

1 **Nutrient enrichment effects are conditional on upstream nutrient concentrations:**

2 **Implications for bioassessment in multi-use catchments**

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

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

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

34 **1. Introduction**

35 Streams support and regulate important ecosystem processes (e.g., nutrient cycling, water
36 purification, and waste assimilation) and provide provisional and cultural services (e.g., food,
37 water, and recreation) to sustain societal needs (Lowe & Likens, 2005). However, human
38 impacts on stream ecosystems are diverse (e.g., habitat alteration and pollution) and are expected
39 to increase with continued population growth and climate change (Søndergaard & Jeppesen,
40 2007). Thus, there is an ongoing need to assess the impacts of human activities to provide
41 managers and policy makers with scientifically defensible information on how to better protect
42 the structure and function of stream ecosystems.

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

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

63 Past studies have shown that although BACI assessments employ a factorial approach to
64 provide information on how the presence of an impact influences stream ecosystems,
65 quantitative differences in environmental stressors (e.g., chemical and sediment concentrations,
66 and thermal pollution) associated with an impact can govern the size of the perturbation from
67 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

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

106 Nutrient (i.e., nitrogen and phosphorus) enrichment is a pervasive chemical stressor that
107 affects many streams draining developed regions (Dodds & Smith, 2016; Wurtsbaugh et al.,
108 2019). Human impacts from point (e.g., sewage effluent) and nonpoint (e.g., agriculture) sources
109 of pollution can cumulatively increase instream nitrogen and phosphorus concentrations and
110 degrade the structure and function of stream ecosystems through accelerated eutrophication
111 (Mainstone & Parr, 2002; Withers & Jarvie, 2008). Eutrophication is characterized by excessive
112 primary production (i.e., sestonic and benthic algae, and macrophytes) and while enriched
113 nutrient concentrations can increase the biomass of primary producers, past studies have often

114 reported threshold or breakpoint type relationships between stream degradation by eutrophication
115 and nutrient concentrations (Evans-White et al., 2013; Jarvie et al., 2013; Heiskary & Bouchard,
116 2015). The ecological implications of stream nutrient enrichment have been well established
117 (e.g., Biggs, 2000; Friberg et al., 2010; Woodward et al., 2012) and because point sources can be
118 readily identified, there has been a substantive amount of research on the nutrient impacts of
119 treated sewage effluents (see Hamdhani et al., 2020). For example, many studies have reported
120 an increase in tolerant and decrease in sensitive benthic macroinvertebrate taxa downstream of
121 sewage outfalls compared to upstream reference reaches (e.g., Quinn & Hickey, 1993; Ortiz et
122 al., 2005; Englert et al., 2013). Assemblage-level differences have also been observed in benthic
123 algal communities (e.g., Chambers et al., 1997; Scrimgeour & Chambers, 2000; Bowman et al.,
124 2005). Most of these past studies occurred in streams with either low upstream nutrient
125 concentrations or a pronounced spike in concentration due to input of sewage with little or no
126 treatment. However, in developed regions sewage effluent is often discharged into streams
127 already enriched with nutrients from upstream nonpoint sources, such as agricultural lands,
128 resulting in more pronounced cumulative effects and nonlinear responses. The point source
129 discharge of effluent into agricultural streams thus provides an opportunity to determine if the
130 cumulative effects of nutrient enrichment can be detected with general and more nuanced
131 assessment approaches that account for nonlinearities associated with environmental stressors.

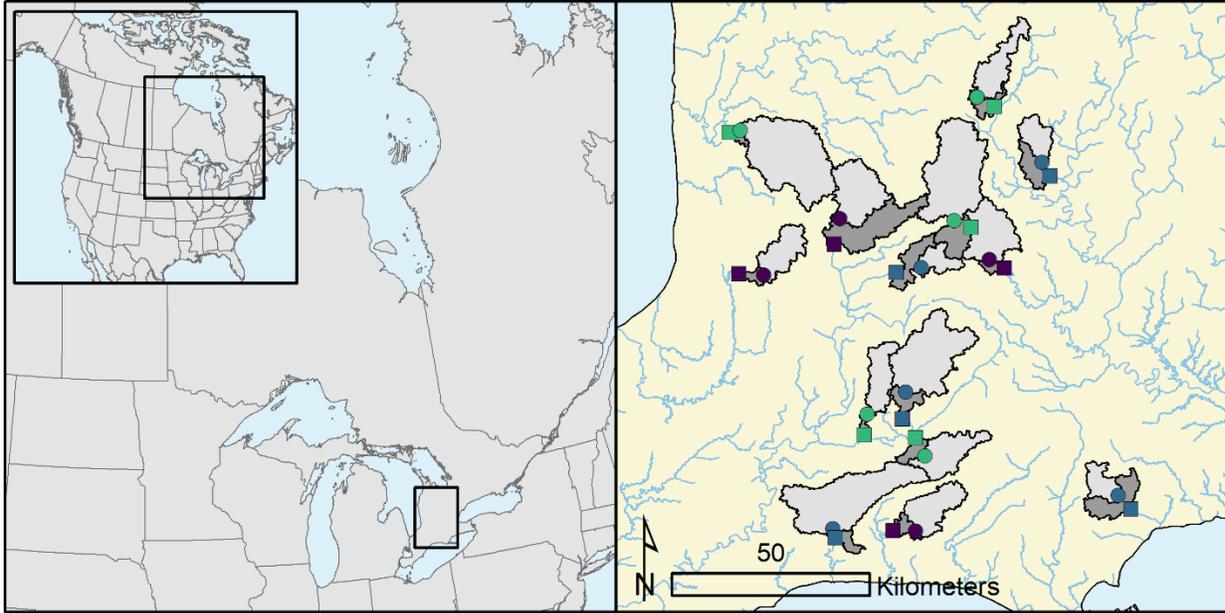
132 Our goal was to demonstrate the challenges of impact assessments in highly developed
133 regions by evaluating the cumulative effects of nutrient enrichment on instream ecological
134 communities using both a general and more targeted study design. First, a multiple before-after
135 (upstream-downstream)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

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

144 **2. Methods**

145 *2.1 Study Area and Design*

146 We assessed the cumulative impact of nutrients from agriculture and sewage effluent on
147 14 streams in the mixed-wood plains ecozone (Crins et al., 2009) of southern Ontario, Canada
148 (Figure 1). Southern Ontario is characterized by a humid, temperate climate with an average
149 annual temperature between 4.9 and 7.8 °C and total precipitation from 759 to 1087 mm (Crins et
150 al., 2009). The underlying physiography consists of post-glacial surface geologies dominated by
151 mixed agricultural land covers (i.e., crop and livestock operations) with patches of natural (i.e.,
152 forest and wetland) and urban areas throughout the landscape (Yates & Bailey, 2010b).



153

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

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

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

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

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

| Stream Name | Dist. | Upstream | | | | Downstream | | | |
|---------------------------|------------------|---------------------------|--------------|-------------|-------------|---------------------------|--------------|-------------|-------------|
| | | Area (km^2) | Agri. (%) | Urb. (%) | Nat. (%) | Area (km^2) | Agri. (%) | Urb. (%) | Nat. (%) |
| Agriculture | (km) | (km^2) | (%) | (%) | (%) | (km^2) | (%) | (%) | (%) |
| 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 |

182

183 *2.2 Sample Collection and Processing*

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

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

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

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

208 *2.3 Data Analysis*

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

222 *2.4 Diatom and Macroinvertebrate Indices*

223 The composition of benthic diatom and macroinvertebrate assemblages were described
224 using common bioassessment indices (see Barbour et al., 1999; Table 2). Diatom assemblages
225 were described by the Eastern Canadian Diatom Index (IDEC; Lavoie et al., 2014), pollution
226 tolerance index (PTI_D; Barbour et al., 1999; Muscio, 2002), percent abundance of high nutrient
227 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.

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

| Indicator | Description | Nutrient Enrichment Effect |
|------------------------------|---|----------------------------|
| <i>Diatoms</i> | | |
| IDECE | 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 | + |

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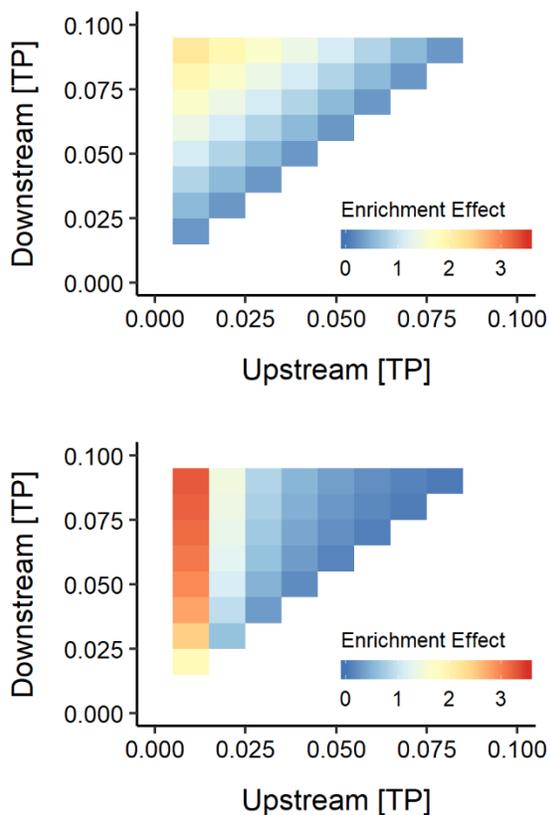
241 *2.5 Statistical Analyses*

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

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

266 Second, the potential cumulative effects of increased nitrogen and phosphorus
267 concentrations on the upstream-downstream differences in stream communities were evaluated
268 through a gradient assessment approach with data from streams exposed to agriculture only and
269 sewage effluent. Multiple regression analyses were used to determine the association ($\alpha = 0.05$)
270 between longitudinal differences (downstream – upstream) in TP and TN (independent variables)
271 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).



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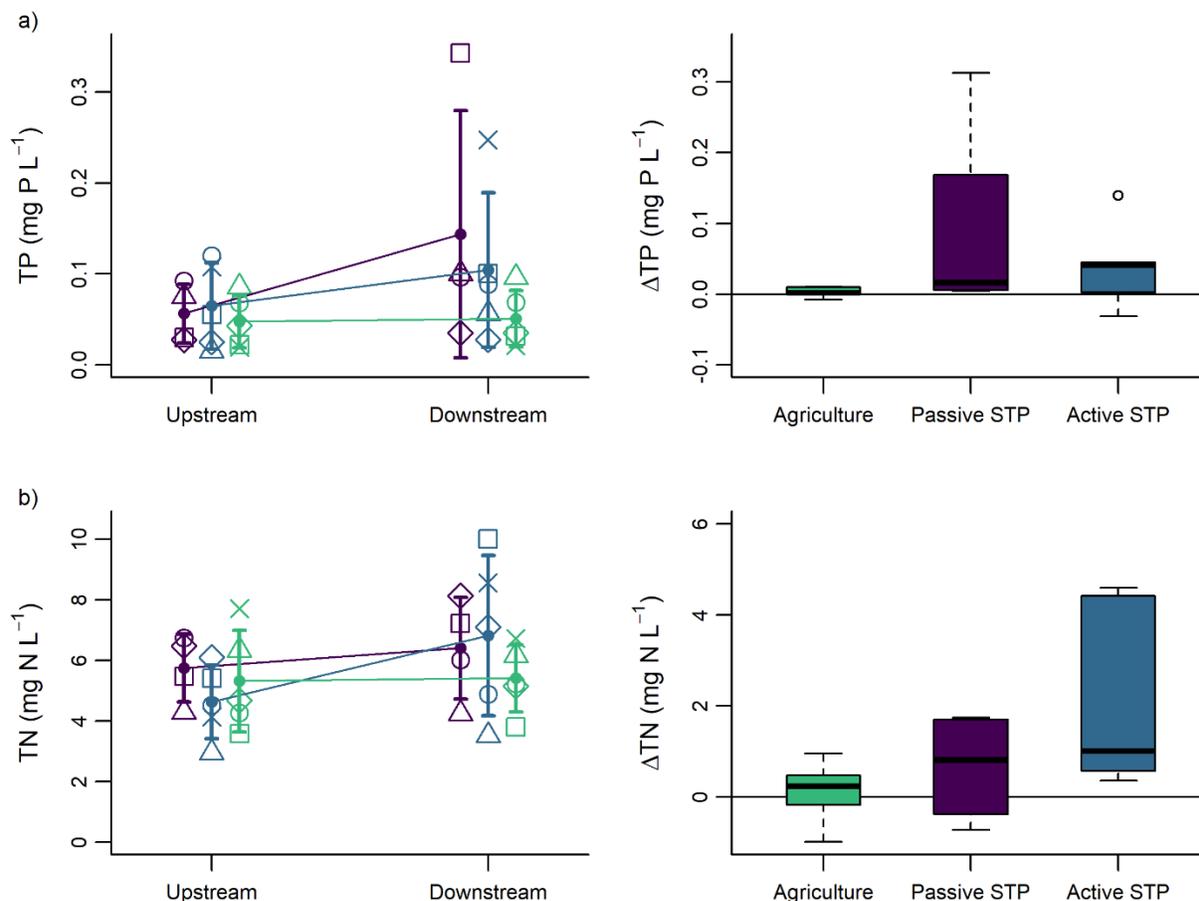
289 **Figure 2:** Illustration of the mean normalized enrichment effect of nutrients (colour gradient) based on (a) absolute
 290 concentration differencing and (b) inverse concentration differencing of hypothetical total phosphorus (TP)
 291 concentrations (mg L^{-1}) between upstream and downstream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

292 3. Results and Discussion

293 3.1 Multiple Before-After Control-Impact Assessment

294 Nutrient concentrations in agricultural streams and at upstream sampling locations in
 295 effluent impacted streams were on average (\pm standard deviation) $0.055 \pm 0.034 \text{ mg P L}^{-1}$ for TP
 296 and $5.25 \pm 1.27 \text{ mg N L}^{-1}$ for TN (Figure 3). Past studies in southern Ontario have reported
 297 similar total nutrient concentrations in streams that drain agricultural catchments (Raney &
 298 Eimers, 2014; Thomas et al., 2018; DeBues et al., 2019; Pearce et al., 2020) and compared to
 299 regional nutrient criteria ($0.024 \text{ mg P L}^{-1}$ for TP; 1.07 mg N L^{-1} for TN) most control streams in
 300 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.



311

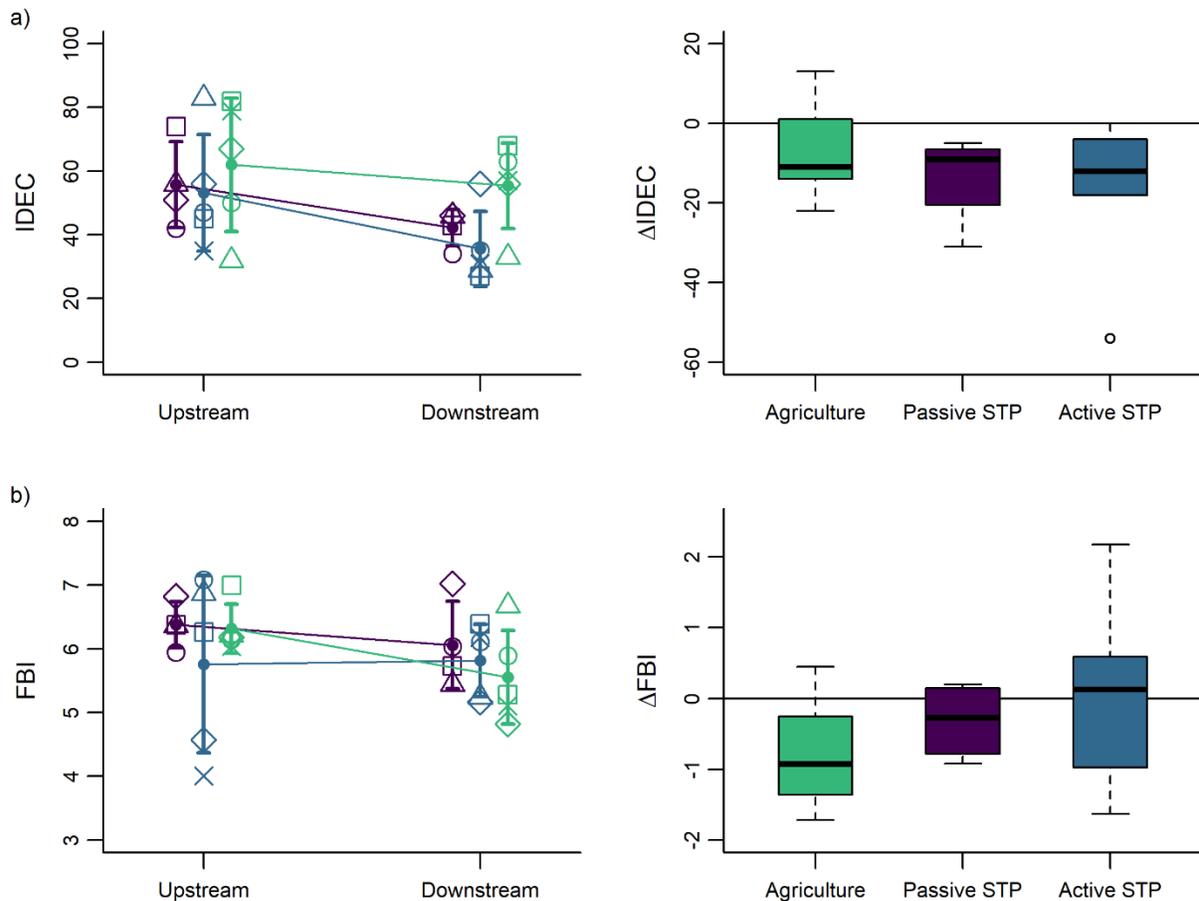
312 **Figure 3:** Mixed model analysis of variance for multiple before-after (upstream-downstream) control-impact
 313 (agriculture-sewage effluent) assessment of (a) total phosphorus (TP) and (b) total nitrogen (TN) concentrations.
 314 Green, purple, and blue colours correspond to streams exposed to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

320 Longitudinal differences (downstream – upstream) in nutrient concentrations were highly
 321 variable among streams that received sewage effluent (-0.03 to 0.31 mg P L⁻¹ for TP; -0.73 to
 322 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
 323 0.95 mg N L⁻¹ for TN) that received no point source inputs of effluent (Figure 3). Variable water
 324 quality impacts of sewage effluent have also been reported in past studies (Quinn & Hickey,
 325 1993; Chambers et al., 1997; Scrimgeour & Chambers, 2000; Bowman et al., 2005) and are often

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

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

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



351

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

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

364 access). Small sample sizes likely accentuated the variability observed in nutrient concentrations
365 downstream of sewage treatment facilities leading to large variation in ecological indices among
366 sampling locations. Increased variability in stream conditions within human impact categories
367 likely resulted in the lack of significant BACI interaction effects among streams. Although the
368 multiple BACI experimental design is an effective method to establish generalizable conclusions
369 on the effects of an identifiable impact on stream ecosystem conditions (e.g., Keough & Quinn,
370 2000; Ried & Quinn, 2004; Roberts et al., 2007; Moreno et al., 2007), suboptimal study designs
371 or assessments with high variability have the potential to result in the false conclusion of an
372 effect (i.e., false positive/type I error) or the false conclusion that there is no effect (i.e., false
373 negative/type II error) (Smokorowski & Randall, 2016). Because of the importance of impact
374 assessments in guiding management interventions, both types of error can have severe
375 implications for environmental management (Keough & Mapstone, 1997). For example, falsely
376 concluding that there is no effect of an impact can result in the further degradation of ecological
377 conditions. Post hoc improvements to study designs (i.e., amalgamation of related human impact
378 levels) and increasing the alpha value (i.e., probability of type I error) in hypothesis tests can
379 lower the probability of type II error and increase the statistical power to detect differences in
380 ecological variables. Where study design is limited by sample size, post hoc leniency is common
381 when extra caution regarding an environmental impact is warranted (Hanson, 2011; Murtaugh,
382 2014). Thus, it is important to critically interpret the results from BACI experiments, especially
383 experiments with suboptimal study designs or high variability (Smokorowski & Randall, 2016).

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

387 $F_{(2,11)} = 3.34, p = 0.073$). No additional significant BACI interaction terms were identified in
388 two-level models that combined passive and active sewage effluent impact categories (Table S1),
389 but the probability of type I error decreased for both s_{BMI} (Sampling Location : Human Impact,
390 $F_{(1,12)} = 3.90, p = 0.072$) and dissimilarity_D (Sampling Location : Human Impact, $F_{(1,12)} = 6.60, p$
391 $= 0.025$). Richness of benthic macroinvertebrate assemblages (s_{BMI}) decreased downstream of
392 point sources with streams that received actively treated sewage effluent having a significantly
393 larger reduction in richness compared to streams exposed to agricultural activities alone
394 (Cohen's $d, 1.451 [0.129, 2.773]$; Table S2). However, longitudinal differences in benthic
395 macroinvertebrate richness in agricultural streams did not differ in comparison to streams that
396 received passively treated sewage effluent. Likewise, Bray Curtis dissimilarity of diatom
397 assemblages (dissimilarity_D) increased downstream of point sources with streams that received
398 actively treated sewage effluent having assemblages that were significantly more dissimilar than
399 streams exposed to agricultural activities alone (Cohen's $d, 1.833 [0.432, 3.235]$), but
400 longitudinal differences in agricultural streams again did not differ from streams that received
401 passively treated effluent. Although the response observed in these two ecological indicators
402 corresponds to the predicted effect of increased nutrient enrichment, our findings do not provide
403 strong support for a generalizable conclusion on the cumulative impact of sewage effluent on
404 ecological communities of agricultural streams.

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

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

419 *3.2 Nutrient Enrichment Gradient*

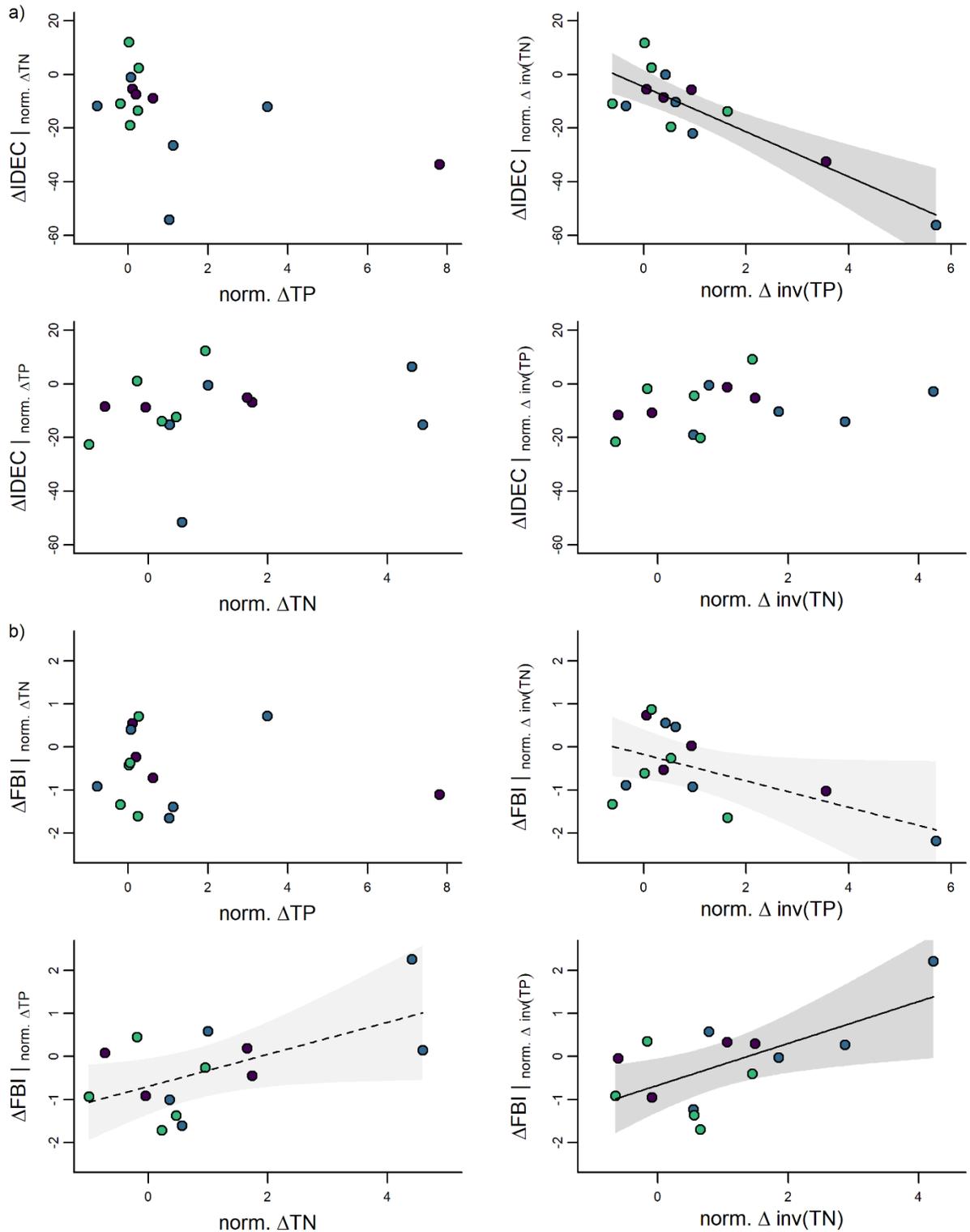
420 Multiple regression analyses between longitudinal differences (downstream – upstream)
421 in absolute nutrient concentrations (normalized) and measured ecological indicators revealed two
422 significant ($\alpha = 0.05$) models (Table S3; Figure 5; Figure S2). Increased inputs of TP were
423 negatively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

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

441 A potential explanation for weak statistical models between stream specific differences in
442 nutrient concentration and ecological indicators is the nonlinear response of instream ecological
443 communities to environmental stressors. Many past studies have reported threshold or breakpoint
444 type relationships between ecological communities and stream nutrient concentrations or
445 associated land use gradients (e.g., Stevenson et al., 2008; Black et al., 2011; Evans-White et al.,
446 2013; Heiskary & Bouchard, 2015; Grimstead et al., 2019). For example, Taylor et al., (2014)
447 observed increased dissimilarity in diatom assemblage composition associated with increasing
448 TP concentrations among oligotrophic to mesotrophic streams, but no difference in dissimilarity
449 among mesotrophic to eutrophic streams. Longitudinal differences in absolute nutrient
450 concentrations do not account for the background or ambient nutrient status of streams that
451 receive nutrient enrichments. Thus, the expectation that larger nutrient inputs should be
452 associated with greater differences in ecological conditions may be confounded by the pre-
453 existing effect of upstream nutrient concentrations on stream communities (Lacoursière et al.,
454 2011; Taylor et al., 2018). For example, Burdon et al. (2016) found that benthic
455 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).



467

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

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

476

477 Three indicators of diatom assemblage composition had significant models: IDEC,
478 %high-NP, and Dissimilarity_D. Longitudinal differences in the IDEC were negatively associated
479 with greater enrichment of TP and indicative of increased pollution ($p < 0.001$, adjusted $R^2 =$
480 0.68). In contrast, longitudinal differences in the relative abundance of high nutrient taxa
481 (%high-NP; $p = 0.052$, adjusted $R^2 = 0.32$) and assemblage dissimilarity (Dissimilarity_D; $p =$
482 0.024, adjusted $R^2 = 0.44$) were positively associated with cumulative inputs of TP.

483 Five indicators of benthic macroinvertebrate assemblage composition had significant
484 models: FBI, PTI_{BMI}, %EPT, %Dipt., and _{BMI}. Longitudinal differences in the Hilsenhoff family
485 biotic index (FBI; $p = 0.021$, adjusted $R^2 = 0.36$), pollution tolerance index (PTI_{BMI}; $p = 0.037$,
486 adjusted $R^2 = 0.24$), and Diptera relative abundance (%Dipt.; $p = 0.038$, adjusted $R^2 = 0.24$) were
487 positively associated with TN enrichment and indicative of poorer ecological conditions.
488 Similarly, TN enrichment was associated with poorer ecological conditions through negative
489 associations with longitudinal differences in Ephemeroptera, Plecoptera, and Trichoptera relative
490 abundance (%EPT; $p = 0.038$, adjusted $R^2 = 0.32$) and richness (_{BMI}; $p = 0.014$, adjusted $R^2 =$
491 0.34). However, in contrast to the associations observed with TN, TP enrichment was negatively
492 associated with longitudinal differences in the Hilsenhoff family biotic index (FBI; $p = 0.056$,
493 adjusted $R^2 = 0.36$) and positively associated with differences in the relative abundances of
494 Ephemeroptera, Plecoptera, and Trichoptera taxa ($p = 0.052$, adjusted $R^2 = 0.32$). However,
495 contrasting directions of association within these two models may be a result of leniency in error

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

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

509

510 **Table 3:** Summary of significant models for the three statistical approaches used to evaluate the effect of nutrient
 511 enrichment and on indices of benthic diatom and macroinvertebrate assemblages. Positive (+) and negative (-)
 512 symbols indicate the direction of association for significant models with one and two symbols indicating
 513 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. | | | | | ++ |
| S _{BMI} | - | | -- | | -- |
| Dissimilarity _{BMI} | | | | | |

514

515 3.3 Summary and Conclusions

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

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

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

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

560 From a management perspective, our finding that nutrient enrichment effects were only
561 evident in our transformed nutrient enrichment gradient assessment indicates that the cumulative
562 effects of nutrient enrichment are highly conditional on upstream ecosystem conditions. Thus,
563 while communities of more degraded streams may be comprised of tolerant taxa and unaffected
564 by further nutrient enrichment, communities of less degraded streams may change dramatically
565 from increased nutrient inputs. Managers should therefore consider the trophic status of streams
566 on a case-by-case basis to ensure that the most effective strategies are employed to conserve or
567 improve stream ecosystem conditions. For example, nutrient load reductions from point sources
568 (e.g., sewage treatment plants) would provide the most benefit to communities of streams with
569 lower nutrient concentrations (oligotrophic/mesotrophic), but in streams with high nutrient
570 concentrations (eutrophic) prioritizing the reduction of nutrient loading from nonpoint sources
571 (e.g., agriculture) may be required before the benefits of cumulative load reductions can be
572 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 $\alpha = 0.10$ (italicic).

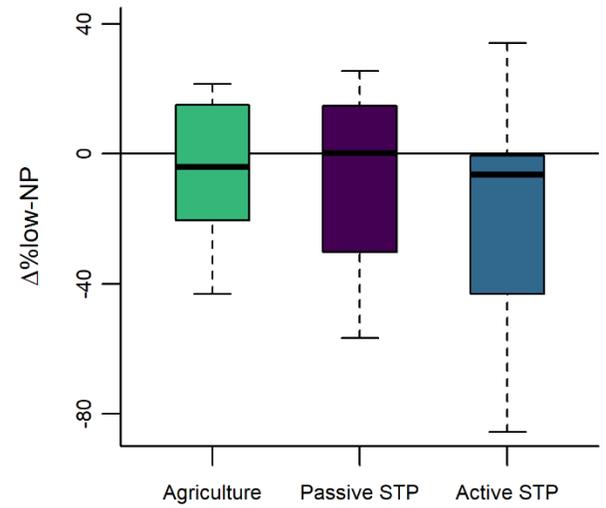
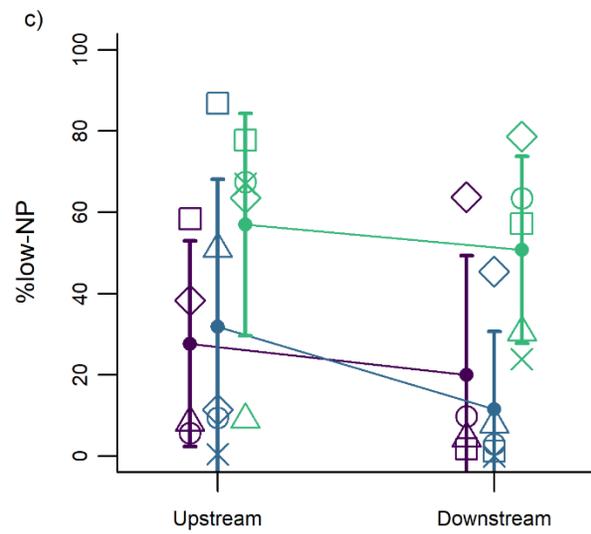
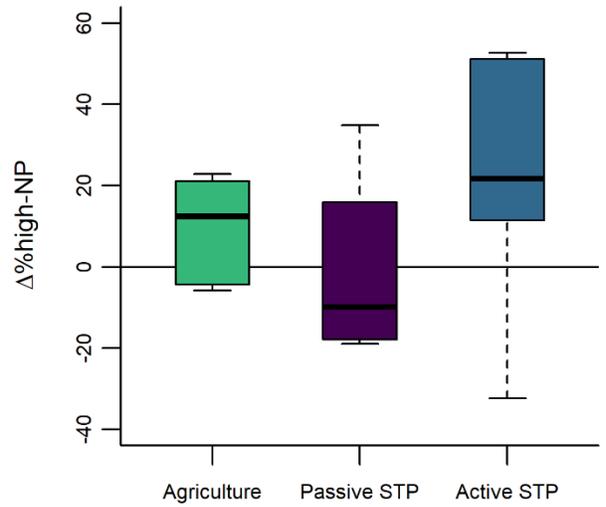
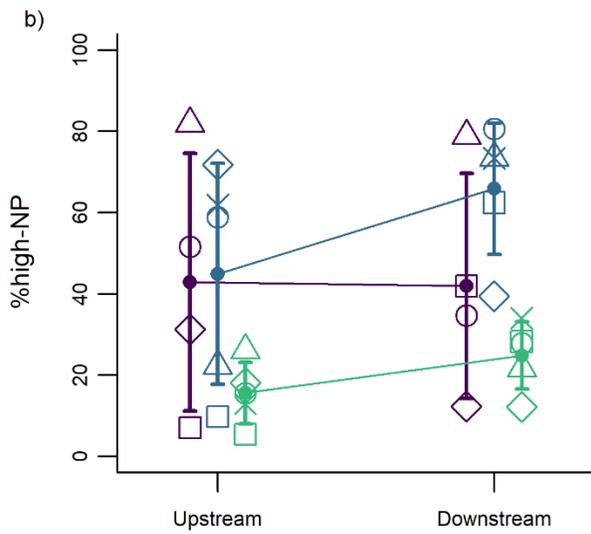
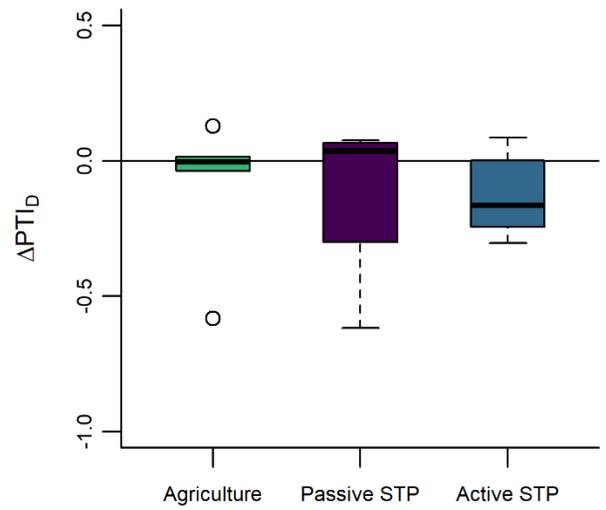
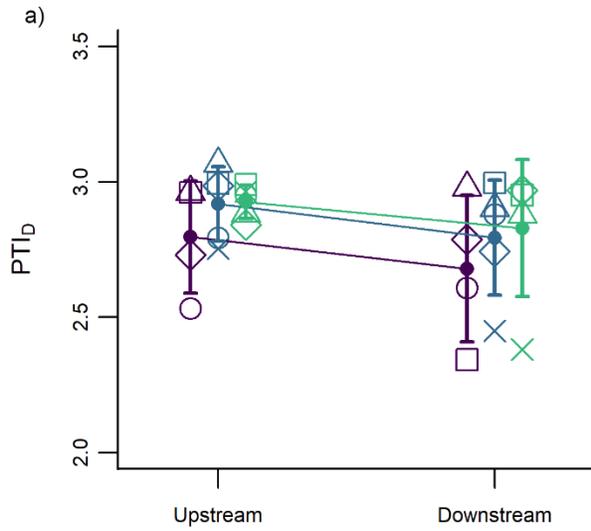
| Indicator | Predictor | Analysis of Variance | |
|-----------------------------------|-------------------------|--|--|
| | | 3 Human Impact Levels | 2 Human Impact Levels |
| <i>Nutrients</i> | | | |
| TP | Location | $F_{(1,11)} = 2.92, p = 0.116$ | $F_{(1,12)} = 3.00, p = 0.109$ |
| | Human Impact | $F_{(2,11)} = 1.12, p = 0.361$ | $F_{(1,12)} = 2.19, p = 0.165$ |
| | 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 | $F_{(2,11)} = 3.84, p = 0.054$ | $F_{(1,12)} = 8.34, p = 0.014^*$ |
| | 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 | $F_{(2,11)} = 3.34, p = 0.073$ | $F_{(1,12)} = 6.60, p = 0.025^*$ |
| <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 | $F_{(2,11)} = 0.23, p = 0.796$ | $F_{(1,12)} = 0.19, p = 0.668$ |
| s _{BMI} | 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 | $F_{(2,11)} = 3.27, p = 0.077$ | $F_{(1,12)} = 3.90, p = 0.072$ |
| Dissimilarity _{BMI} | Location | – | – |
| | Human Impact | – | – |
| | Location : Human Impact | $F_{(2,11)} = 0.06, p = 0.938$ | $F_{(1,12)} = 0.13, p = 0.726$ |

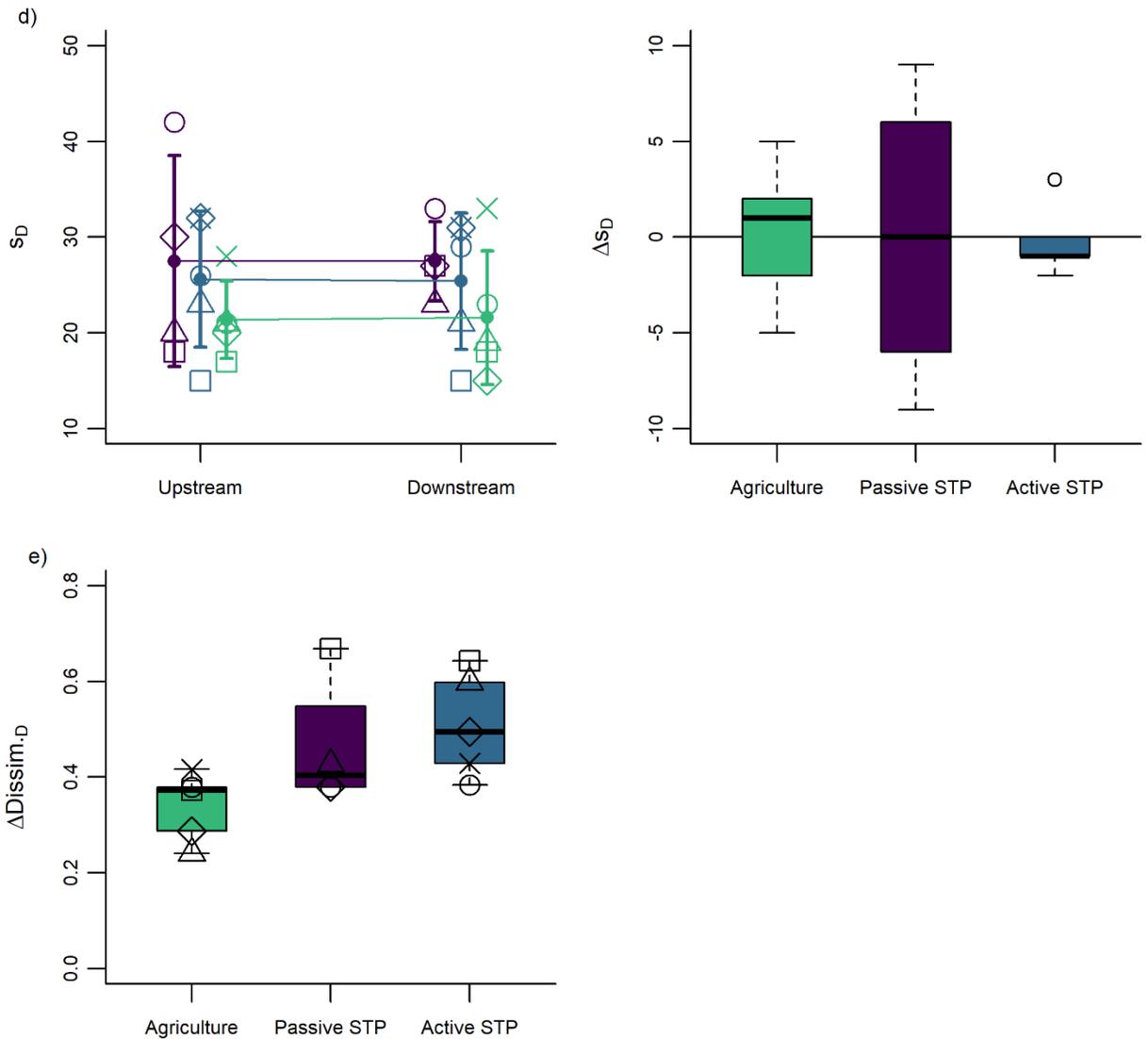
829 **Table S2:** Cohen's d effect size and 90% confidence interval (CI) for comparisons of stream
 830 specific differences (downstream – upstream) in each dependent variable between agricultural
 831 and sewage effluent receiving streams. Bolded values indicate effect sizes that have 90%
 832 confidence 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] |
| S _{BMI} | 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.

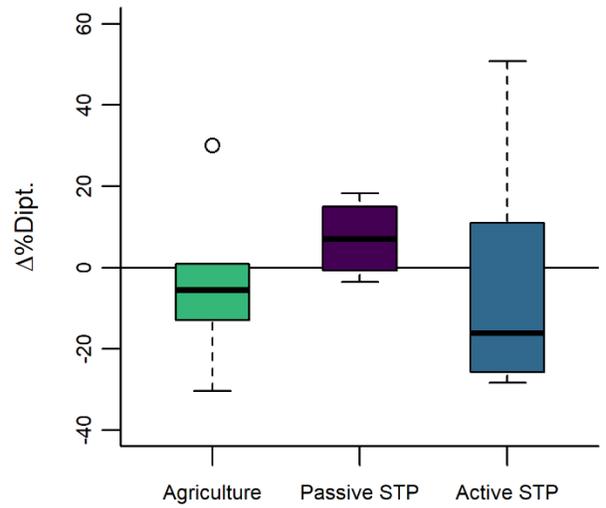
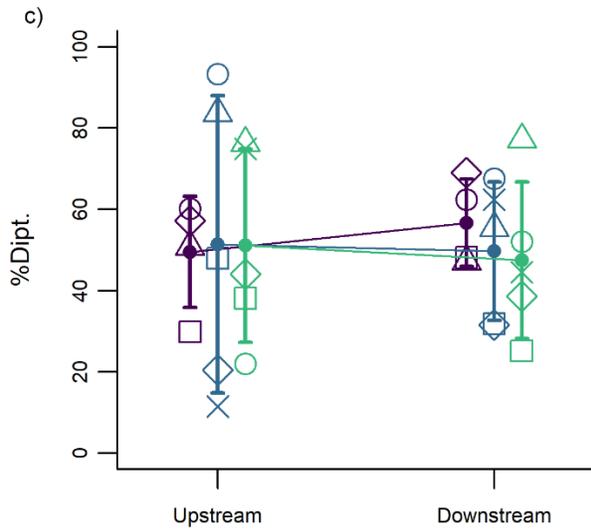
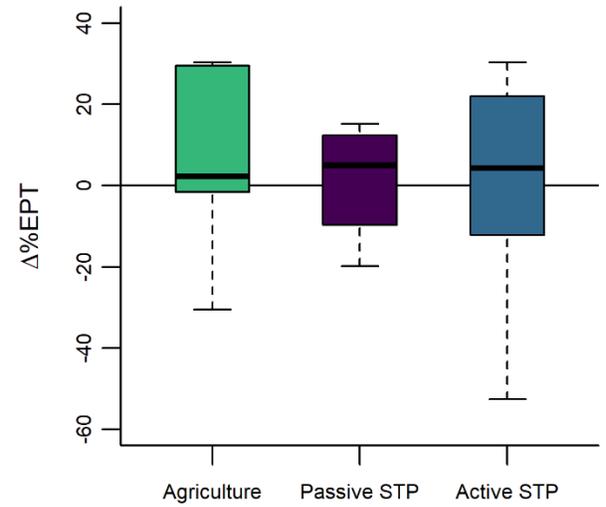
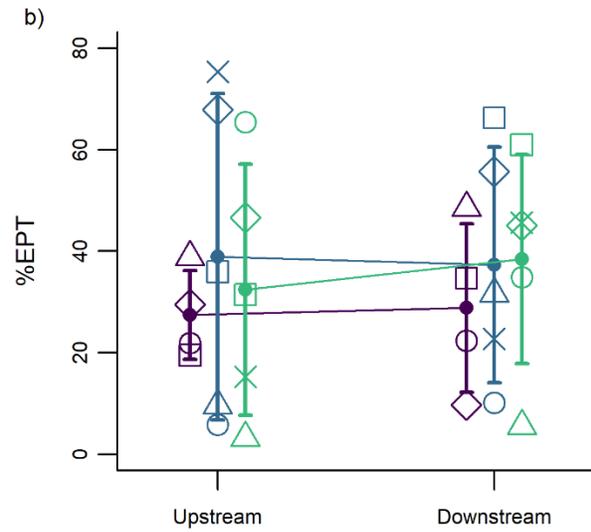
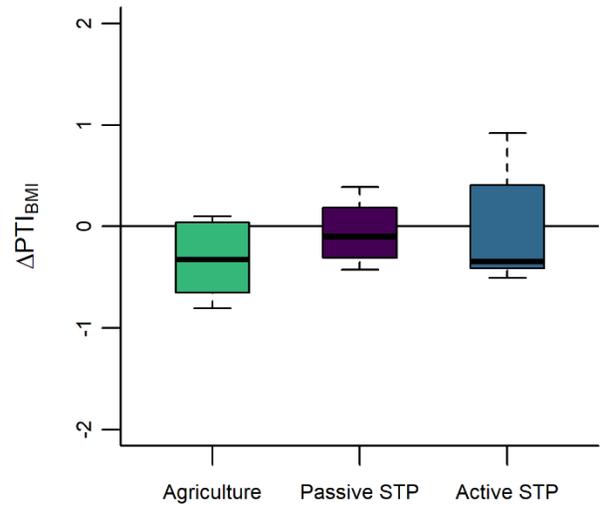
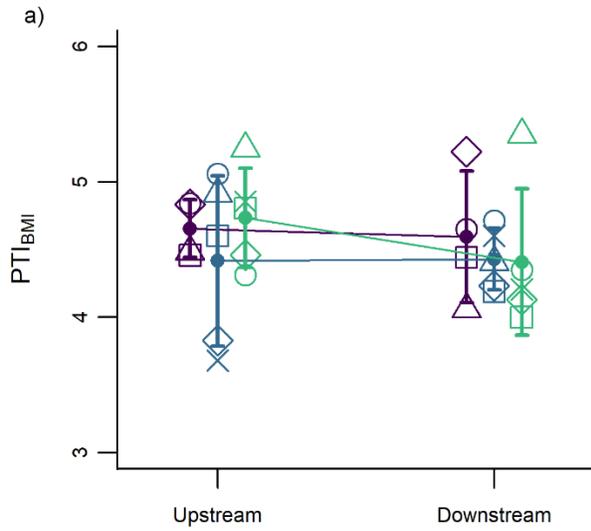
| Indicator | Predictor | Untransformed | | | | | Inverse Transformed | | | | |
|------------------------------|-----------|-------------------------------------|---------------|---------------------|--------------|---------------------|--------------------------------------|-----------------|---------------------|----------------|---------------------|
| | | 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 | | | |

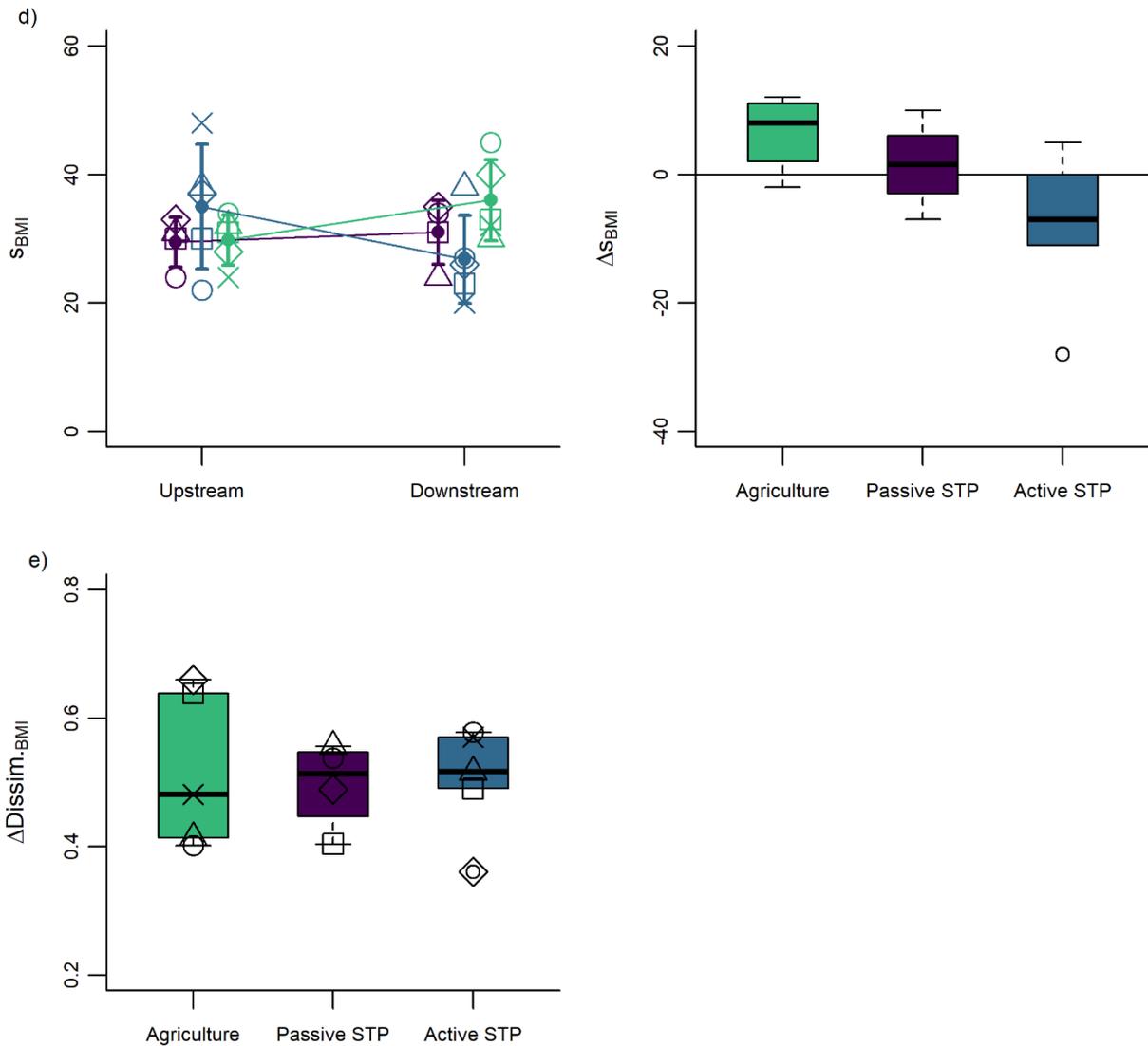




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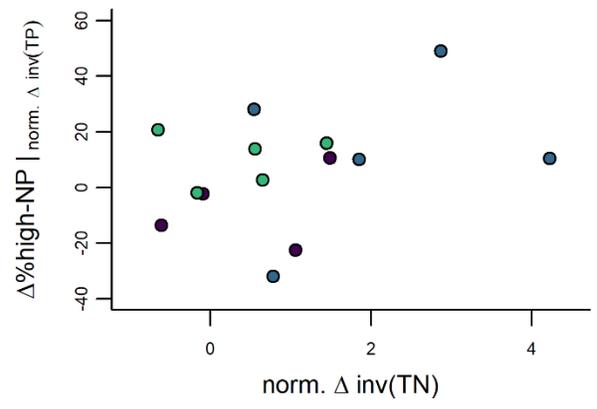
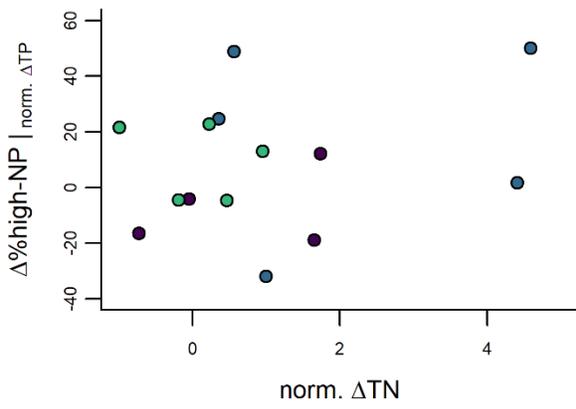
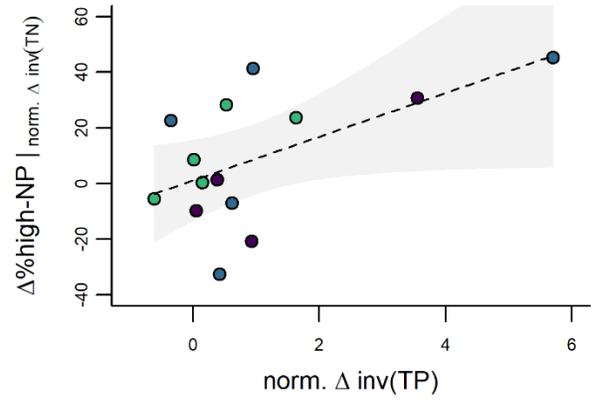
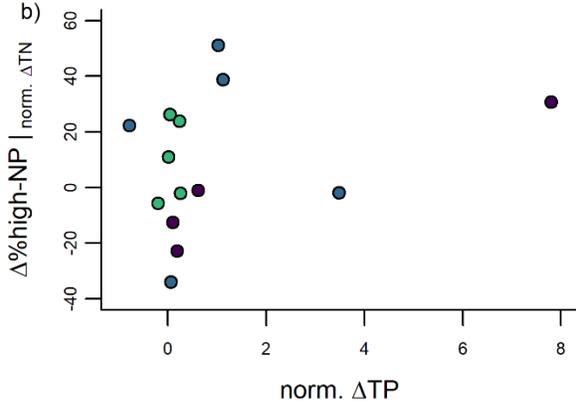
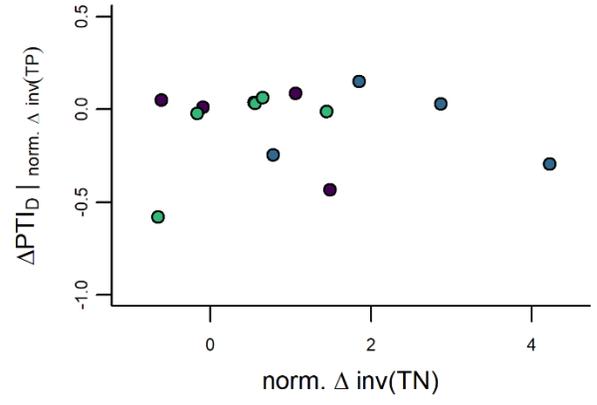
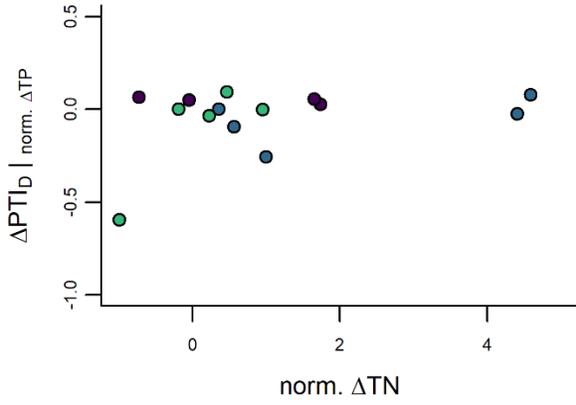
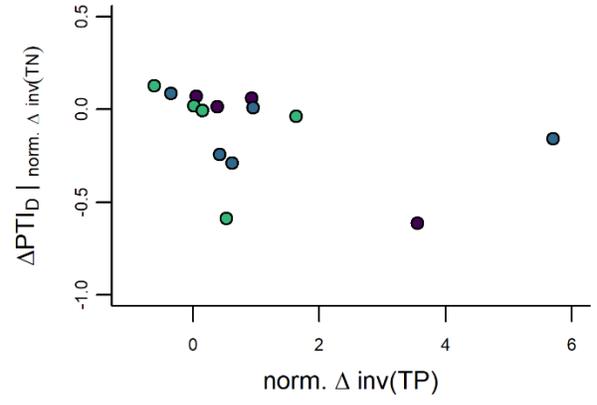
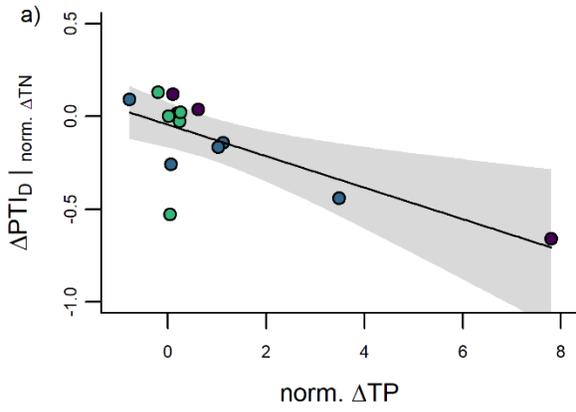
Figure S1-1: Mixed model analysis of variance for modified before-after (upstream-downstream) control-impact (agriculture-sewage effluent) assessment for ecological indicators of diatom assemblages: (a) pollution tolerance index (PTI_D), (b) percent abundance of high nutrient taxon (%high-NP), (c) percent 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.

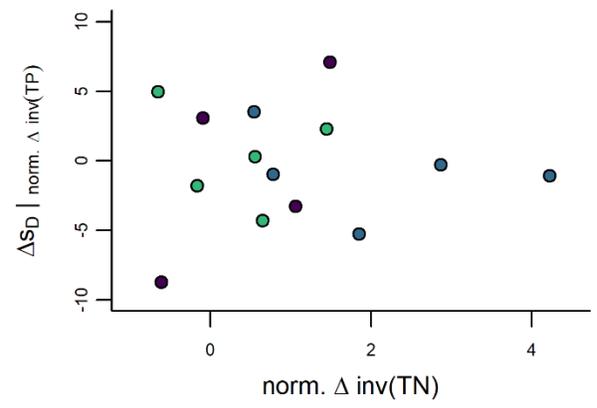
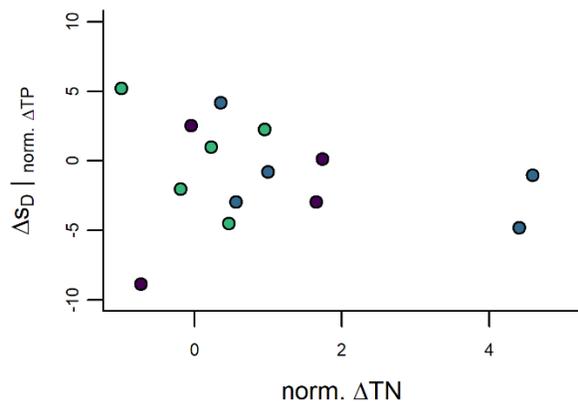
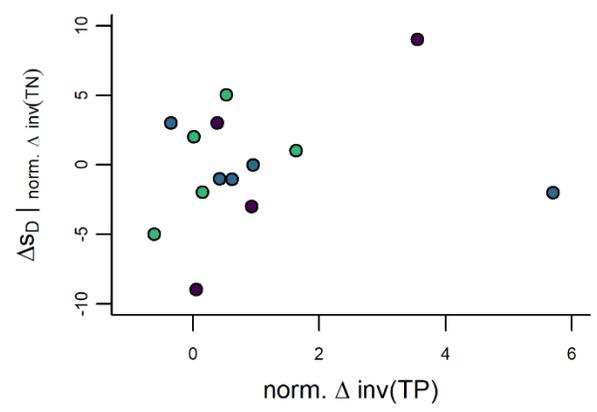
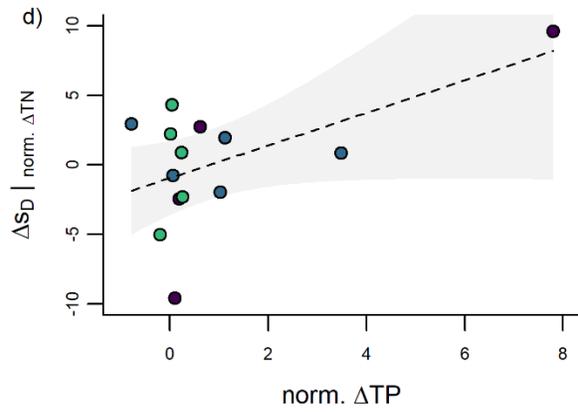
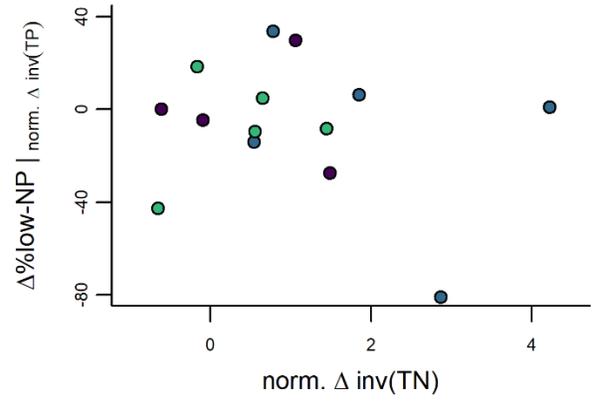
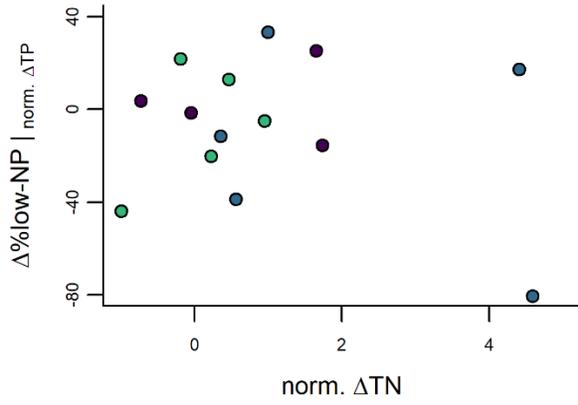
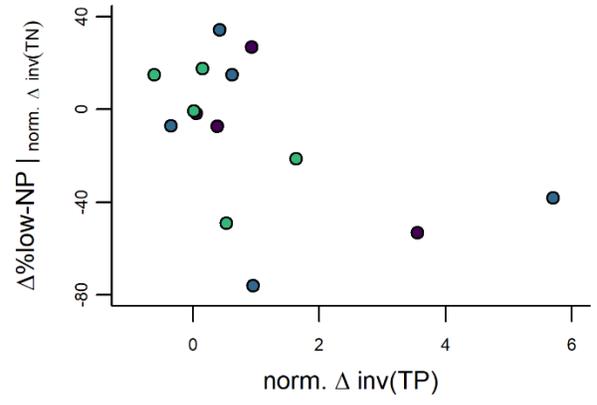
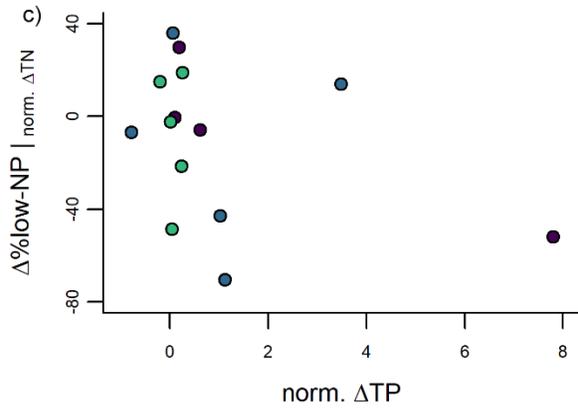


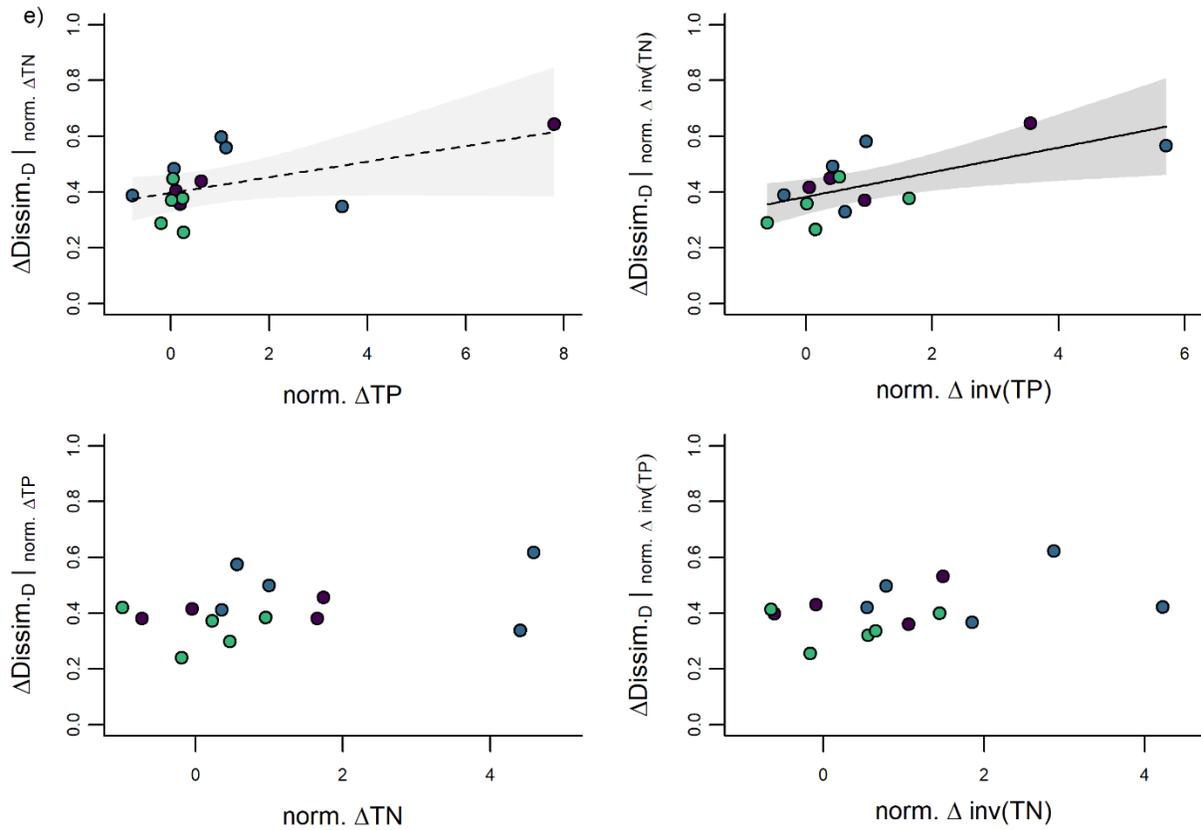


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Figure S1-2: Mixed model analysis of variance for modified before-after (upstream-downstream) control-impact (agriculture-sewage effluent) assessment for ecological indicators of benthic macroinvertebrate assemblages: (a) pollution tolerance index (PTI_{BMI}), (b) percent Ephemeroptera, Plecoptera, and Tricoptera 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.

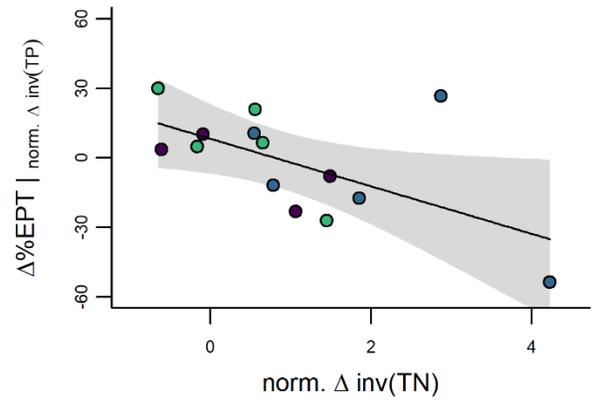
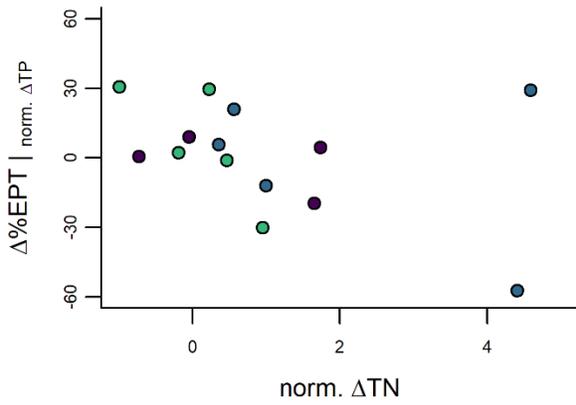
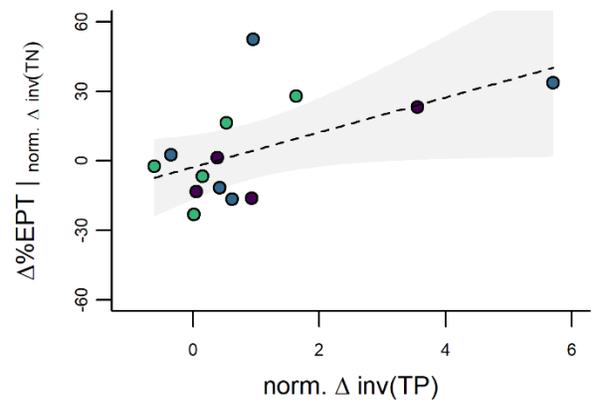
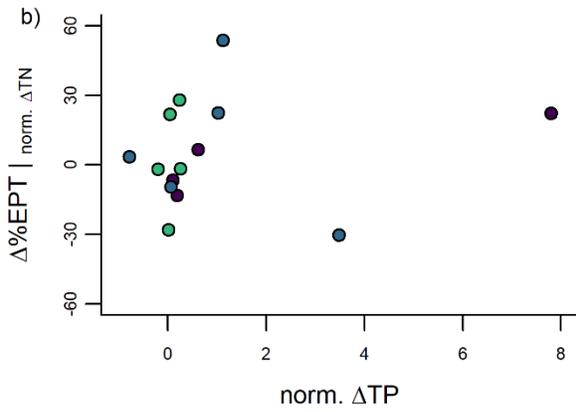
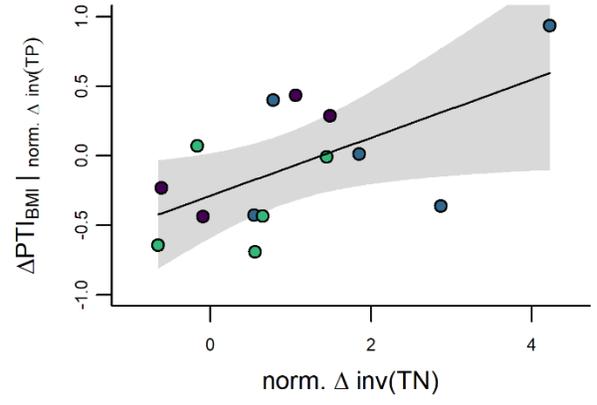
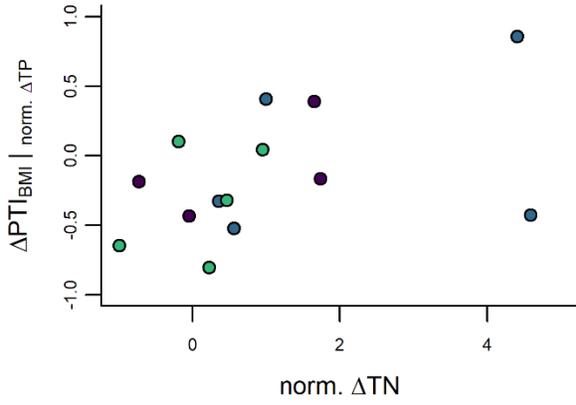
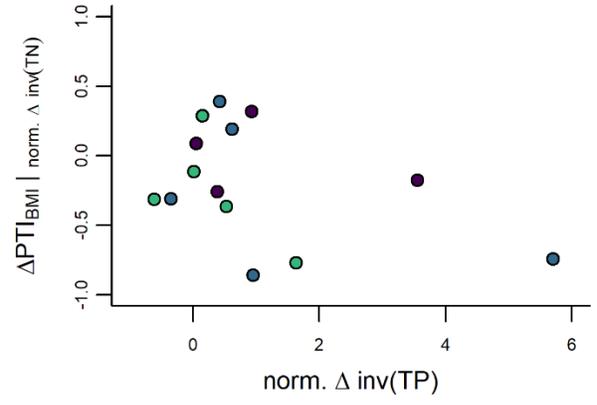
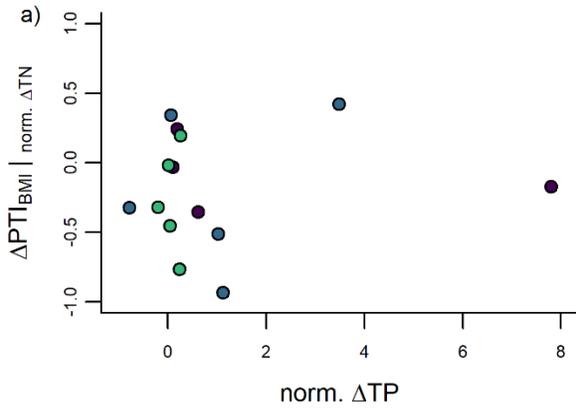


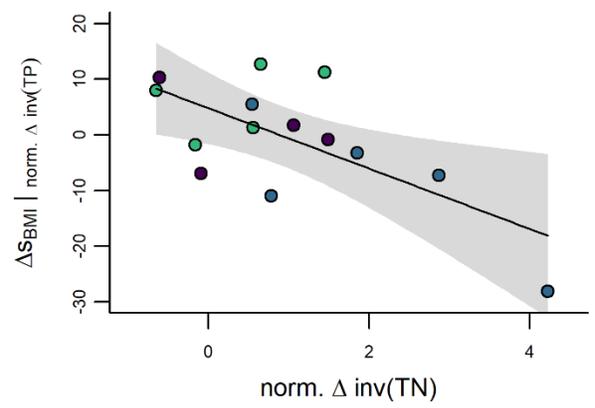
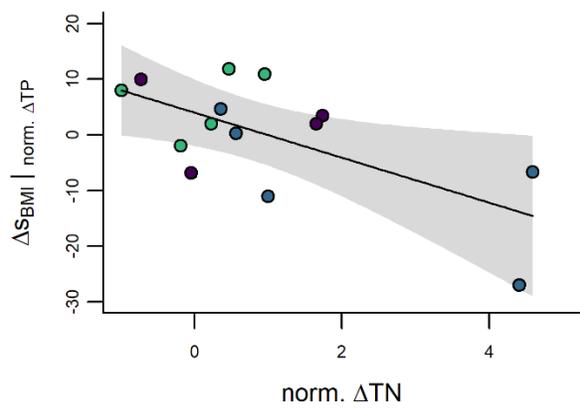
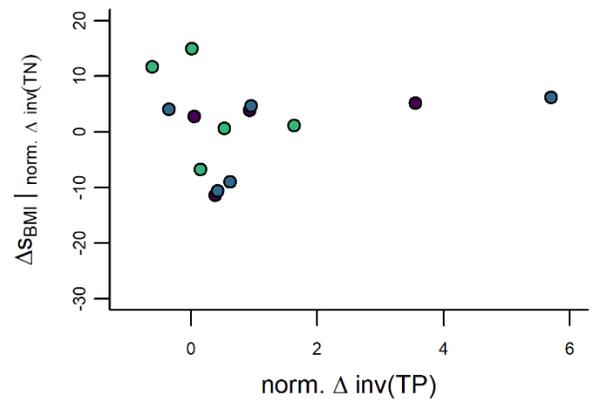
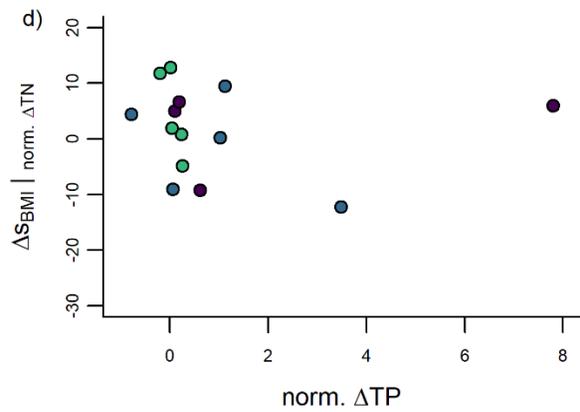
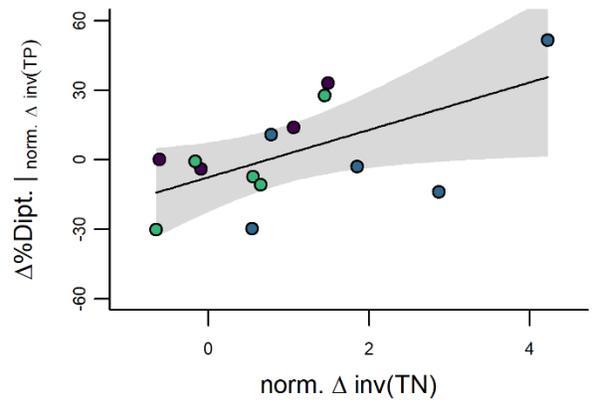
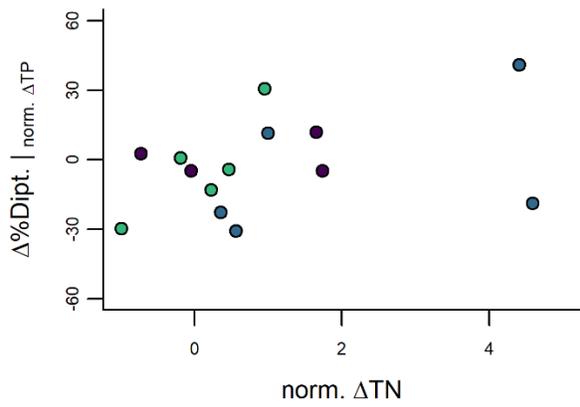
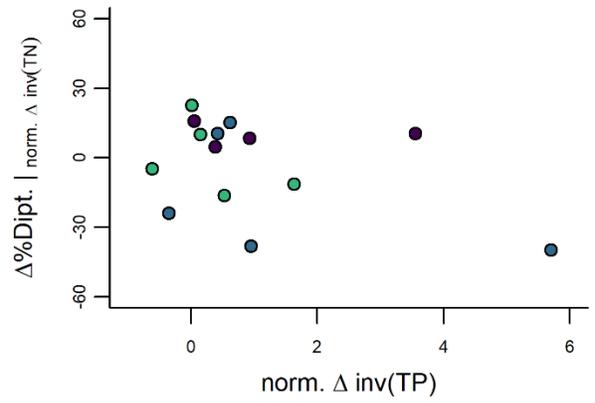
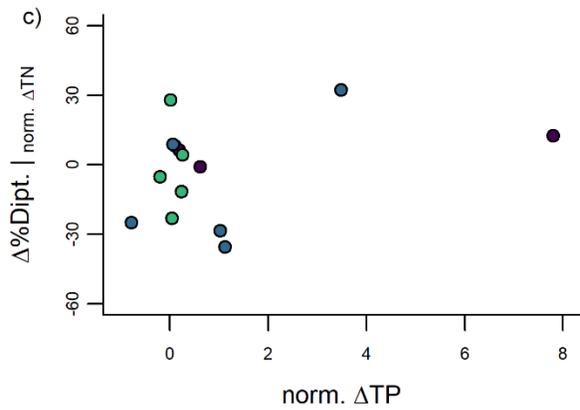


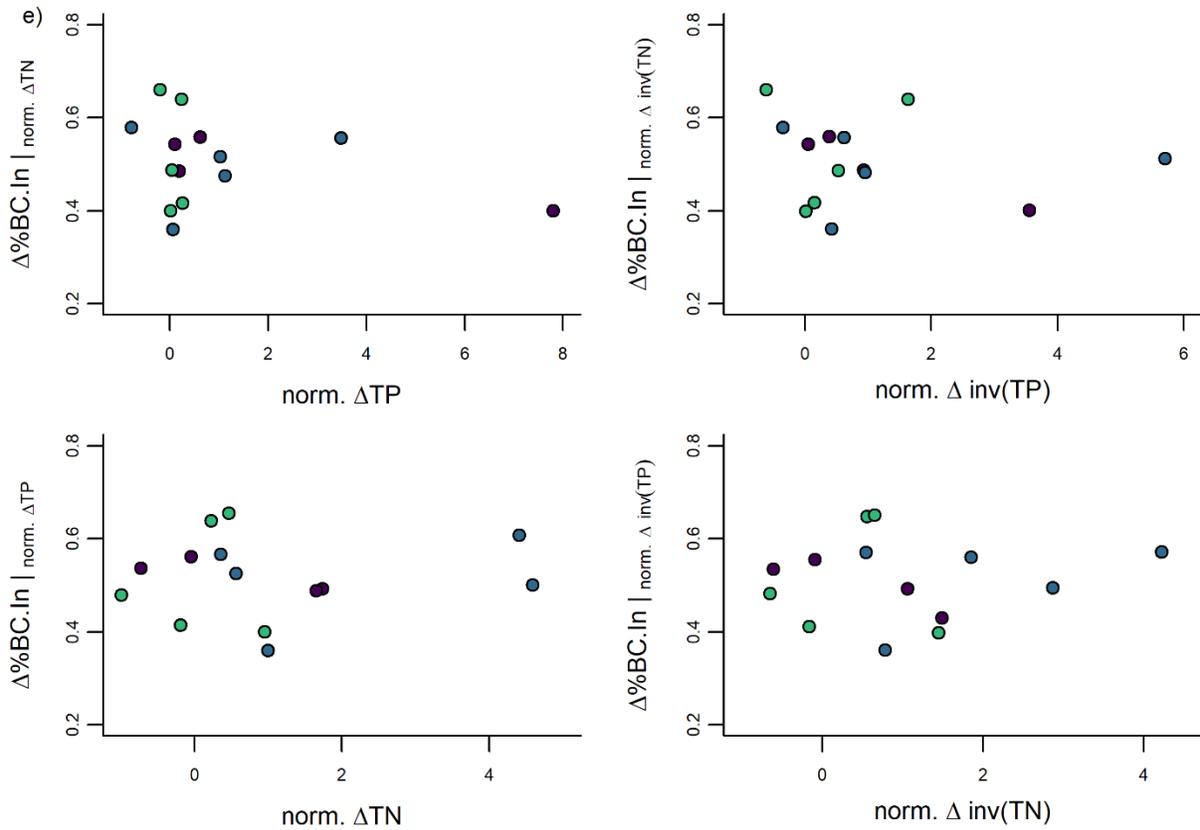


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Figure S2-1: Multiple regression models of the association between normalized (norm.) longitudinal differences (downstream – upstream) in untransformed and inverse transformed (inv.) total phosphorus (TP) and total nitrogen (TN) concentrations and 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 $\alpha = 0.10$.







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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_{BMI}), (b) percent Ephemeroptera, Plecoptera, and Tricoptera abundance (%EPT), (c) percent Diptera abundance (%Dipt.), (d) taxon richness (s_{BMI}), and (e) Bray-Curtis dissimilarity ($dissimilarity_{BMI}$). 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 $\alpha = 0.10$.

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