinvertebrate taxa sampled upstream and downstream of eight

sewage treatment lagoons differed from about 75 to 15% in association with a 6 to 484-fold difference in effluent dilution factor of downstream reaches. Although upstream sampling locations in these past studies often had lower nutrient concentrations than were measured in our study, the among stream variation that we observed in the longitudinal patterns of benthic diatom and macroinvertebrate assemblages may also be explained by the extent that downstream changes in agricultural activities and the discharge of sewage effluent increase stream nutrient concentrations. Therefore, a gradient study design targeting quantitative changes in stream nutrient enrichment may be better suited to evaluate the cumulative effects of sewage effluent on ecological communities of agricultural streams than a more generalized factorial approach.

3.2 Nutrient Enrichment Gradient

Multiple regression analyses between longitudinal differences (downstream – upstream) in absolute nutrient concentrations (normalized) and measured ecological indicators revealed two significant ($\alpha = 0.05$) models (Table S3; Figure 5; Figure S2). Increased inputs of TP were negatively associated with the diatom pollution tolerance index (PTI_D), indicative of a greater increase in abundance of pollution tolerant taxa ($p = 0.008$, adjusted $R^2 = 0.40$). Likewise, richness of benthic macroinvertebrate assemblages ($SBMI$) was observed to decrease downstream of cumulative TN inputs ($p = 0.027$, adjusted $R^2 = 0.35$). Increasing the leniency in model evaluation ($\alpha = 0.1$), two other ecological indicators were found to be associated with increased nutrient enrichment. In contrast to benthic macroinvertebrate assemblages, diatom assemblage richness (s_D) increased in association with TP enrichment ($p = 0.065$, adjusted $R^2 = 0.15$), but assemblages were observed to be more dissimilar ($Dissimilarity_D$) between sampling locations with greater differences in TP concentration ($p = 0.071$, adjusted $R^2 = 0.35$). Longitudinal differences in the Hilsenhoff family biotic index (FBI) also increased (indicative of increased

organic pollution) with TN enrichment ($p = 0.052$, adjusted $R^2 = 0.21$). Indeed, our finding of multiple significant associations between stream nutrient enrichment and longitudinal differences in ecological indicators provides greater support for the cumulative effects of nutrients on instream ecological communities compared to the factorial evaluation of human impact categories. However, the statistical support for multiple regression models between absolute nutrient concentrations and measured ecological indicators is likely not strong enough to provide defensible conclusions on the effects of nutrient enrichment due to the relaxed error assumptions in the majority of models that were considered significant.

A potential explanation for weak statistical models between stream specific differences in nutrient concentration and ecological indicators is the nonlinear response of instream ecological communities to environmental stressors. Many past studies have reported threshold or breakpoint type relationships between ecological communities and stream nutrient concentrations or associated land use gradients (e.g., Stevenson et al., 2008; Black et al., 2011; Evans-White et al., 2013; Heiskary & Bouchard, 2015; Grimstead et al., 2019). For example, Taylor et al., (2014) observed increased dissimilarity in diatom assemblage composition associated with increasing TP concentrations among oligotrophic to mesotrophic streams, but no difference in dissimilarity among mesotrophic to eutrophic streams. Longitudinal differences in absolute nutrient concentrations do not account for the background or ambient nutrient status of streams that receive nutrient enrichments. Thus, the expectation that larger nutrient inputs should be associated with greater differences in ecological conditions may be confounded by the pre-existing effect of upstream nutrient concentrations on stream communities (Lacoursière et al., 2011; Taylor et al., 2018). For example, Burdon et al. (2016) found that benthic macroinvertebrate communities in streams that drained catchments with intensive agricultural

456 land uses were more resistant to eutrophication-mediated disturbance from sewage effluent.
457 Therefore, the environmental context of upstream, pre-exposed reaches may need to be
458 quantitatively accounted for to fully understand the anthropogenic pressures associated with
459 nutrient enrichment on stream ecosystems.

460 Inverse (reciprocal) transformation of stream nutrient concentrations prior to upstream-
461 downstream differencing was used to increase the numerical weight of cumulative nutrient inputs
462 to streams with lower upstream concentrations compared to the same cumulative nutrient input
463 in streams with enriched upstream nutrient concentrations. Multiple regression analyses between
464 longitudinal differences in inverse transformed nutrient concentrations and measured ecological
465 indicators revealed seven significant ($\alpha = 0.05$) models with one additional ecological indicator
466 having a significant model at an alpha value of 0.1 (Table S3; Figure 5; Figure S2).

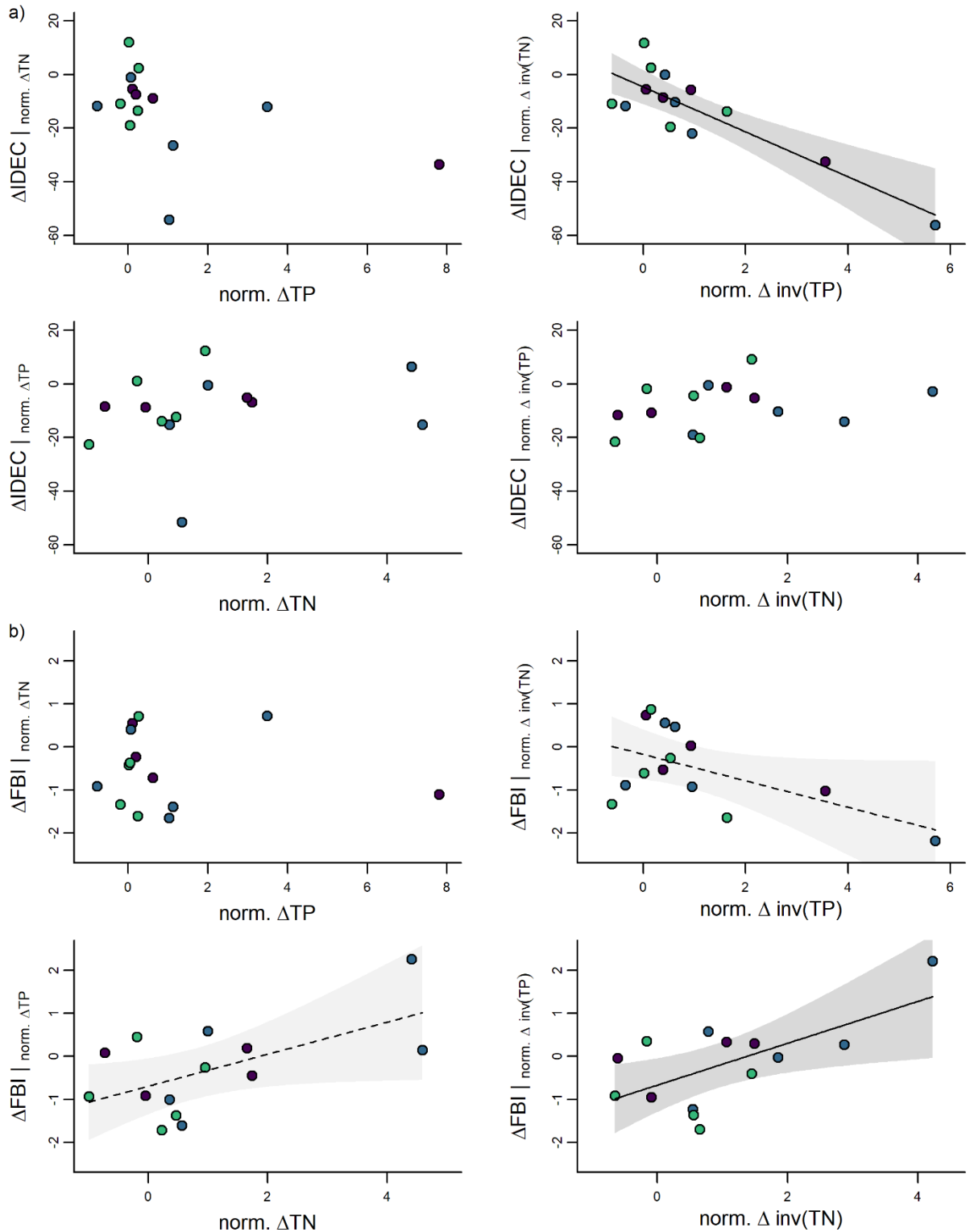


Figure 5: Multiple regression models of the association between normalized (norm.) longitudinal differences (downstream – upstream) in untransformed and inverse transformed (inv.) total phosphorus (TP) and total nitrogen (TN) concentrations and measures of (a) the eastern Canadian diatom index (IDEC) and (b) the Hilsenhoff family biotic index (FBI). Green, purple, and blue points correspond to streams exposed to agriculture, agriculture +

passively treated lagoon sewage effluent, and agriculture + actively treated mechanical sewage effluent, respectively. Solid trend lines correspond to significant associations at $\alpha = 0.05$ and dashed trend lines correspond to significant associations at $\alpha = 0.10$. Figures for other ecological indices are located in the supplementary material (Figure S2).

Three indicators of diatom assemblage composition had significant models: IDEC, %high-NP, and Dissimilarity_D. Longitudinal differences in the IDEC were negatively associated with greater enrichment of TP and indicative of increased pollution ($p < 0.001$, adjusted $R^2 = 0.68$). In contrast, longitudinal differences in the relative abundance of high nutrient taxa (%high-NP; $p = 0.052$, adjusted $R^2 = 0.32$) and assemblage dissimilarity (Dissimilarity_D; $p = 0.024$, adjusted $R^2 = 0.44$) were positively associated with cumulative inputs of TP.

Five indicators of benthic macroinvertebrate assemblage composition had significant models: FBI, PTI_{BMI}, %EPT, %Dipt., and s_{BMI} . Longitudinal differences in the Hilsenhoff family biotic index (FBI; $p = 0.021$, adjusted $R^2 = 0.36$), pollution tolerance index (PTI_{BMI}; $p = 0.037$, adjusted $R^2 = 0.24$), and Diptera relative abundance (%Dipt.; $p = 0.038$, adjusted $R^2 = 0.24$) were positively associated with TN enrichment and indicative of poorer ecological conditions. Similarly, TN enrichment was associated with poorer ecological conditions through negative associations with longitudinal differences in Ephemeroptera, Plecoptera, and Trichoptera relative abundance (%EPT; $p = 0.038$, adjusted $R^2 = 0.32$) and richness (s_{BMI} ; $p = 0.014$, adjusted $R^2 = 0.34$). However, in contrast to the associations observed with TN, TP enrichment was negatively associated with longitudinal differences in the Hilsenhoff family biotic index (FBI; $p = 0.056$, adjusted $R^2 = 0.36$) and positively associated with differences in the relative abundances of Ephemeroptera, Plecoptera, and Trichoptera taxa ($p = 0.052$, adjusted $R^2 = 0.32$). However, contrasting directions of association within these two models may be a result of leniency in error

assumptions as both associations with TP had increased probabilities of concluding a false positive (type I error).

Accounting for nonlinearities in associations with cumulative effects of nutrients resulted in an improvement in the number and strength of significant models between longitudinal differences in nutrient enrichment and ecological indicators (Table 3). In contrast to more general approaches used to evaluate our dataset, the inverse transformed nutrient gradient provided stronger evidence to support the negative effects of cumulative nutrient enrichment in streams. Our finding that ecological indicators measured in streams of lower nutrient status changed more substantially downstream of nutrient inputs compared to streams of higher nutrient status is consistent with the nonlinear response of stream communities to increased stress (Burdon et al., 2016; 2019). Therefore, the cumulative effects of sewage effluent and/or increased nutrient loading from agricultural activities appear to be context dependent based on ambient (upstream) stream conditions.

Table 3: Summary of significant models for the three statistical approaches used to evaluate the effect of nutrient enrichment and on indices of benthic diatom and macroinvertebrate assemblages. Positive (+) and negative (–) symbols indicate the direction of association for significant models with one and two symbols indicating significance at $\alpha = 0.10$ and $\alpha = 0.05$, respectively.

| Indicator | Multiple-BACI | Untransformed Enrichment Gradient | | Inv. Transformed Enrichment Gradient | |
|------------------------------|---------------|--------------------------------------|----|-----------------------------------------|----|
| | | TP | TN | TP | TN |
| <i>Diatoms</i> | | | | | |
| IDEC | | | | -- | |
| PTI _D | | -- | | | |
| %high-NP | | | | + | |
| %low-NP | | | | | |
| S _D | | + | | | |
| Dissimilarity _D | + | + | | ++ | |
| <i>Macroinvertebrates</i> | | | | | |
| FBI | | | + | - | ++ |
| PTI _{BMI} | | | | | ++ |
| %EPT | | | | + | -- |
| %Dipt. | | | | | ++ |
| SBMI | - | | -- | | -- |
| Dissimilarity _{BMI} | | | | | |

3.3 Summary and Conclusions

We demonstrated that cumulative nutrient inputs to streams can negatively influence biological communities, but that this conclusion would not have been realized with more general assessment approaches. Results from our multiple BACI assessment suggested that sewage effluent had no effect on measured ecological indicators and that upstream agricultural activities likely resulted in degradation to stream communities (e.g., loss of sensitive taxa). However, we expect that quantitative differences in environmental stressors associated with the operation of sewage treatment facilities resulted in our inability to qualitatively detect a generalizable, system-wide impact of sewage effluent. Moreover, although results from our gradient assessment that quantitatively evaluated the influence of longitudinal differences in nitrogen and phosphorus concentrations on stream communities adhered to the anticipated effects of nutrient enrichment, these findings also did not provide strong statistical support to make defensible conclusions

regarding ecological impacts of nutrient enrichment. Given the well-established nonlinear effect of nutrient enrichment on stream communities (e.g., Evans-White et al., 2013; Heiskary & Bouchard, 2015), we further refined our gradient assessment and transformed our nutrient enrichment gradient to increase the numerical weight of nutrient inputs that occurred in streams with low ambient nutrient concentrations. Accounting for the nonlinearity of nutrient enrichment effects reduced the unexplained variability observed in longitudinal differences in ecological indicators among streams and provided a more robust understanding of how stream communities respond to cumulative increases in nutrient concentrations. However, our inverse transformed nutrient enrichment gradient best represents ecological indicators that have a logarithmic-like stressor-response relationship. Different stressors or ecological indicators that have alternative nonlinear associations with increased nutrient concentrations (e.g., state shift) may therefore require independently refined statistical consideration in cumulative effects assessments.

Ecological monitoring and assessment studies are needed to evaluate human impacts on stream ecosystems and provide defensible information to guide mitigation actions. Even though considerable effort was taken during site selection to isolate the impact of sewage effluent, variability associated with complex, multi-use catchments in highly developed regions appears to have limited the ability of factorial assessment approaches to detect differences in diatom and benthic macroinvertebrate indices. Potential differences in other sewage effluent contaminants (e.g., micropollutants, synthetic chemicals, metals, and organic matter) and study design constraints (e.g., sample size and distance between sampling locations) may have further contributed to the residual variance in assessment models and resulted in disconcerted conclusions. However, despite potential covariates, we did observe that nutrient enrichment from sewage effluent was associated with the downstream degradation of biological community

structure independent of human activity categories in our gradient assessment. Simple BACI assessments conducted on ecological indicators replicated spatially or temporally within individual impact streams would therefore have likely yielded significant BACI interaction effects for some of the effluent receiving streams in our study. However, these simple assessments would be subject to limitations associated with pseudo-replication, restricted generalizability, and diagnostic interpretation. To provide basin-wide management recommendations on the effects of human impacts in highly developed regions, researchers may therefore need to go beyond standard assessment approaches and more explicitly consider the environmental stressors and their associated complexities related to the impact under investigation (e.g., Burdon et al., 2016).

From a management perspective, our finding that nutrient enrichment effects were only evident in our transformed nutrient enrichment gradient assessment indicates that the cumulative effects of nutrient enrichment are highly conditional on upstream ecosystem conditions. Thus, while communities of more degraded streams may be comprised of tolerant taxa and unaffected by further nutrient enrichment, communities of less degraded streams may change dramatically from increased nutrient inputs. Managers should therefore consider the trophic status of streams on a case-by-case basis to ensure that the most effective strategies are employed to conserve or improve stream ecosystem conditions. For example, nutrient load reductions from point sources (e.g., sewage treatment plants) would provide the most benefit to communities of streams with lower nutrient concentrations (oligotrophic/mesotrophic), but in streams with high nutrient concentrations (eutrophic) prioritizing the reduction of nutrient loading from nonpoint sources (e.g., agriculture) may be required before the benefits of cumulative load reductions can be observed in local stream communities.

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823 Supplementary Material

824 **Table S1:** Mixed model analysis of variance for modified before-after (Location) control-impact (Human Impact)
 825 assessment of nutrient concentrations, diatom assemblage indicators, and macroinvertebrate assemblage indicators.
 826 Analysis of variance results are reported for models with three (i.e., agriculture, passive sewage effluent, and active
 827 sewage effluent) and two (i.e., agriculture and sewage effluent) human impact factor levels. F and p values in bold
 828 represent significant models at $\alpha = 0.05$ (*) and $\alpha = 0.10$ (italic).

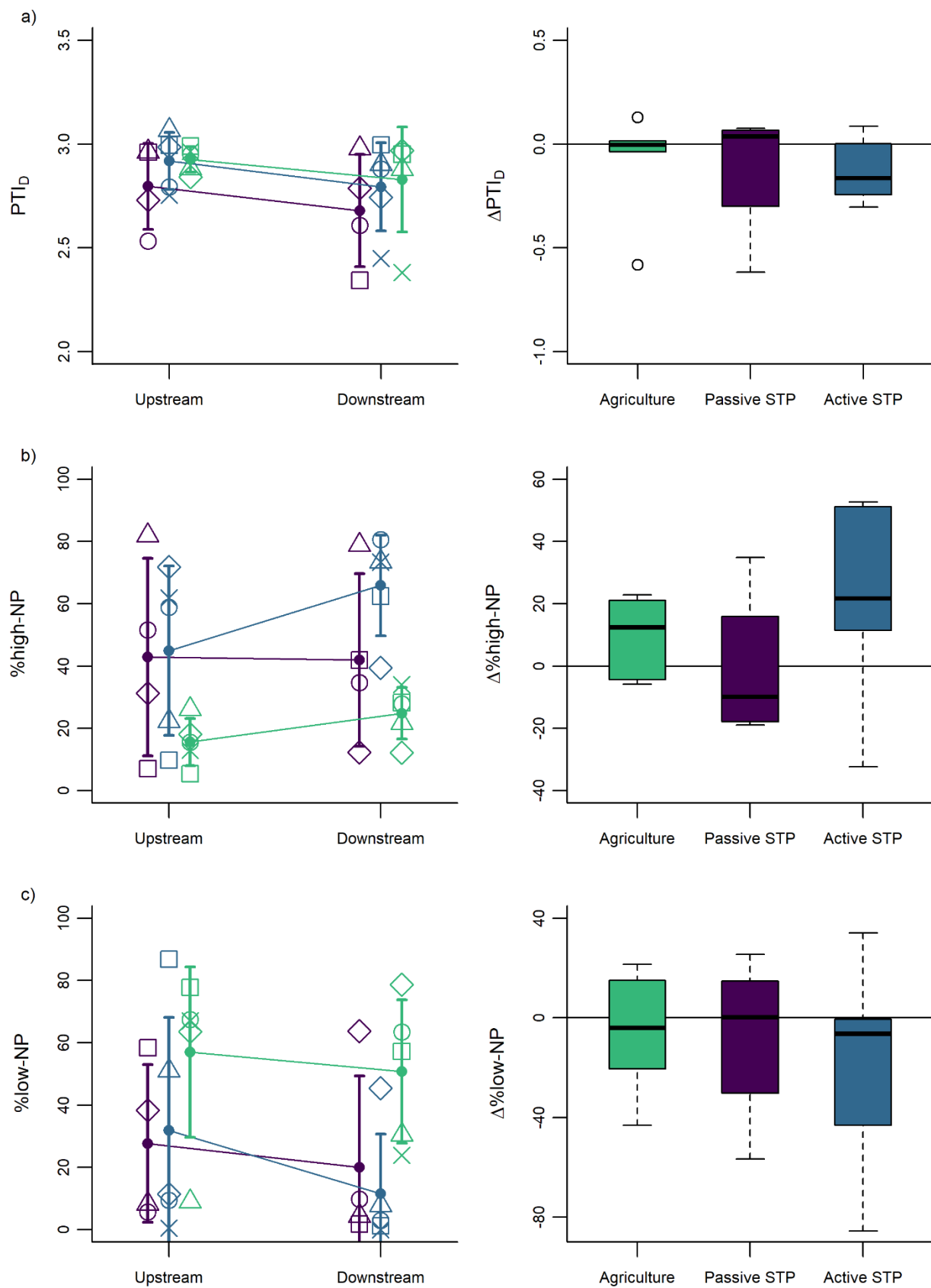
| Indicator | Predictor | Analysis of Variance | |
|-----------------------------------|-------------------------|---------------------------------------------------------|---------------------------------------------------------|
| | | 3 Human Impact Levels | 2 Human Impact Levels |
| <i>Nutrients</i> | | | |
| TP | Location | $F_{(1,11)} = 2.92, p = 0.116$ | $F_{(1,12)} = 3.00, p = 0.109$ |
| | Human Impact | $F_{(2,11)} = 1.12, p = 0.361$ | $F_{(1,12)} = 2.19, p = 0.165$ |
| | Location : Human Impact | $F_{(2,11)} = 1.03, p = 0.389$ | $F_{(1,12)} = 1.43, p = 0.255$ |
| TN | Location | $F_{(1,11)} = 6.22, p = 0.030^*$ | $F_{(1,12)} = 5.61, p = 0.036^*$ |
| | Human Impact | $F_{(2,11)} = 0.25, p = 0.785$ | $F_{(1,12)} = 0.40, p = 0.538$ |
| | Location : Human Impact | $F_{(2,11)} = 2.57, p = 0.121$ | $F_{(1,12)} = 2.56, p = 0.136$ |
| <i>Diatoms</i> | | | |
| IDEC | Location | $F_{(1,11)} = 7.95, p = 0.017^*$ | $F_{(1,12)} = 8.57, p = 0.013^*$ |
| | Human Impact | $F_{(2,11)} = 1.67, p = 0.233$ | <i>$F_{(1,12)} = 3.23, p = 0.098$</i> |
| | Location : Human Impact | $F_{(2,11)} = 0.56, p = 0.587$ | $F_{(1,12)} = 1.06, p = 0.323$ |
| PTI _D | Location | $F_{(1,11)} = 2.58, p = 0.136$ | $F_{(1,12)} = 2.82, p = 0.119$ |
| | Human Impact | $F_{(2,11)} = 1.07, p = 0.377$ | $F_{(1,12)} = 0.74, p = 0.405$ |
| | Location : Human Impact | $F_{(2,11)} = 0.02, p = 0.984$ | $F_{(1,12)} = 0.03, p = 0.860$ |
| %high-NP | Location | $F_{(1,11)} = 2.27, p = 0.160$ | $F_{(1,12)} = 2.17, p = 0.167$ |
| | Human Impact | $F_{(2,11)} = 5.71, p = 0.020^*$ | $F_{(1,12)} = 9.78, p = 0.009^*$ |
| | Location : Human Impact | $F_{(2,11)} = 0.80, p = 0.475$ | $F_{(1,12)} = 0.02, p = 0.896$ |
| %low-NP | Location | $F_{(1,11)} = 1.43, p = 0.258$ | $F_{(1,12)} = 1.52, p = 0.241$ |
| | Human Impact | <i>$F_{(2,11)} = 3.84, p = 0.054$</i> | $F_{(1,12)} = 8.34, p = 0.014^*$ |
| | Location : Human Impact | $F_{(2,11)} = 0.22, p = 0.808$ | $F_{(1,12)} = 0.18, p = 0.676$ |
| s _D | Location | $F_{(1,11)} = 0.00, p = 1.000$ | $F_{(1,12)} = 0.00, p = 1.000$ |
| | Human Impact | $F_{(2,11)} = 0.99, p = 0.402$ | $F_{(1,12)} = 1.90, p = 0.193$ |
| | Location : Human Impact | $F_{(2,11)} < 0.01, p = 0.991$ | $F_{(1,12)} = 0.01, p = 0.906$ |
| Dissimilarity _D | Location | — | — |
| | Human Impact | — | — |
| | Location : Human Impact | <i>$F_{(2,11)} = 3.34, p = 0.073$</i> | $F_{(1,12)} = 6.60, p = 0.025^*$ |
| <i>Benthic Macroinvertebrates</i> | | | |
| FBI | Location | $F_{(1,11)} = 1.43, p = 0.258$ | $F_{(1,12)} = 1.52, p = 0.242$ |
| | Human Impact | $F_{(2,11)} = 0.63, p = 0.552$ | $F_{(1,12)} = 0.02, p = 0.896$ |
| | Location : Human Impact | $F_{(2,11)} = 0.73, p = 0.502$ | $F_{(1,12)} = 1.26, p = 0.283$ |
| PTI _{BMI} | Location | $F_{(1,11)} = 0.08, p = 0.787$ | $F_{(1,12)} = 1.12, p = 0.313$ |
| | Human Impact | $F_{(2,11)} = 0.33, p = 0.726$ | $F_{(1,12)} = 0.09, p = 0.776$ |
| | Location : Human Impact | $F_{(2,11)} = 0.11, p = 0.899$ | $F_{(1,12)} = 1.42, p = 0.257$ |
| %EPT | Location | $F_{(1,11)} = 0.02, p = 0.905$ | $F_{(1,12)} = 0.08, p = 0.777$ |
| | Human Impact | $F_{(2,11)} = 0.63, p = 0.552$ | $F_{(1,12)} = 0.03, p = 0.867$ |
| | Location : Human Impact | $F_{(2,11)} = 0.24, p = 0.789$ | $F_{(1,12)} = 0.20, p = 0.661$ |
| %Dipt. | Location | $F_{(1,11)} < 0.01, p = 0.979$ | $F_{(1,12)} < 0.01, p = 0.978$ |
| | Human Impact | $F_{(2,11)} = 0.05, p = 0.955$ | $F_{(1,12)} = 0.06, p = 0.817$ |
| | Location : Human Impact | $F_{(2,11)} = 0.23, p = 0.796$ | $F_{(1,12)} = 0.19, p = 0.668$ |
| SBMI | Location | $F_{(1,11)} = 0.01, p = 0.908$ | $F_{(1,12)} = 0.01, p = 0.909$ |
| | Human Impact | $F_{(2,11)} = 0.43, p = 0.658$ | $F_{(1,12)} = 0.08, p = 0.388$ |
| | Location : Human Impact | <i>$F_{(2,11)} = 3.27, p = 0.077$</i> | <i>$F_{(1,12)} = 3.90, p = 0.072$</i> |
| Dissimilarity _{BMI} | Location | — | — |
| | Human Impact | — | — |
| | Location : Human Impact | $F_{(2,11)} = 0.06, p = 0.938$ | $F_{(1,12)} = 0.13, p = 0.726$ |

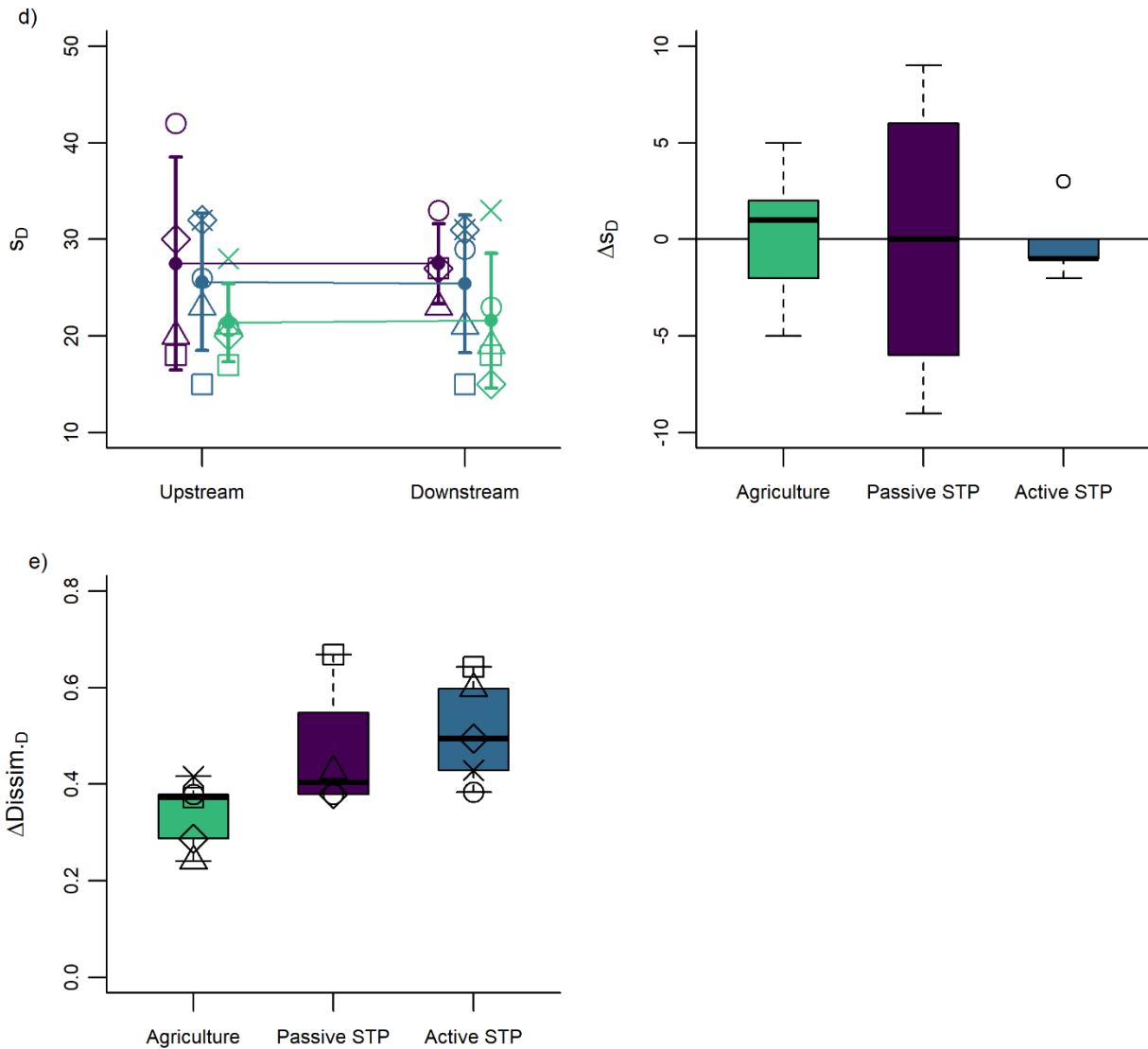
Table S2: Cohen's d effect size and 90% confidence interval (CI) for comparisons of stream specific differences (downstream – upstream) in each dependent variable between agricultural and sewage effluent receiving streams. Bolded values indicate effect sizes that have 90% confidence intervals that do not overlap zero.

| Indicator | Predictor | Cohen's d [90% CI] |
|-----------------------------------|-------------|-----------------------------|
| <i>Nutrients</i> | | |
| TP | All STP | 0.667 [-0.352, 1.686] |
| | Passive STP | 0.855 [-0.472, 2.182] |
| | Active STP | 0.796 [-0.426, 2.018] |
| TN | All STP | 0.892 [-0.147, 1.930] |
| | Passive STP | 0.572 [-0.725, 1.868] |
| | Active STP | 1.314 [0.017, 2.611] |
| <i>Diatoms</i> | | |
| IDEC | All STP | 0.574 [-0.438, 1.587] |
| | Passive STP | 0.533 [-0.760, 1.826] |
| | Active STP | 0.610 [-0.593, 1.813] |
| PTI _D | All STP | 0.100 [-1.116, 1.316] |
| | Passive STP | 0.069 [-1.202, 1.340] |
| | Active STP | 0.125 [-1.052, 1.302] |
| %high-NP | All STP | 0.074 [-1.141, 1.290] |
| | Passive STP | 0.529 [-0.764, 1.822] |
| | Active STP | 0.443 [-0.747, 1.634] |
| %low-NP | All STP | 0.239 [-0.980, 1.458] |
| | Passive STP | 0.048 [-1.222, 1.320] |
| | Active STP | 0.378 [-0.809, 1.564] |
| S _D | All STP | 0.068 [-1.148, 1.283] |
| | Passive STP | 0.034 [-1.237, 1.305] |
| | Active STP | 0.132 [-1.045, 1.309] |
| Dissimilarity _D | All STP | 1.433 [0.082, 2.784] |
| | Passive STP | 1.174 [-0.201, 2.549] |
| | Active STP | 1.833 [0.432, 3.235] |
| <i>Benthic Macroinvertebrates</i> | | |
| FBI | All STP | 0.627 [-0.615, 1.869] |
| | Passive STP | 0.589 [-0.709, 1.886] |
| | Active STP | 0.677 [-0.532, 1.886] |
| PTI _{BMI} | All STP | 0.664 [-0.581, -1.910] |
| | Passive STP | 0.703 [-0.606, 2.012] |
| | Active STP | 0.649 [-0.558, 1.855] |
| %EPT | All STP | 0.251 [-0.969, 1.470] |
| | Passive STP | 0.215 [-1.060, 1.489] |
| | Active STP | 0.261 [-0.920, 1.442] |
| %Dipt. | All STP | 0.245 [-0.974, 1.465] |
| | Passive STP | 0.600 [-0.698, 1.900] |
| | Active STP | 0.068 [-1.108, 1.245] |
| SBMI | All STP | 1.034 [-0.253, 2.322] |
| | Passive STP | 0.703 [-0.582, 2.043] |
| | Active STP | 1.451 [0.129, 2.773] |
| Dissimilarity _{BMI} | All STP | 0.200 [-1.418, 1.018] |
| | Passive STP | 0.217 [-1.058, 1.491] |
| | Active STP | 0.149 [-1.029, 1.327] |

834 **Table S3:** Independent cumulative effects multiple regression models between normalized stream specific differences (downstream – upstream) nutrient
835 concentrations (untransformed and inverse transformed) and measured ecological indicators of benthic diatom and macroinvertebrate assemblages. Bolded
836 coefficients (Coef.) and p values correspond to significant models at $\alpha = 0.05$ (*) and $\alpha = 0.10$ (italic). SE = standard error; Adj. = adjusted.

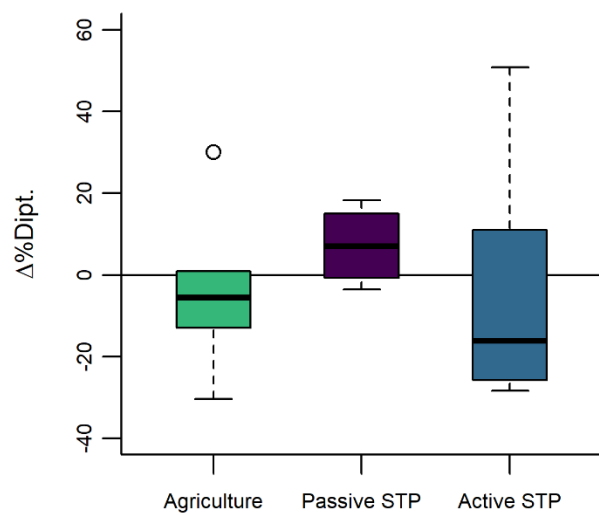
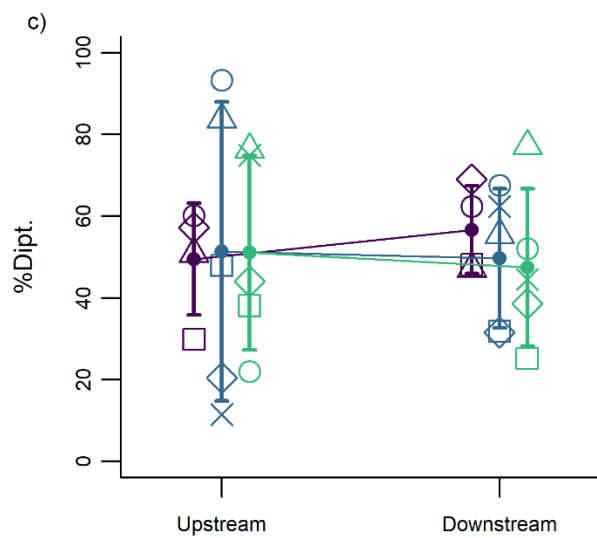
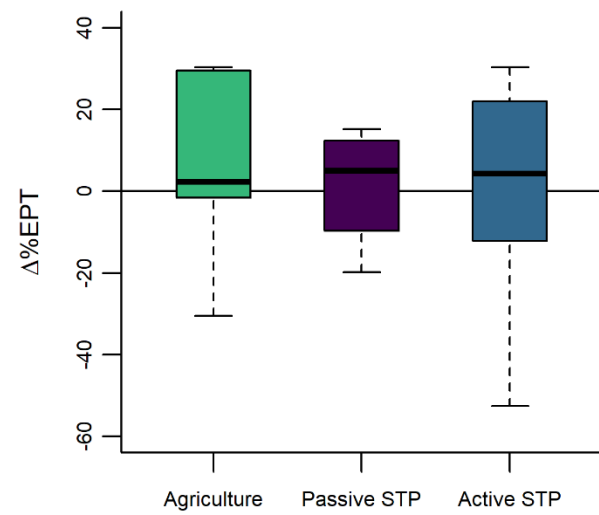
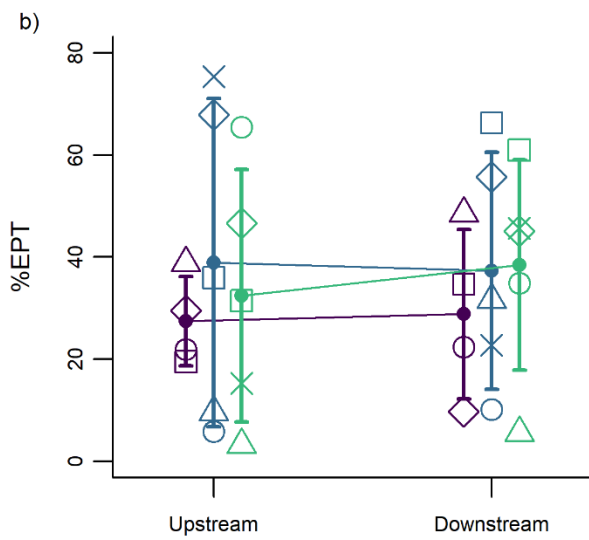
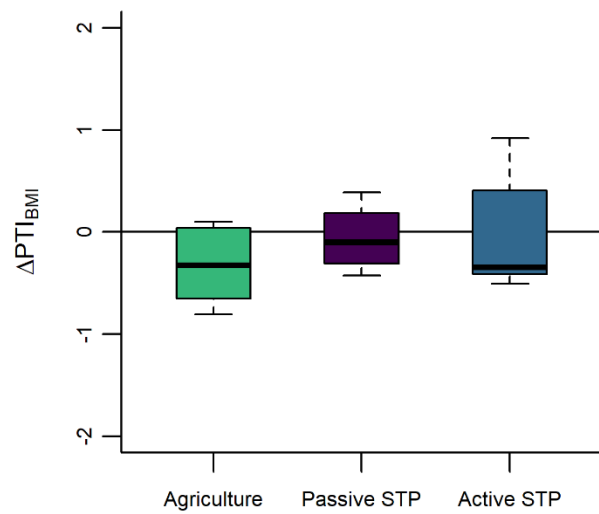
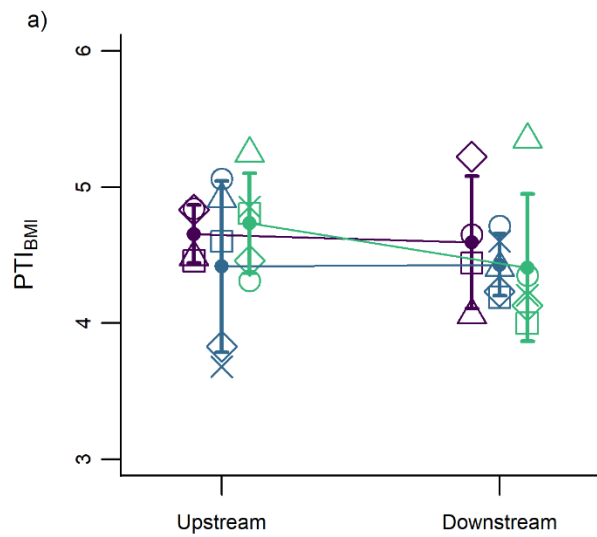
| | | Untransformed | | | | | Inverse Transformed | | | | |
|------------------------------|-----------|-------------------------------------|---------------|---------------------|--------------|---------------------|--------------------------------------|--------------------|---------------------|-------------------|---------------------|
| Indicator | Predictor | Coef. (\pm SE) | p-value | F _(2,11) | P | Adj. R ² | Coef. (\pm SE) | p-value | F _(2,11) | P | Adj. R ² |
| <i>Diatoms</i> | | | | | | | | | | | |
| IDEC | TP | -3.18 (\pm 2.24) | 0.183 | 1.01 | 0.394 | 0.002 | -8.36 (\pm1.57) | < 0.001* | 14.47 | < 0.001 | 0.675* |
| | TN | 2.06 (\pm 2.94) | 0.497 | | | | 1.81 (\pm 1.97) | 0.378 | | | |
| PTI _D | TP | -0.08 (\pm0.03) | 0.008* | 5.35 | 0.024 | 0.400* | -0.06 (\pm 0.04) | 0.167 | 1.23 | 0.330 | 0.034 |
| | TN | 0.04 (\pm 0.03) | 0.325 | | | | -0.00 (\pm 0.05) | 0.942 | | | |
| %high-NP | TP | 2.98 (\pm 3.56) | 0.421 | 1.08 | 0.373 | 0.012 | 7.86 (\pm3.61) | 0.052 | 4.12 | 0.046 | 0.324* |
| | TN | 3.43 (\pm 4.68) | 0.479 | | | | 5.30 (\pm 4.54) | 0.268 | | | |
| %low-NP | TP | -5.40 (\pm 4.66) | 0.271 | 1.42 | 0.282 | 0.061 | -9.46 (\pm 5.38) | 0.107 | 2.27 | 0.149 | 0.164 |
| | TN | -3.72 (\pm 6.11) | 0.555 | | | | -4.42 (\pm 6.77) | 0.527 | | | |
| SD | TP | 1.17 (\pm0.57) | 0.065 | 2.14 | 0.164 | 0.150 | 0.63 (\pm 0.80) | 0.449 | 0.34 | 0.719 | -0.113 |
| | TN | -0.47 (\pm 0.75) | 0.541 | | | | 0.01 (\pm 1.00) | 0.989 | | | |
| Dissimilarity _D | TP | 0.03 (\pm0.01) | 0.071 | 4.48 | 0.038 | 0.349* | 0.04 (\pm0.02) | 0.024 | 6.02 | 0.017 | 0.436* |
| | TN | 0.02 (\pm 0.02) | 0.275 | | | | 0.03 (\pm 0.02) | 0.181 | | | |
| <i>Macroinvertebrates</i> | | | | | | | | | | | |
| FBI | TP | -0.03 (\pm 0.13) | 0.845 | 2.70 | 0.111 | 0.207 | -0.31 (\pm0.14) | 0.056 | 4.65 | 0.034 | 0.360* |
| | TN | 0.37 (\pm0.17) | 0.052 | | | | 0.49 (\pm0.18) | 0.021* | | | |
| PTI _{BMI} | TP | 0.02 (\pm 0.06) | 0.754 | 1.88 | 0.198 | 0.119 | -0.10 (\pm 0.07) | 0.186 | 3.09 | 0.086 | 0.244 |
| | TN | 0.13 (\pm 0.08) | 0.141 | | | | 0.21 (\pm0.09) | 0.037* | | | |
| %EPT | TP | 1.42 (\pm 3.47) | 0.691 | 0.80 | 0.474 | -0.032 | 7.52 (\pm3.45) | 0.052 | 4.01 | 0.049 | 0.316* |
| | TN | -5.71 (\pm 4.10) | 0.235 | | | | -10.25 (\pm4.34) | 0.038* | | | |
| %Dipt. | TP | 2.58 (\pm 2.98) | 0.327 | 2.24 | 0.153 | 0.160 | -4.83 (\pm 3.45) | 0.189 | 3.06 | 0.088 | 0.241 |
| | TN | 5.44 (\pm 3.91) | 0.245 | | | | 10.23 (\pm2.36) | 0.038* | | | |
| S _{BMI} | TP | -0.32 (\pm 1.20) | 0.794 | 4.42 | 0.039 | 0.345* | 0.61 (\pm 1.48) | 0.687 | 4.34 | 0.041 | 0.339* |
| | TN | -4.04 (\pm1.58) | 0.027* | | | | -5.42 (\pm1.86) | 0.014* | | | |
| Dissimilarity _{BMI} | TP | -0.01 (\pm 0.01) | 0.399 | 0.04 | 0.679 | -0.102 | -0.01 (\pm 0.02) | 0.624 | 0.128 | 0.881 | -0.155 |
| | TN | < 0.01 (\pm 0.02) | 0.836 | | | | < 0.01 (\pm 0.02) | 0.859 | | | |

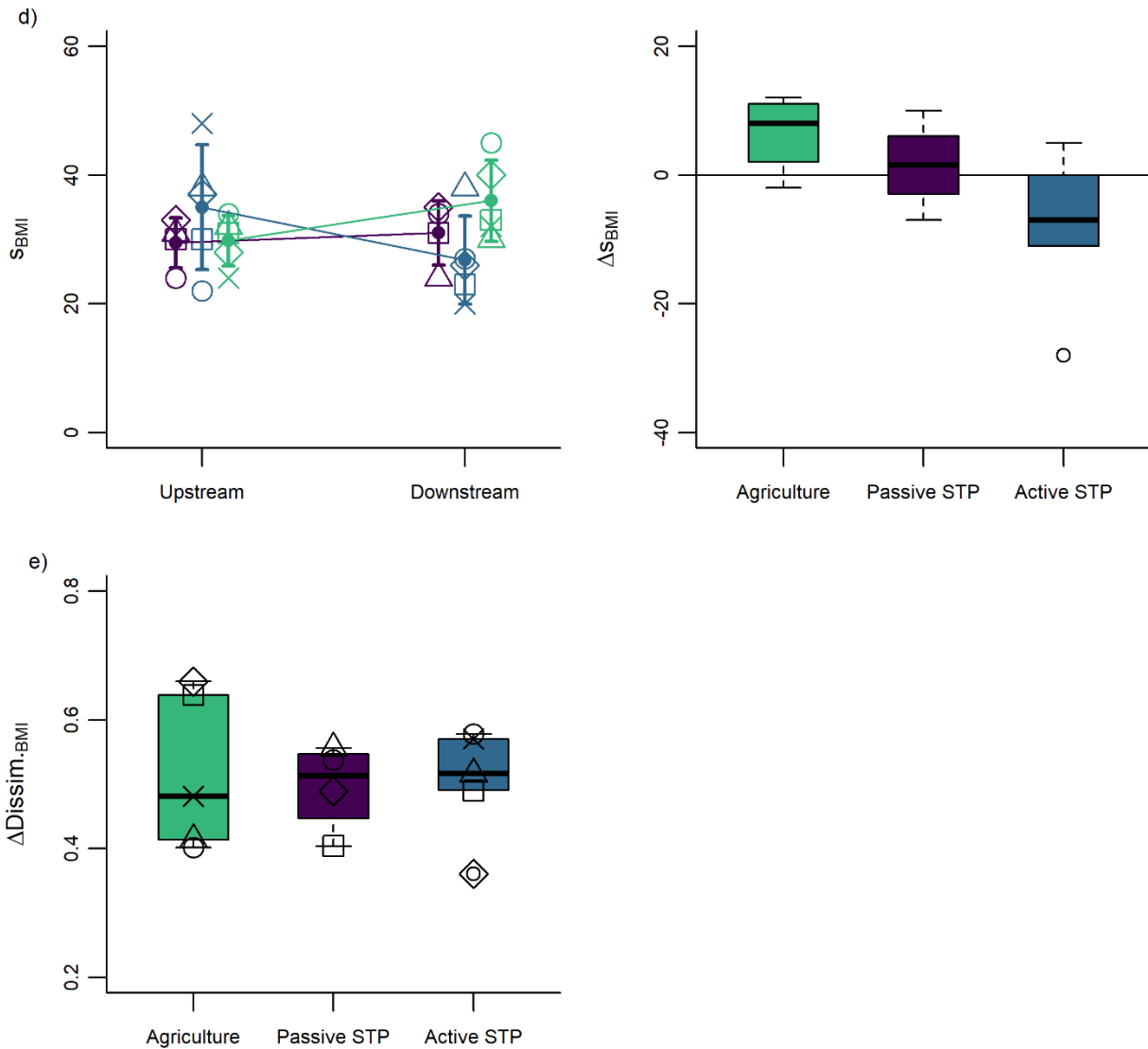




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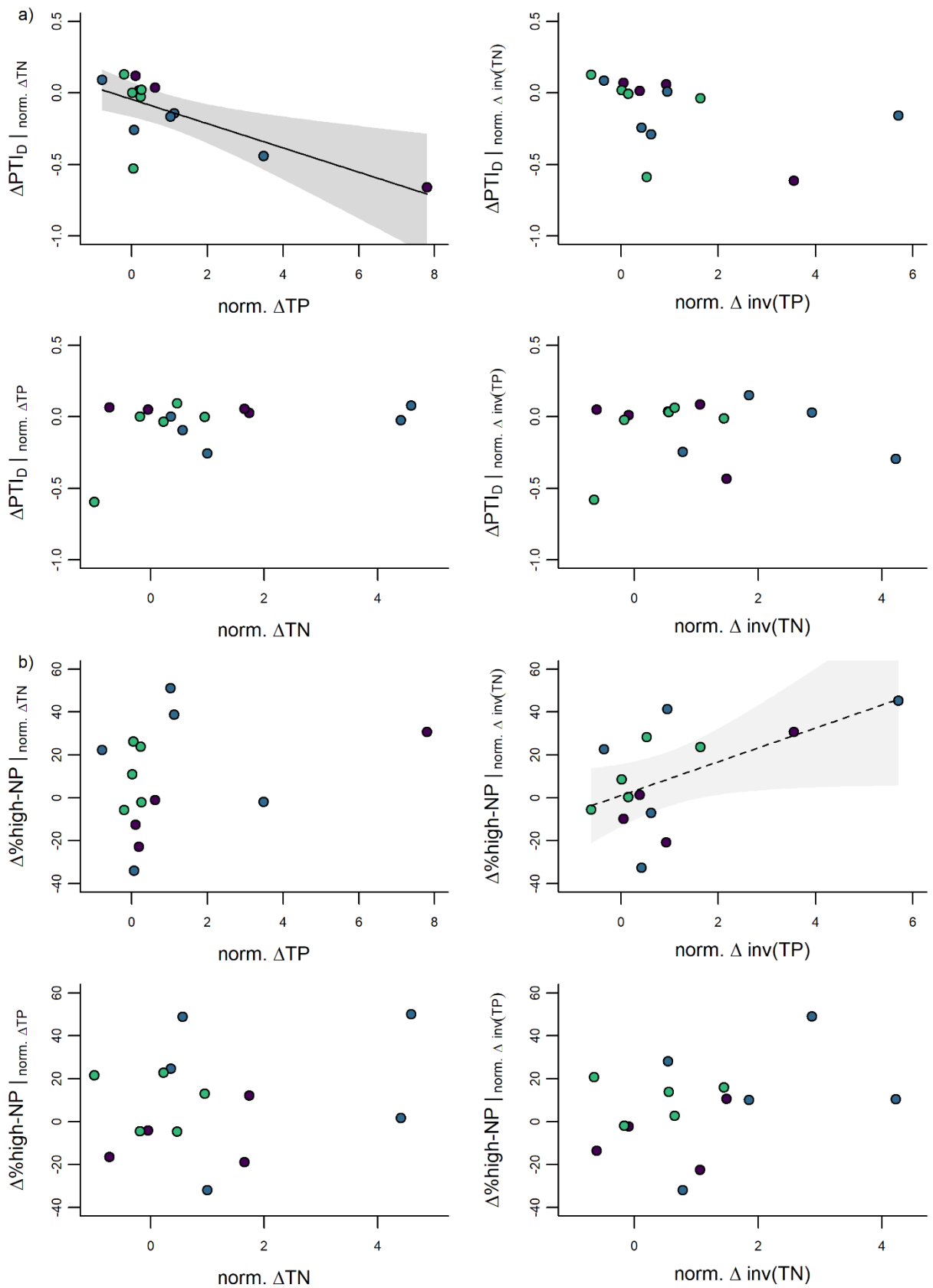
Figure S1-1: Mixed model analysis of variance for modified before-after (upstream-downstream) control-impact (agriculture-sewage effluent) assessment for ecological indicators of diatom assemblages: (a) pollution tolerance index (PTI_D), (b) percent abundance of high nutrient taxon (%high-NP), (c) percent abundance of low nutrient taxon (%low-NP), (d) taxon richness (s_D), and (e) Bray-Curtis dissimilarity (dissimilarity_D). Green, purple, and blue colours correspond to streams exposed to agriculture, agriculture + passively treated lagoon sewage effluent, and agriculture + actively treated mechanical sewage effluent, respectively. Open symbols correspond to indicator values for individual study streams and filled points represent the mean \pm one standard deviation within each human impact category. Longitudinal differences (downstream – upstream) in ecological indicators are summarized by boxplots (min, 25th percentile, 50th percentile, 75th percentile, max) for each human impact category.

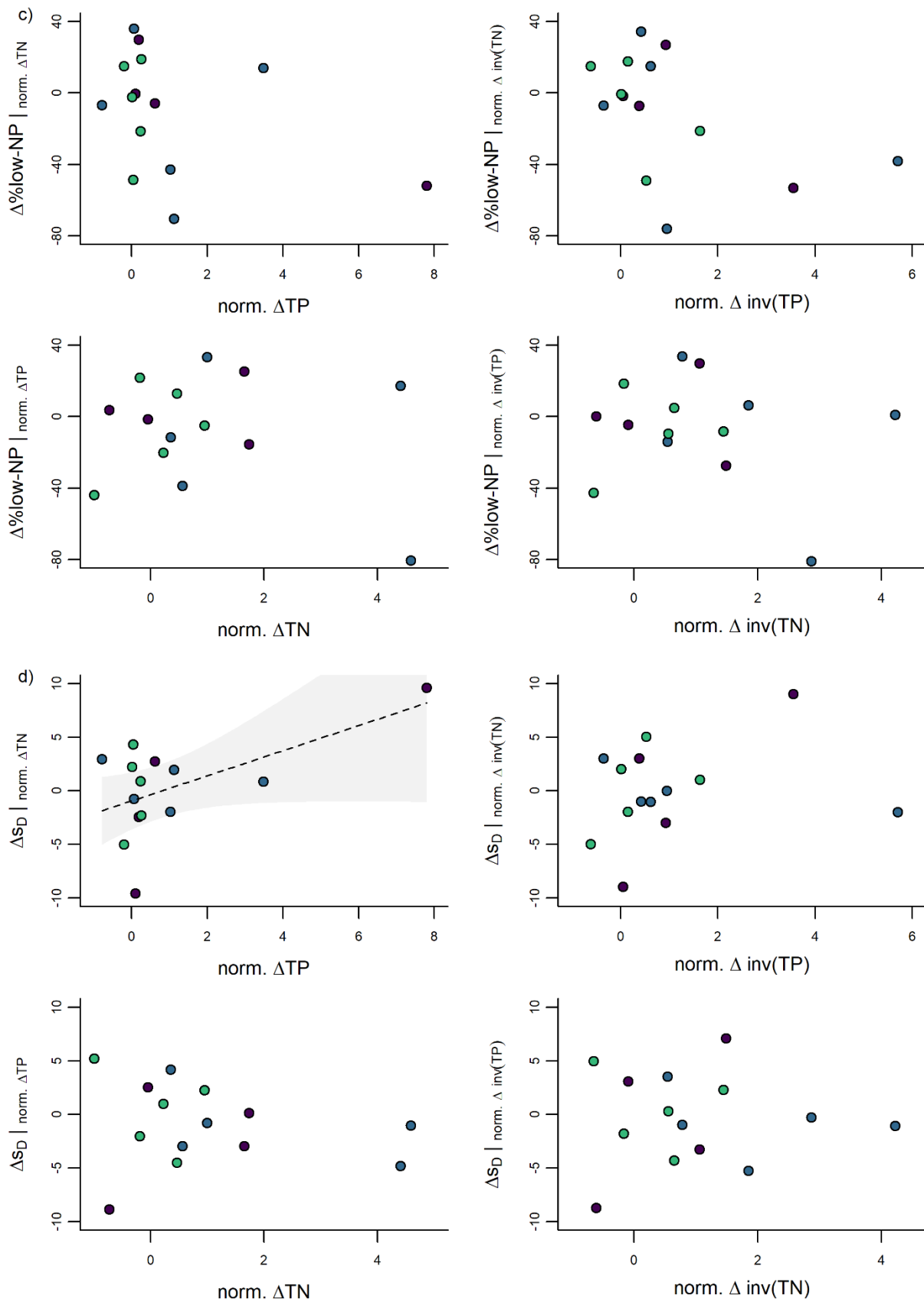




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Figure S1-2: Mixed model analysis of variance for modified before-after (upstream-downstream) control-impact (agriculture-sewage effluent) assessment for ecological indicators of benthic macroinvertebrate assemblages: (a) pollution tolerance index (PTI_{BMI}), (b) percent Ephemeroptera, Plecoptera, and Trichoptera abundance (%EPT), (c) percent Diptera abundance (%Dipt.), (d) taxon richness (s_{BMI}), and (e) Bray-Curtis dissimilarity ($dissimilarity_{BMI}$). Green, purple, and blue colours correspond to streams exposed to agriculture, agriculture + passively treated lagoon sewage effluent, and agriculture + actively treated mechanical sewage effluent, respectively. Open symbols correspond to indicator values for individual study streams and filled points represent the mean \pm one standard deviation within each human impact category. Longitudinal differences (downstream – upstream) in ecological indicators are summarized by boxplots (min, 25th percentile, 50th percentile, 75th percentile, max) for each human impact category.





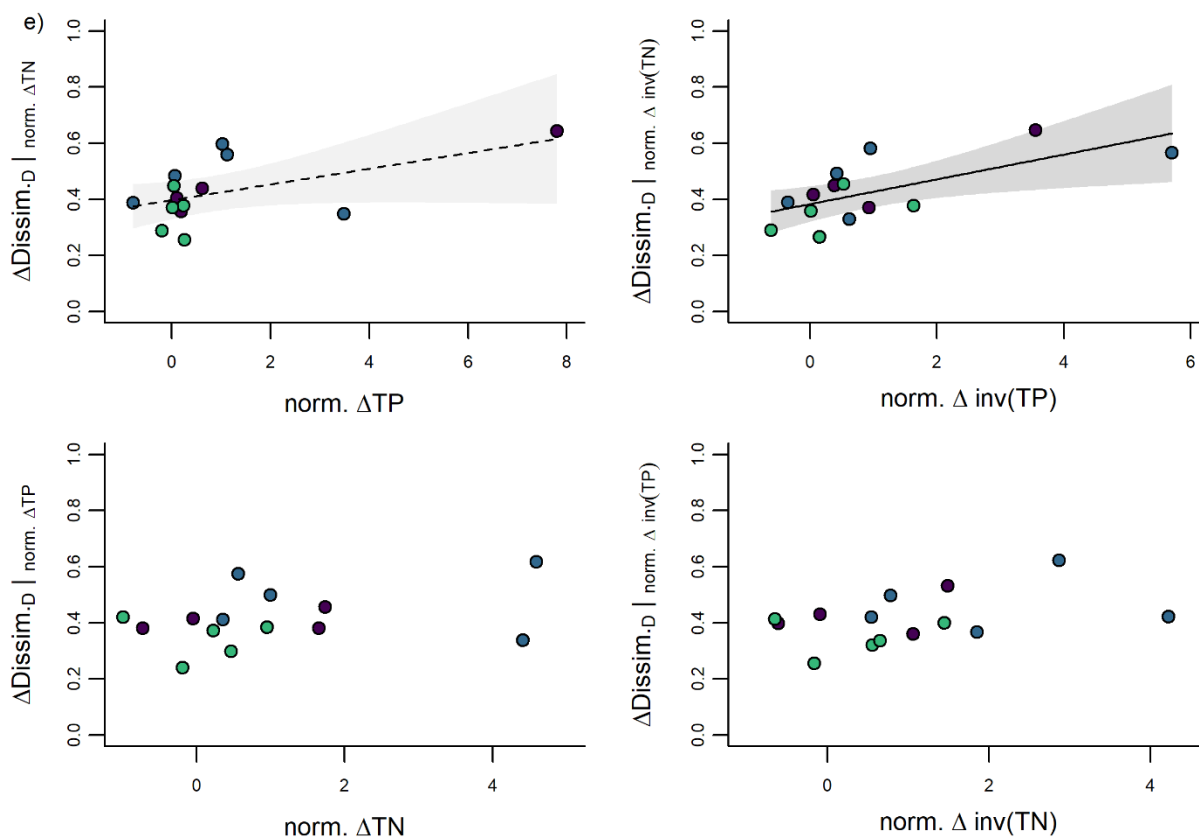
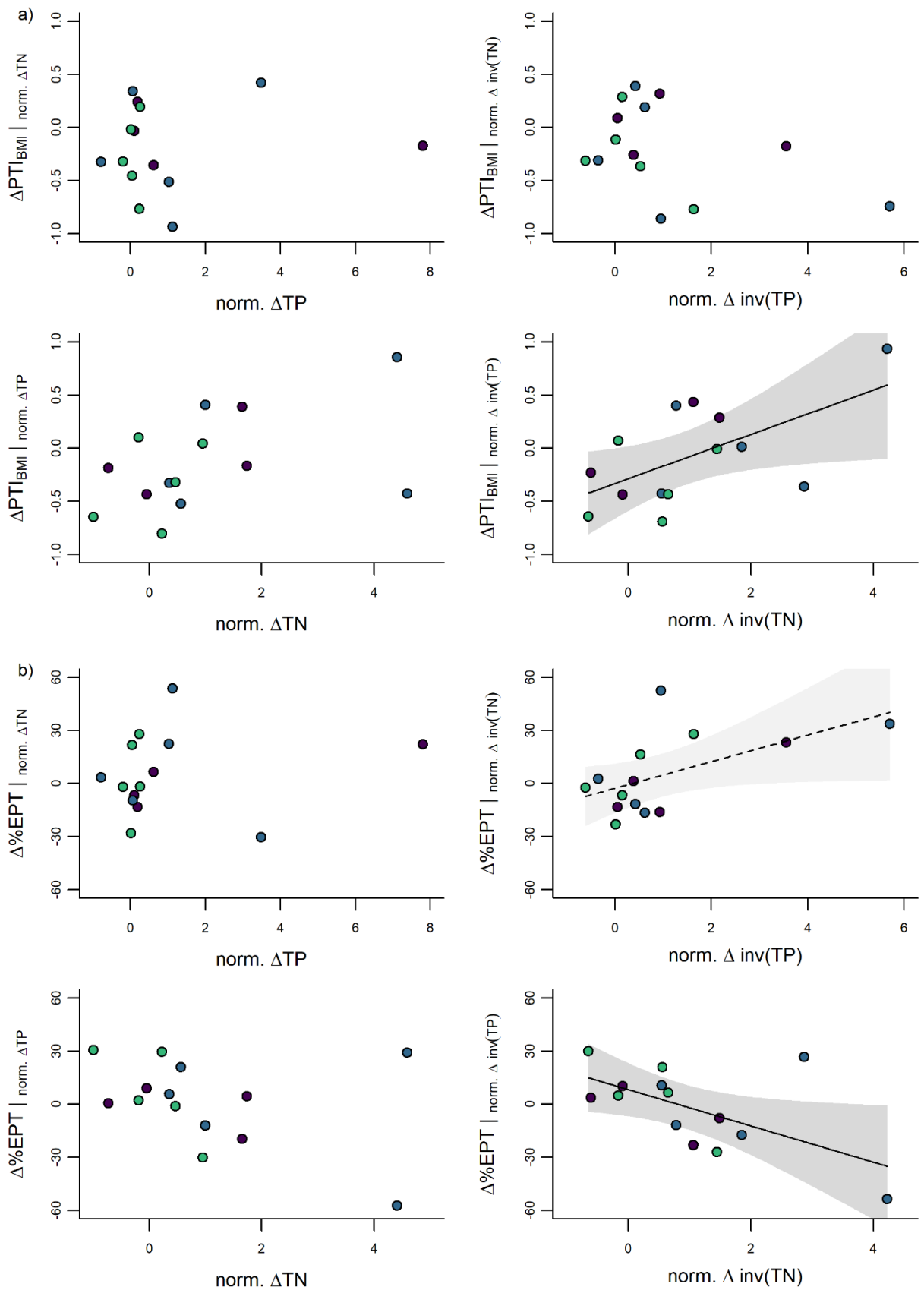
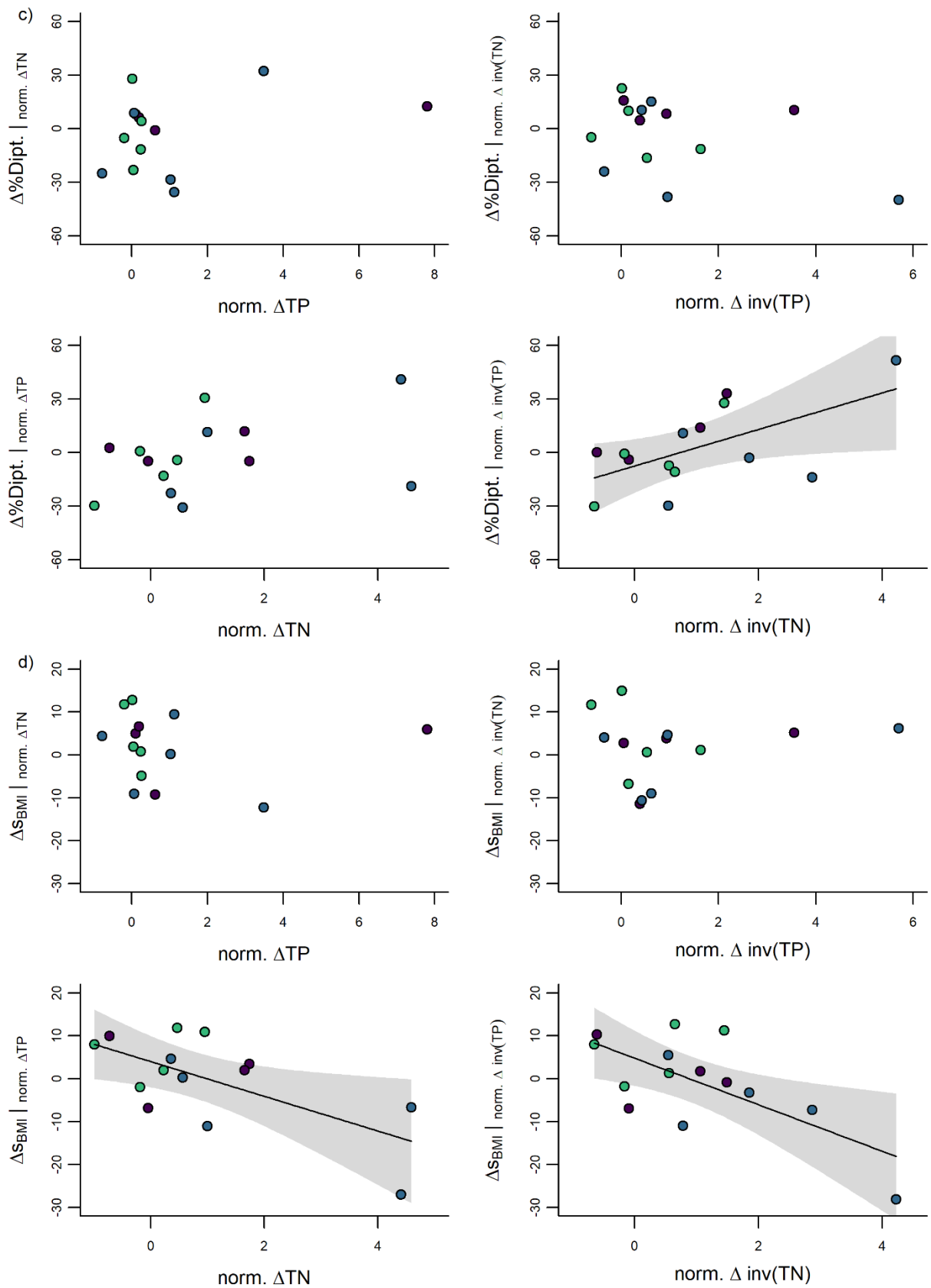


Figure S2-1: Multiple regression models of the association between normalized (norm.) longitudinal differences (downstream – upstream) in untransformed and inverse transformed (inv.) total phosphorus (TP) and total nitrogen (TN) concentrations and measures of ecological indicators of diatom assemblages: (a) pollution tolerance index (PTI_D), (b) percent abundance of high nutrient taxon (%high-NP), (c) percent abundance of low nutrient taxon (%low-NP), (d) taxon richness (s_D), and (e) Bray-Curtis dissimilarity (dissimilarity_D). Green, purple, and blue points correspond to streams exposed to agriculture, agriculture + passively treated lagoon sewage effluent, and agriculture + actively treated mechanical sewage effluent, respectively. Solid trend lines correspond to significant associations at $\alpha = 0.05$ and dashed trend lines correspond to significant associations at $\alpha = 0.10$.





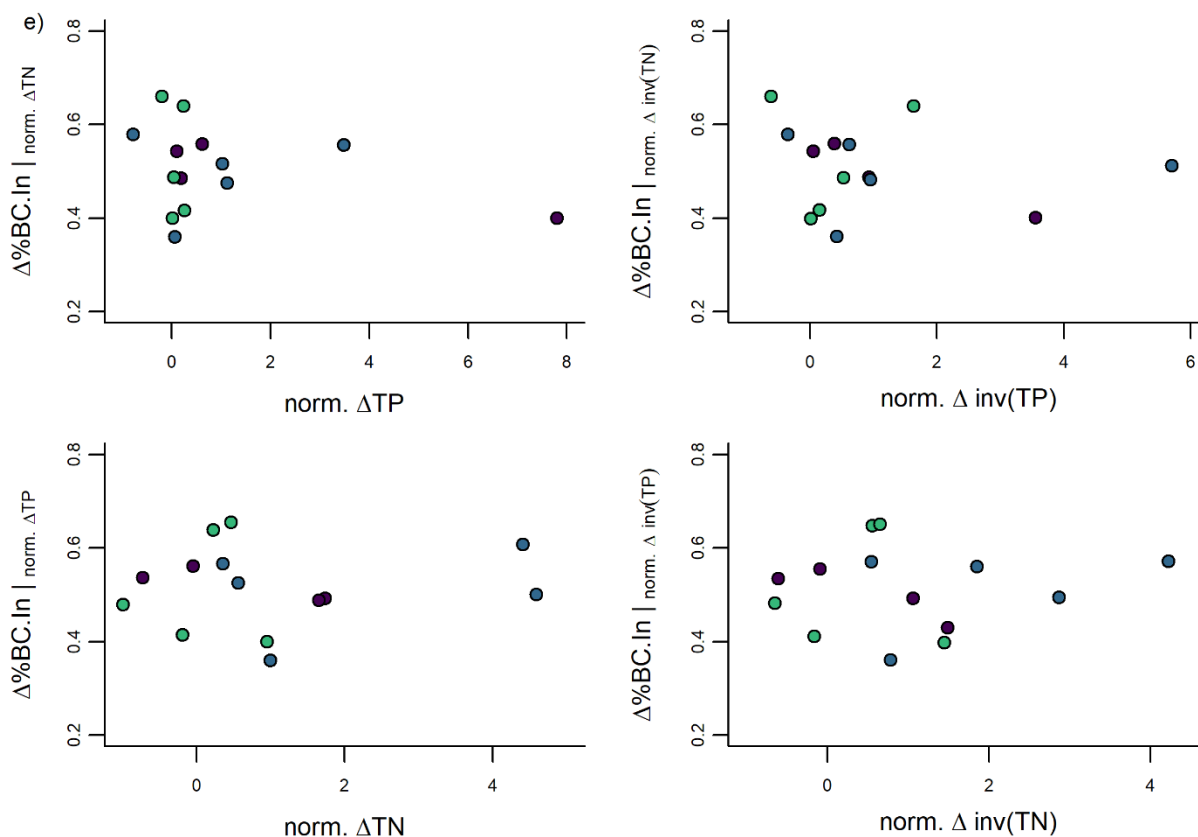


Figure S2-2: Multiple regression models of the association between normalized (norm.) longitudinal differences (downstream – upstream) in untransformed and inverse transformed (inv.) total phosphorus (TP) and total nitrogen (TN) concentrations and measures of ecological indicators of benthic macroinvertebrate assemblages: (a) pollution tolerance index (PTI_{BML}), (b) percent Ephemeroptera, Plecoptera, and Trichoptera abundance (%EPT), (c) percent Diptera abundance (%Dipt.), (d) taxon richness (s_{BML}), and (e) Bray-Curtis dissimilarity ($dissimilarity_{BML}$). Green, purple, and blue points correspond to streams exposed to agriculture, agriculture + passively treated lagoon sewage effluent, and agriculture + actively treated mechanical sewage effluent, respectively. Solid trend lines correspond to significant associations at $\alpha = 0.05$ and dashed trend lines correspond to significant associations at $\alpha = 0.10$.