



## AN ABSTRACT OF THE THESIS OF

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Title: Response of Stream Macroinvertebrate Community to Canopy-opening  
Manipulations

Abstract approved: \_\_\_\_\_

Dana Warren

Stream light availability is an important factor influencing aquatic food webs. In forested headwaters, stream algal production is highly light-limited, and an increase in light often enhances benthic algal growth, which in turn increases food availability for primary consumers in the stream. In forested headwater streams, light availability is almost entirely mediated by the canopy structure of streamside vegetation. Over the last century, many streamside forests in the Pacific Northwest were heavily harvested, leaving dense regenerating stands for the time being. Under current conditions, the dense closed canopies, allow for limited primary production, and a low abundance of invertebrates that feed on stream algae. We investigated the response of benthic periphyton, stream macroinvertebrates, and prey consumption by trout to a release from light limitation in a paired-reach study design. We hypothesized that increased light availability promotes elevated

algal production which causes the invertebrate community to shift toward scraper dominance, and predicted that this change in community structure is detectable in the diets of trout. We found that the presence of a canopy gap had little influence on the invertebrate community, and this lack of change was not being masked by increased consumption of grazing invertebrates in summer trout diets.

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Response of Stream Macroinvertebrate Community to  
Canopy-opening Manipulations

by

Cedar Mackaness

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I understand that my thesis will become part of the permanent collection of Oregon State University libraries. My signature below authorizes release of my thesis to any reader upon request.

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# Introduction

In forested systems, streams and their biota are intrinsically linked to riparian vegetation (Vannote, Minshall, Cummins, Sedell, & Cushing (1980)). Stream food webs depend on direct carbon subsidies from the terrestrial environment in the form of both leaf litter and terrestrial invertebrates (Wipfli (1997)), but riparian controls on stream systems aren't limited to organic inputs. Riparian canopy cover also has an indirect effect on stream food webs through the control of light available for benthic primary production. In the Pacific Northwest (PNW) region of North America, riparian forests have changed substantially in the past half century. In response to a legacy of heavy harvesting (Pan et al. (2011)), riparian forest protections have created dense second-growth vegetation along streams in contrast with structurally complex old-growth forests containing multiple canopy gaps (D. R. Warren et al. (2016)). The dense vegetation in these regenerating forests decreases light availability and limits benthic primary production (M. J. Kaylor, Warren, & Kiffney (2017)). As forest stand development continues natural disturbances and individual tree mortalities will increase canopy heterogeneity through the introduction of gaps. To understand how aquatic food webs respond

to an increase in light associated with canopy gaps, we investigate the response of macroinvertebrates and fish feeding to canopy-opening manipulations.

Light, and its impact on primary productivity in streams is of particular interest because autochthonous carbon can be disproportionately represented in consumer biomass relative to its availability in aquatic systems (Lau, Leung, & Dudgeon (2009), McCutchan & Lewis (2002)). In forested headwaters specifically, basal carbon availability is dominated by leaf litter (McCutchan & Lewis (2002)); however, energetically, algae is a higher quality food source and is preferentially assimilated into higher trophic levels (Macarelli & others (2011)). Primary consumers mediate basal carbon availability for higher trophic levels, and in streams, primary consumers are dominated by macroinvertebrates, a major food resource for insectivorous fish. Because macroinvertebrates play a crucial role in mediating food web interactions, understanding their community dynamics and functional diversity can provide key insights into broader ecosystem functioning. Invertebrates in the scraper functional feeding group in particular have evolved specialized mouthparts for consuming benthic algal biofilms (periphyton), and increases in algal production in high light areas can elicit a positive response among these scraping taxa (Liess, Le Gros, Wagenhoff, Townsend, & Matthaei (2012)).

Macroinvertebrate community data have historically been used to evaluate stream health. Indices such as the B-IBI (benthic index of biological integrity) rely on total taxa richness and taxa richness of key families, such as Plecoptera, Ephemeroptera and Trichoptera, to evaluate the biological condition of streams. More broadly, an assessment of the whole community can be used to evaluate over-

all food web and ecosystem responses to a multitude of variables. For example, studies using nonmetric multidimensional scaling (NMS) has been used to assess community responses along a variety of environmental gradients (M. B. Cole, Russell, & Mabee (2003), Purcell et al. (2009)).

In headwater streams, the benthic invertebrate community represents the primary food source for trout, although this may vary seasonally (Syrjänen, Korsu, Louhi, Paavola, & Muotka (2011)). Interspecific interactions between trout and invertebrates can alter the benthic community and cascade to lower trophic levels (Peckarsky & McIntosh (1998)). In headwater streams trout are opportunistic foragers, eating whatever is available in their habitat. Cutthroat trout, the dominant fish species in Cascade headwaters, typically feed from the water column using visual cues to capture prey. Because salmonids are visual predators, their feeding efficiency can be influenced by light conditions and visibility (Wilzbach, Cummins, & Hall (1986)), therefore gaps have the potential to affect fish feeding not only through potential increases in scraper invertebrate food resources, but also by increasing foraging capture rates of all taxa and functional feeding groups.

Clear-cut harvests without buffers and the resultant reach-level increase in stream light can increase stream primary and secondary productivity, but increases in light also lead to increases in temperature, and cutting to the stream edge can increase sediment loads. Given these negative impacts, clear-cutting adjacent to streams is no longer a common practice in the Pacific Northwest. In unmanaged forests, and in recently implemented riparian buffers, stands are recovering from past land clearing, and are currently in the early to mid-seral stage of develop-

ment with dense homogenous canopy cover and low stream light (M. J. Kaylor et al. (2017)). Canopy gaps will begin developing naturally along streams as stands mature, and restoration efforts focused on emulating natural disturbance may expedite forest shifts toward late-succession and old-growth structural conditions (Kreutzweiser, Sibley, Richardson, & Gordon (2012)). While studies on reach-scale forest clearing demonstrate a clear response in benthic primary producers, invertebrates, and trout to release from light limitation (Murphy & Hall (1981)), this does not reflect future riparian conditions in most forested landscapes. Rather, as stands progress toward late successional forest structure, localized light patches (as opposed to large openings) will become increasingly prevalent. The more moderate influence of these small canopy gaps on stream ecosystems has not been widely investigated, especially in an experimental context.

We implemented a two-year long before-after, control-impact study designed to detect and accurately capture the effect of canopy gaps on aquatic ecosystems. In this work, we hypothesized that primary production would increase when canopy gaps were created, and this would cause the invertebrate community to shift toward more scraping taxa. However, we expect the response to be dampened in comparison to observed responses in large scale riparian clearing studies. In addition to evaluating the macroinvertebrate community, we quantified trout diets with the expectation that shifts in the invertebrate community would be reflected proportionally in the diet of opportunistic foraging of trout, but also to ensure that a potential signal in invertebrate scraper taxa was not being masked by increased foraging of apex consumers.

# Methods

## *Study location*

The study consisted of five reach pairs on five replicate streams in the western Cascade Mountains of Oregon. Each reach pair consisted of one treatment reach and one reference reach. Two of the reach pairs (W-100, W-113) are located on private Weyerhaeuser Co. land, and three (LOON, CHUCK, MCTE) are located on U.S. Forest Service land, one of which (MCTE) is situated in the HJ Andrews Experimental Forest. Stream reaches were 90 meters in length and treatment gaps were 20 to 40 meters in diameter and situated approximately around meter thirty of treatment reaches. Sites had a buffer between stream reach pairs to limit any effects of the upstream reach on downstream conditions.

All of the streams are wadeable, fish-bearing streams with bankfull widths of 1-6 meters. Fish-bearing streams were purposefully selected to provide management-relevant results for key species such as salmonids. Additionally, streams of this size comprise roughly 70% of total stream length in forested catchments. The streams run through 40-60-year-old riparian forests regenerating from previous harvest. These forests have a homogenous canopy structure with heavy understory shading,



Table 1: Study site attributes

Stream	Elevation (m)	Bankful Width (m)	Base flow (L s <sup>-1</sup> )	Latitude	Longitude
CHUCK	833.0	5.19825	21.0	43.95362	-122.1136
LOON	721.2	4.13170	12.5	43.95362	-122.1833
MCTE	867.0	2.20000	5.0	44.25454	-122.1667
W-100	441.0	5.39000	43.9	44.19813	-122.4930
W-113	537.0	3.30000	9.1	44.19289	-122.5107

as defined by their early to mid seral stage. Small streams also provide ease of sampling and maximize the effect of a canopy opening manipulation since small streams may be completely shaded by overhead vegetation due to their high edge to area ratio.

### *Study Design*

The before-after, control-impact (BACI) study design lends itself to experimental field studies by accounting for natural variations between sites. By taking the difference of a given variable between the paired reaches and comparing the change in the difference from pre to post-treatment years, we account for both spatial and temporal variation. For the BACI analyses, a sample unit refers to a whole stream including both treatment and reference reaches because the metric of interest for BACI is the difference between the two reaches. Therefore, we have five sample units with two repeated measures, pre and post-treatment. To test for effects of the gap treatment, we quantify and assess changes in the reach differences between the two years. Samples were collected during summer 2017 and summer 2018 with pre-treatment data gathered during summer 2017 and post-treatment

data gathered during summer 2018. Canopy gaps were cut in the treatment reach during the winter of 2017-18 to permit adequate time for response to the canopy manipulation at all sites besides MCTE. At MCTE gaps were cut at the end of summer 2017 after data collection.

### *Data Collection*

#### Light

Daily, photosynthetically active radiation (PAR) was estimated from fluorescence decay rate over a twenty-four hour period following methods in Warren et al. Fluorescein dyes were prepared by diluting to 400 g L<sup>-1</sup> with DI water and buffering with 40 g L<sup>-1</sup> of aquarium salt. Once the dye was prepared, we filled 3.7mL glass vials and stored them in the dark until deployment. At each study reach three replicate vials were deployed every five meters, and retrieved twenty-four hours later. Because fluorescence of fluorescein changes with temperature (Bechtold, Rosi-Marshall, Warren, & Cole (2012)), vials were left in the dark until they reached room temperature. Fluorescence was then measured using a fluorometer (Turner Designs, San Jose, California), and the twenty-four hour decay rate was converted to daily photosynthetically active radiation (PAR) using the relationship in (D. R. Warren, Collins, Purvis, Kaylor, & Bechtold (2017)).

## Chlorophyll *a*

In each study reach, three ceramic tiles (15 cm x 15cm) were placed every 10 meters and left for 4 weeks before they were collected to allow periphyton communities to establish. Tiles were placed in riffle sections at a depth of 10-25 cm to keep them from silting over. All tiles were deployed in mid-July, and the tiles were deployed at the control and treatment reaches of each stream at the same time to keep within unit measures consistent. After collection, tiles were kept in the dark, submerged in water for two hours to avoid potential photosaturation issues with the *in situ* chlorophyll *a* measurements. Chlorophyll *a* (abbreviated as Chla for the remainder of this text) concentrations were then quantified using a BenthosTorch™ (BBE Moldaenke GmbH), a portable field instrument used for the quantification of chlorophyll *a* fluorescence on different substrates.

## Benthic Invertebrate Sampling

Three benthic invertebrate samples were taken at each stream reach at meters 15, 45, 75, or the closest area with non-boulder substrate. Samples were collected once per year over the course of one week using a Surber sampler with a .09 m<sup>2</sup> sampling area. Substrate was disturbed to a depth of approximately four inches for one minute. The sample was then preserved in 95% ethanol for identification and enumeration in the lab.

In the lab, the three benthic samples per reach were combined into a single pooled sample for each reach. The pooled sample was then subsampled using a

Caton tray. Squares  $\frac{1}{30}$  the area of the Caton tray were randomly sampled until the cutoff of 300 individuals or greater was reached. Benthic invertebrates were then identified down to genus or the lowest taxonomic unit (LTU) for cryptic taxa such as Chironomidae primarily using Merritt, Cummins, & Berg (2008). Counts from subsamples were then converted to densities using the following formula:

$$\frac{1}{3 * s * 0.09} \quad (1)$$

where  $s$  is the fraction subsampled, 0.09 is the area of the Surber sampler in square meters, and the result is divided by three because three samples from meters fifteen, forty-five and seventy-five were pooled.

For NMS and other community analyses, singleton taxa (taxa occurring in only one reach) were removed from the original matrix and density values were log transformed to reduce the effect of abundant taxa (Chironomidae, *Baetis*, *Microcrasema*) on community relationships by applying the formula  $\ln(n + 1)$  where  $n$ . The resulting matrix of benthic invertebrates at the LTU level of identification (20 reaches by 64 taxa) was then used for analysis. Functional feeding groups were assigned using the trophic relationships of each taxon as identified in Merritt et al. (2008), and raw density values were used for FFG analyses because sparse densities were not a concern with aggregate functional groups.

During Chla tile collection at the two streams with snails as the dominant scraper, the number of snails (Juga) and cased caddisfly (observed taxa being Uenoidae and Glossosomatidae primarily) on each tile were recorded and then

removed. before taking readings with a BenthosTorch™.

## Trout Diets

Trout diets were collected during the post-treatment year. Trout diets were collected during three-pass depletion of fish population estimates and were only taken from a subset of fish greater than 100 mm in length. Fish were anesthetized using AQUA-S and gastric lavaged. Stomach contents were evacuated by injecting water into the fish stomach using a piece of small plastic tubing attached to a syringe. Diet samples were collected in filter paper and preserved in 95% ethanol for lab processing.

All trout diets were processed (9 to 13 diets per reach) with aquatic invertebrates identified down to the family level and terrestrial invertebrates identified to order. Because the number of fish dieted in each reach varied, the average of all fish diets was used. The resulting matrix was then filtered for aquatic species and appended to a matrix of 2018 benthic invertebrate families (10 reaches by 38 families), producing a matrix of 20 sample units (SU's) by 40 families consisting of both fish diets and benthic samples. Singleton taxa were then removed to create a matrix of combined diet and benthic families of 20 SU's by 36 families. At this point, the combined matrix was relativized by row maxima to compensate for the difference between benthic sampling—measured in density per m<sup>2</sup>—and fish diets.

## *Data Analysis*

### BACI Analysis

The BACI analysis was performed in R (R Core Team (2018)), and consisted of calculating reach-pair differences by subtracting the control reach value from the treatment reach value. Reach differences were calculated for light, chl<sub>a</sub>, total invertebrate density and invertebrate densities by functional feeding group. A paired t-test with 4 degrees of freedom was then performed for each metric by subtracting the reach difference from the pre-treatment year from the difference value in the post-treatment year for each stream assuming the difference between the two reach ratios should be zero.

### Community Analysis

Community analyses were performed in PC-ORD (McCune & Mefford (2016)) and R (R Core Team (2018)) using the Vegan package (Oksanen et al. (2018)). Blocked multi-response permutation procedure (MRBP) was used to assess differences between treatment and control reaches in the pre and post treatment years. MRBP was followed up with blocked indicator species analysis (ISA) to determine underlying taxa driving any grouping detected by MRBP. The combined benthic and diet community matrix was subsequently tested for any differences between treatment and control reaches and benthic versus diet taxa representation using the same MRBP and ISA methods.

To test for any pre-treatment reach differences in 2017, MRBP was run on 2017 data only with Treatment as the two a priori groups and blocked by Stream. The 2018 post-treatment data was then assessed using the same MRBP grouping and blocking. MRBP is a nonparametric method used to test for differences between groups. This method accommodates paired or blocked study designs by accounting for variation related to study design variables that have little bearing on the question being addressed. In this case, MRBP accounts for any between-stream variation. MRBP outputs a p-value for the observed within-group distance (smaller distances constituting stronger grouping) by shuffling SU's between groups to generate a distribution of possible within-group distances (McCune, Grace, & Urban (2002)).

The follow-up ISA calculates an indicator value (IV) for each species. The IV is a composite of a taxon's fidelity and exclusivity to a group. If a taxon is consistently abundant in one group and never present in any other, then it would receive a high IV. Conversely, a taxon rarely abundant in SU's of one group and present in other groups would receive a low IV (McCune et al. (2002)). A Monte Carlo test of 1,000 permutations of the taxa matrix was used to generate a p-value for each taxon's IV.

Nonmetric multidimensional scaling (Kruskal (1964)) was used to visually assess differences between the treatment and control reach communities, and quantify the relationship between the synthetic community variable and Chla . Sorensen distance was used for both ordinations to reduce the impact of outliers. The ordination was rotated to maximize the environmental variable Chla along axis 1. A

random start was used and the real data were run 250 times to ensure an absolute stress minima was reached. A Monte Carlo test with 100 permutations was used to generate a p-value for the probability that the final ordination has a lower than expected stress value.

#### Analysis of Trout Diets

Trout diets were collected in the post-treatment year, which limits analysis to a comparison of reference and treatment reaches without the BACI control on inherent reach differences. We performed paired t-tests for the abundance of each functional feeding group represented in the diets of trout in the reference and the treatment reach, and on the modified Ivlev's selectivity index (as defined in Jacobs (1974)) for each FFG.



# Results

## *Light and Chlorophyll*

In 2017, before treatment, the average daily PAR reaching the stream benthos among the five streams was consistently low in both reference and treatment reaches and there was an average difference between the treatment and reference reach of  $-0.16 \text{ mol m}^{-2} \text{ day}^{-1}$ . In 2018, after gaps were cut, light went up by  $2.60 \text{ mol m}^{-2} \text{ day}^{-1}$  on average in the treatment reach compared to the reference reach (Figure 1) resulting in a final yearly difference between reach differences of  $2.77 \text{ mol m}^{-2} \text{ day}^{-1}$  (p-value = 0.019, t-value = -3.83).

Again, for chl<sub>a</sub>, values across all sites in the pre-treatment year were low (mean =  $0.095 \text{ ug cm}^{-2}$ ), and there was little difference between reaches. After gaps were cut in the post-treatment year, Chl<sub>a</sub> values went up by  $0.44 \text{ ug cm}^{-2}$  in the gap reach, and only  $0.175 \text{ ug cm}^{-2}$  in the reference reach (final BACI difference =  $0.265 \text{ ug cm}^{-2}$ , p-value = 0.002).

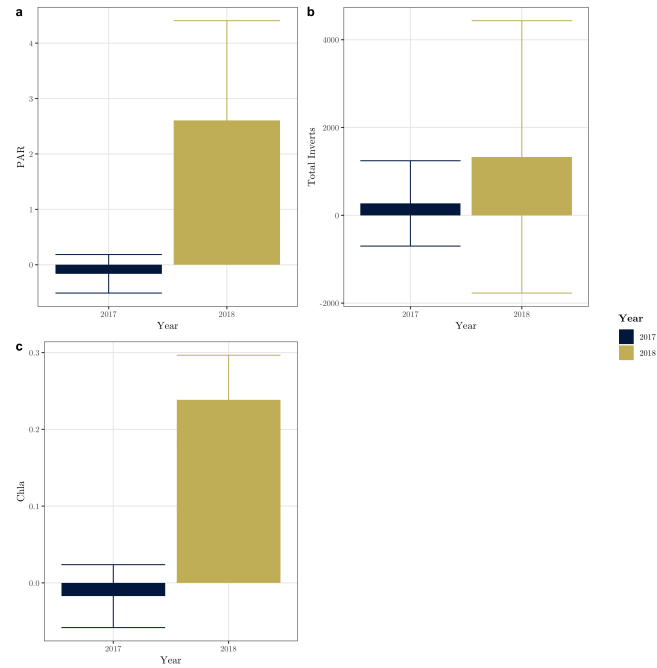


Figure 1: Light and Chla reach differences in the pre and post-treatment years

### *Juga on Tiles*

In the pre-treatment year, the average density of *Juga* on tiles among the two streams with *Juga* present was 24.44 snails per m<sup>2</sup> with little difference between the control and treatment reaches. In the post treatment year the average snail density in the treatment reach increased by 204.44 snails per m<sup>2</sup>, whereas snail density in the control reach only increased by 88.89 snails per m<sup>2</sup>. Snail abundance at these two streams was moderately associated with Chla ( $r^2 = 0.3204$ ,  $p = 0.00547$ ), but saw the largest BACI response in meters ten and twenty, slightly upstream of the gap treatment.

### *Benthic Invertebrate Community*

There was little difference between benthic invertebrate communities in the treatment and reference reaches in the pre-treatment year (MRBP:  $A = 0.041$ ,  $p = 0.071$ ), or the post-treatment year ( $A = -0.022$ ,  $p = 0.838$ ). The results from the NMS ordinations support the results of the MRBP (Figure 2).

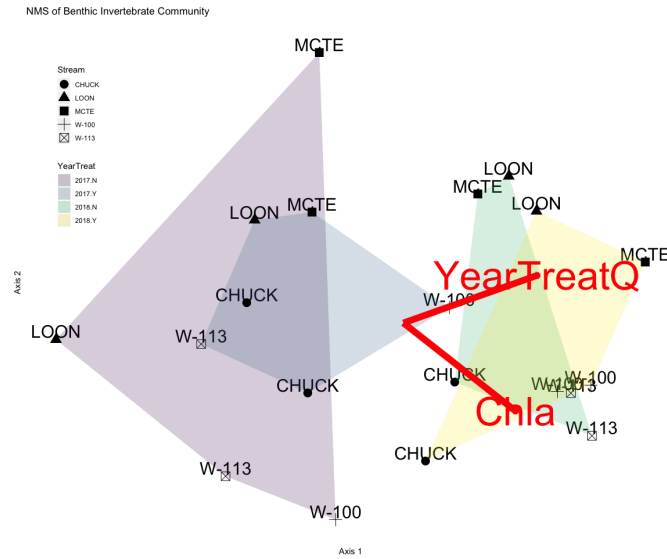


Figure 2: NMS of each reach in invertebrate community space. Each point represents a single stream reach, shapes identify stream, and color identifies treatment and year.

The NMS ordination of benthic invertebrates converged on a 2D solution with a final stress of 12.031. BenthoTotal (total chlorophyll values from the BenthosTorch) and YearTreatQ (a binary variable coded with 1's for 2018 treated reaches and 0's for all other reaches) both had positive  $r^2$  values with axis 1 (YearTreatQ  $r^2 = 0.272$ , BenthoTotal  $r^2 = 0.304$ ).

### **Add relationship of individual taxa to axes and ISA analysis.**

#### *Invertebrate Functional Feeding Groups*

Collector gatherers were by far the most abundant functional feeding group in the post-treatment year for both reaches at all sites. This does not appear to be due to the treatment of the gaps since we see heightened collector gatherer response in the reference reach as well. Collector filterers were typically the least abundant FFG in any stream or year. No FFG had a significant response across all streams. Scraping invertebrates only showed a positive response to the gap in MCTE with all other streams having a moderately negative BACI response. When we treat streams as independent replicates and perform a t-test of total invertebrate density response and the density response of each FFG individually, we find that only collector filterers had a statistically significant response (Table 2).

Table 2: BACI t-test results for various metrics

FFG	t-value	p-value
SH	0.0711617	0.9450224
P	0.5161519	0.6224931
SCe	-1.5465440	0.1624379
CG	0.5978468	0.5666133
SCi	0.8552979	0.4216123
CF	2.1321982	0.0656781
All Bugs	0.8400000	0.4300000

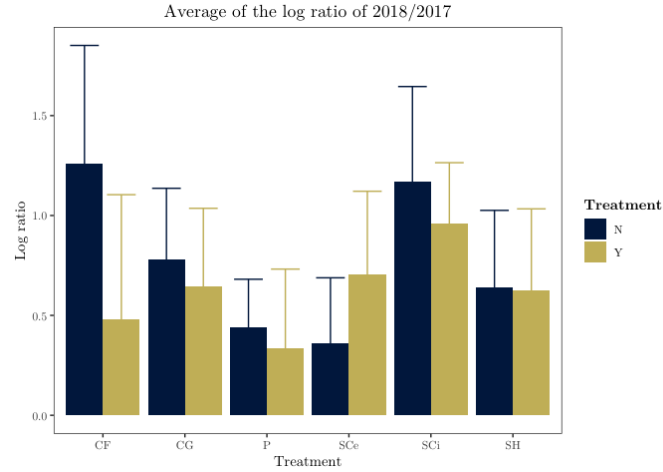
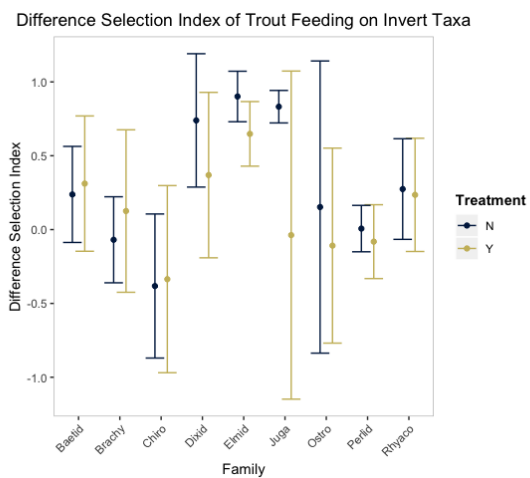
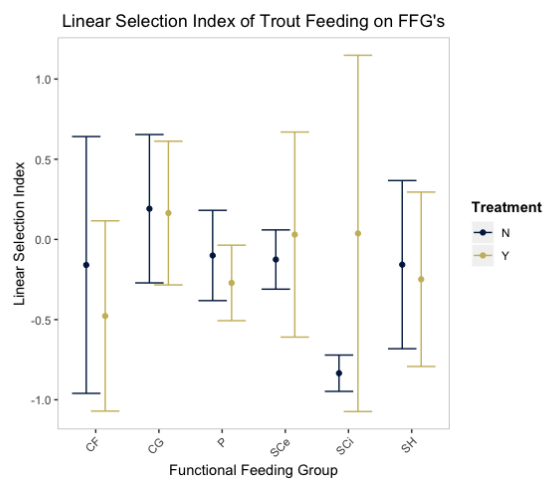


Figure 3: Log ratio of treatment reach divided by control reach for each FFG. CF = Collector-filterer, CG = Collector-gatherer, SH = Shredder, SC = Scrapers, SCe = Edible Scrapers, SCi = Inedible Scrapers, P = Predators, and All Bugs is the total macroinvertebrate density

# *Trout Diet*



# Discussion

*large idea to keep in mind*

Gaps are, by definition, open canopy patches in a larger forested system. While localized responses beneath a gap may occur, we were particularly interested in whether the effect of an individual canopy gap could be measured at the stream reach scale. Studies have found that systems with multiple gaps (Wootton (2012)), or with patches of high shade (Heaston, Kaylor, & Warren (2018)), had an effect on the overall invertebrate community. Yet, significant localized responses to increases in light beneath a single gap may not translate to significant system-wide responses at the stream or even the reach level. Our study design emphasizes the effects of gaps that comprise only a fraction of a stream reach, focusing on the integrated effect of small gaps embedded in a larger forested environment.

*General overview*

While light and Chla responded as expected to an opening of the riparian canopy, our reach-scale metrics for the invertebrate community do not show a response

to increases in primary production. This does not match our original hypothesis that increases in high quality algae would lead to increased abundance of scraping invertebrate taxa. Instead, we found the magnitude of the invertebrate response was not large enough to manifest at the reach scale and that fish consumption was most likely not masking changes in community composition or functional feeding group relative abundance. In the one taxa group for which we did evaluate local responses, snails, we saw a strong response to the treatment in meters adjacent to the gap treatment, but our reach-scale metrics for other invertebrate community members showed little BACI response.

#### *light and Chla*

While light increases in our study were large due to previous heavy shading, they were not outside the realm of what occurs naturally in these systems. While these small-scale disturbances may have localized impacts on stream biota, the reach scale response seems to be limited. Previous studies on forest shading, indicate a strong linear relationship between light and GPP, yet our increases in light show limited response. This may indicate potential photosaturation limiting algal response to gaps.

#### *Invertebrates FFG response*

The reach-scale metrics seem to be in contradiction with previous studies on stream light (Heaston et al. (2018), Kaylor & Warren (2017), Wootton (2012)), but these



studies focused on the immediate, within-treatment response of invertebrates and fish to various alterations to light availability. In that regard, our snails responded as expected, but the relative size of our canopy manipulations, one similar to small scale natural disturbances and individual tree mortality, limited reach-level trophic responses.

*Community, why we feel justified using FFG's*

Community PCA results what have others (in Dave's lab) found and what did we find here? How might these results change if you were to look at other seasons? Overall communities were more similar within a stream than across treatments. This is to be expected but we thought there would be a shift in the same direction that would show up. We did see trends in SC that fit our hypothesis, that were not significant, however what is interesting is that with such different communities between streams the shifts that we saw were a result of changes across a wide range of taxa, not just one taxa group changing. So even though there is skepticism about the FFG's and how they relate to diet (Revisiting RCC sites in ID Rosi and Baxter are co-authors), the trends from our sites suggest that applying an FFG here may be reasonable.

*Fish diets*

*Our goal is to relate light to various biotic responses using a bottom-up framework, what are limitations to this*

Complex trophic dynamics and limits on primary productivity such as photosaturation and nutrient limitation make it hard to capture a per-unit-light biotic response. Studies demonstrate clear differences between old-growth and second-growth light dynamics and system productivity, but how frequently or how large of gaps are necessary to get a response?

## Conclusion

Gaps do have an effect but it appears to be local – at least for gaps of this size and the effect is muted at the whole reach scale. But more gaps may give us a different answer since there is clearly some local impact.

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