

1 Direct and indirect influences of macrophyte cover on abundance and growth of  
2 juvenile Atlantic salmon

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20 Abstract

- 21 1. The relationships between macrophytes and the physical and biological characteristics  
22 of the environments that aquatic organisms inhabit are complex. Previous studies have  
23 shown that the macrophytes, *Ranunculus* (subgenus *Batrachium*), which are dominant  
24 in lowland chalk streams and widespread across Europe, can enhance juvenile  
25 Atlantic salmon abundance and growth to a greater degree than other physical and  
26 biological habitat characteristics. However, mechanistic understanding of how this  
27 effect might arise requires consideration of the direct and indirect relationships among  
28 habitat characteristics that are likely to be influenced by the presence of macrophyte  
29 cover.
- 30 2. We applied structural equation modelling to data collected during a two-year in-river  
31 manipulative experiment in the river Frome (southern England, UK) designed to  
32 quantify the magnitude and the relative importance of direct and indirect influences of  
33 *Ranunculus* cover and other physical and biological variables, including water  
34 velocity, water depth, prey biomass and body size, and abundance of con- and hetero-  
35 specifics, on abundance and somatic growth of 0+ salmon.
- 36 3. Results indicated a strongly positive direct influence of *Ranunculus* cover on salmon  
37 abundance, as well as positive influences of *Ranunculus* on velocity heterogeneity and  
38 water depth that are indirectly related to decreased salmon abundance. Interestingly,  
39 there was no indication of a direct influence of *Ranunculus* cover on salmon growth,  
40 although *Ranunculus* was indirectly related to increased salmon growth through its  
41 positive influence on prey biomass, an effect mediated by velocity heterogeneity and  
42 proportion of fast velocities.

43 4. These findings provide novel mechanistic insights into the key role of *Ranunculus* in  
44 their native lowland rivers to enhance abundance and improve conditions for multiple  
45 food web components. Strategies to maintain or enhance naturally occurring  
46 *Ranunculus* in these rivers are therefore likely to return wide ranging ecosystem  
47 benefits, including for species of high conservation value, such as salmon. These  
48 mechanistic impacts on habitat heterogeneity and ecosystem productivity could  
49 generalise to native macrophytes in other river systems, particularly where habitat is  
50 dominated by vegetation in the absence of large substrates.

51

52 1. Introduction

53

54 Terrestrial and aquatic vegetation form key ecosystem structures that provide resources,  
55 shelter and physical conditions that benefit a variety of species and functional groups (Tews  
56 *et al.*, 2004; Teixeira de Mello *et al.*, 2015; Wilkinson *et al.*, 2021). Density and morphology  
57 of plants often underly habitat heterogeneity (Diehl & Kornijow, 1998; Tews *et al.*, 2004;  
58 Warfe & Barmuta, 2004), which is essential for resilient, abundant and diverse animal  
59 communities (Kovalenko, Thomaz & Warfe, 2012; Teixeira de Mello *et al.*, 2015).

60 Macrophytes are considered to be such foundation species that strongly structure freshwater  
61 ecosystems (Scheffer *et al.*, 2015; Lürig, *et al.*, 2021) and their impacts on trophic  
62 interactions (e.g. Dionne & Folt, 1991; Warfe & Barmuta, 2004), water velocities and  
63 sediment retention (e.g. Wharton *et al.*, 2006) have been well studied. However, previous  
64 research has primarily focused on direct responses of animal abundances to macrophyte  
65 structures, with less consideration for how the habitat heterogeneity created by macrophytes  
66 modulates animal responses (Kovalenko, Thomaz & Warfe, 2012).

67 In lowland rivers, which are typically more stable and less dynamic than upland river systems  
68 (Berrie, 1992), macrophytes are particularly important for juvenile salmonids as they provide  
69 a key source of habitat heterogeneity in the absence of other large substrate structures (Berrie,  
70 1992; Riley *et al.*, 2009). While not studied as extensively as other habitat characteristics,  
71 positive interactions between juvenile salmonids and macrophytes in lowland rivers have  
72 been demonstrated across a wide geographical scale, including southern England (e.g. Riley  
73 *et al.*, 2009; Marsh *et al.*, 2021a), south-west Ireland (McCormick & Harrison, 2011),  
74 northern Finland (Maki-Petays *et al.* 2011), and north-eastern United States (Beland *et al.*

75 2004) and are unlikely to be limited to these areas. Atlantic salmon (*Salmo salar*, hereafter  
76 salmon) reach greater abundances and faster growth rates in habitats with high cover of the  
77 dominant lowland macrophytes known as water crowfoot, *Ranunculus* spp. (Marsh *et al.*,  
78 2020a, 2021a), although the mechanisms underlying these relationships remain unclear.  
79 Cover is important for juvenile salmon (Armstrong *et al.*, 2003; Marsh *et al.*, 2020a, 2021a),  
80 and so the vertical structure of *Ranunculus*, coupled with the floating canopy on the water  
81 surface, offers direct protection from both aerial and aquatic predators (Johnsson, Rydeborg  
82 & Sundstrom, 2004; Diehl & Kornijow, 1998), harsh environmental conditions (Gries &  
83 Juanes, 1998), and potential competitors (Venter *et al.*, 2008).

84 In addition to the direct influence of macrophytes as shelter, the impacts of macrophyte cover  
85 on individual salmon and populations might be mediated by physical habitat characteristics,  
86 inter- and intra-specific competition, and resource quality and quantity. The structural  
87 complexity of submerged macrophytes, particularly those with dense canopies such as  
88 *Ranunculus* (Cotton *et al.*, 2006), dramatically increases the resistance to water flow. This  
89 provides refuges of low water velocity and increased water depths, the latter of which might  
90 be avoided by juvenile salmon due to increased presence of predators and larger conspecifics  
91 (Gibson, 1993; Wharton *et al.*, 2006, Marsh *et al.*, 2020a). As water velocities slow within  
92 the plant stands, water flow is accelerated around the plant beds, forming adjacent areas of  
93 fast velocities (Wharton *et al.*, 2006). This results in heterogeneous water velocities, a  
94 desirable habitat characteristic for juvenile salmonids to exploit for energy-efficient foraging  
95 opportunities, with low velocity refuges alongside fast water velocities providing increased  
96 supply of drifting prey, which might otherwise be energetically demanding environments to  
97 forage in (Wankowski & Thorpe, 1979; Morantz *et al.*, 1987).

98 Heterogeneity in water velocities and depths can promote more abundant and diverse  
99 macroinvertebrate communities, which juvenile salmon feed upon (Keeley & Grant, 1997;

100 Riley *et al.*, 2009). The reduced water velocities within plant stands result in increased  
101 retention of fine sediment and organic material (Gregg and Rose 1982; Cotton *et al.*, 2006),  
102 providing food for macroinvertebrate communities dominated by larger individuals. The  
103 complexity provided by plant stands can also offer macroinvertebrates protection from  
104 predators (Dionne & Folt, 1991) and thus, enables the coexistence of larger prey individuals  
105 and their fish predators. Consequently, *Ranunculus* communities are often associated with  
106 productive and abundant macroinvertebrate communities (Armitage & Cannan, 2000; Riley  
107 *et al.*, 2009).

108 Teasing apart the direct and indirect influences of these habitat components to inform  
109 fisheries management is challenging due to the interdependence among habitat factors  
110 (Armstrong *et al.*, 2003), however, detailed understanding of the complex relationships  
111 between *Ranunculus* and juvenile salmon populations is essential if we are to apply effective  
112 management strategies. Populations of Atlantic salmon have suffered unprecedented declines  
113 in the last few decades (ICES, 2021). Natural recovery of salmon populations is hindered by  
114 a decline in both the numbers of juvenile salmon migrating to sea, and, in some regions, the  
115 condition of the individuals prior to migrating. Recent works have quantified the relationship  
116 between size of salmon pre-migration and the likelihood of individuals returning to  
117 freshwater to reproduce (Armstrong *et al.*, 2018; Gregory *et al.*, 2019). Individual body  
118 condition is strongly influenced by habitat conditions experienced during the first summer for  
119 juvenile salmonids (Zabel & Achord, 2004), prompting calls for a better understanding of the  
120 habitat conditions that maximise numbers and quality of juveniles (Armstrong *et al.*, 2018;  
121 Gregory *et al.*, 2019). Macrophytes, such as *Ranunculus*, have also suffered population  
122 crashes in lowland rivers within the last three decades, likely as a consequence of  
123 anthropogenic impacts including flow modifications and increased agricultural inputs  
124 (Cranston, 2004; Cotton *et al.*, 2006). There is, therefore, an urgency to understanding the

125 role of *Ranunculus* in promoting healthy juvenile salmon populations, to improve ecosystem  
126 management benefiting salmon and anticipate the implications of macrophyte declines on  
127 salmon productivity.

128 Here, we analysed detailed data from an in-field *Ranunculus* manipulation experiment  
129 designed to quantify ecological mechanisms governing the previously observed positive  
130 impact of *Ranunculus* cover on salmon abundances and somatic growth (Figure 1, Marsh *et*  
131 *al.*, 2021a). We aimed to determine whether these relationships represent direct influences  
132 through a source of cover, or indirect via influences on the wider ecosystem dynamics  
133 through impacts on physical habitat, food resources and/or competition. We hypothesised that  
134 abundant *Ranunculus* positively influences heterogenous physical habitat characteristics  
135 (increased water depth and velocity heterogeneity), the amount of basal resources and prey  
136 (organic material and macroinvertebrate prey biomass and body size), but also increases the  
137 potential for intra- and inter-specific competition via increased abundance of juvenile salmon  
138 and brown trout (*Salmo trutta*) respectively (Figure 2). We subsequently postulated indirect  
139 influences of *Ranunculus* on salmon abundance through a positive influence of velocity  
140 heterogeneity, fast velocities and prey biomass, and a negative influence of increased water  
141 depths (and associated higher predation risk) on salmon abundance (Figure 2a). We also  
142 hypothesised indirect influences of *Ranunculus* on salmon growth through a positive  
143 influence of velocity heterogeneity and prey biomass and body size, as well as a negative  
144 influence of fast velocities (and associated reduced foraging efficiency) and higher  
145 abundances of salmon and trout competitors (Figure 2b).

146

## 147 2. Methods

### 148 Experimental design

149 The in-field *Ranunculus* manipulation experiment took place over two years (2016-2017) on  
150 the North Stream, an anabranch of the river Frome, a lowland river situated in Dorset, UK  
151 (Figure 3). Three blocks, approximately 100 m in length, were selected on the North Stream  
152 in locations with similar physical habitat, existing *Ranunculus* beds and no riparian canopy.  
153 At each block, three sites, 20 m in length, were designated and each assigned a different  
154 *Ranunculus* treatment, which was achieved by digging out and replanting *Ranunculus* plants  
155 between sites. To simulate the natural mosaic of *Ranunculus* beds, we replanted plants in  
156 random patches within the site. Following manipulation of the existing *Ranunculus* plant  
157 cover in spring (March/April) of each year, the treatments of low (<10%), medium (30-40%)  
158 and high (>60%) cover were maintained throughout the year, producing a gradient of  
159 *Ranunculus* cover until the plants began natural senescence in autumn (see Marsh et al.,  
160 2021a for full details of the manipulation procedure). On four sample occasions between June  
161 and October of each year, juvenile salmonid populations, aquatic macroinvertebrates and  
162 basal resources were sampled, and *Ranunculus* cover, water velocity and depths were  
163 quantified. Previous findings from this experiment showed that both salmon abundance and  
164 growth were enhanced by *Ranunculus* during summer months (Marsh *et al.*, 2021a, Figure 1).  
165 We used these data to investigate potential direct or indirect effects of *Ranunculus* cover on  
166 juvenile salmon (Figure 2).

### 167 Salmon abundance and growth

168 Juvenile salmon (0+ or young-of-the-year) abundance and biometric data were collected  
169 between June and October of each year at each of the nine sites (see Marsh et al., 2021a for  
170 full fish sampling procedure). We repeatedly electrofished each site in a upstream direction,  
171 capturing and removing all juvenile salmonids encountered, until two consecutive fishing  
172 passes yielded zero salmonid captures. We set stop nets at the downstream and upstream site  
173 limits to maximise capture efficiency. Captured individuals were sedated, identified to



174 species, weighed (to nearest 0.2 g), and marked with a Passive Integrated Transponder (PIT)  
175 tag (12.5 mm length, 2.12 mm diameter, Biomark, Idaho, USA), which was inserted into the  
176 body cavity, and the adipose fin clipped to indicate a tagged individual. Growth of individual  
177 salmon was calculated using change in the weight (g) measurements of tagged individuals  
178 that were recaptured in the same site, with the assumption that the individuals had been  
179 exposed to the site-specific *Ranunculus* cover for the duration of the growing period  
180 measured.

### 181 Physical habitat variables

182 Physical habitat variables were recorded at 25 evenly spaced quadrats (0.5 x 0.5 m) per site,  
183 on each sample occasion. *Ranunculus* cover was estimated as the percentage quadrat cover,  
184 water depth was measured (cm), and water surface velocity was estimated visually as one of  
185 five categories [1 = 0-25 cm s<sup>-1</sup>; 2 = 25-50 cm s<sup>-1</sup>; 3 = 50-75 cm s<sup>-1</sup>; 4 = 75-100 cm s<sup>-1</sup>; 5  
186 >100 cm s<sup>-1</sup>]. We calculated the site-level proportion of fast velocities as the proportion of  
187 velocity categories 3, 4 and 5 recorded in quadrats at a site. Measures of *Ranunculus* cover,  
188 water depth and proportion of fast velocities were averaged to create a site-level measure of  
189 each variable. The steepness in velocity gradients between a quadrat and its neighbouring  
190 quadrats was calculated using the Terrain Ruggedness Index (TRI, see Marsh et al., 2020a for  
191 details). Absolute values of TRI were averaged to represent site-level velocity heterogeneity.

### 192 Prey and basal resources

193 To determine biomass and size of macroinvertebrate prey, and the resources potentially used  
194 by the prey (basal resources), we sampled the benthos at three randomly selected locations  
195 per site, on each sample occasion. Using a Surber sample (0.25 m x 0.25 m, mesh aperture  
196 250 µm), we disturbed the substrate by hand for 30 seconds to collect macroinvertebrates and  
197 organic material. Samples were preserved in 70 % ethanol solution for sorting, identification  
198 and measuring individual macroinvertebrates, and processing of the organic material. All

199 macroinvertebrate specimens were identified to the lowest practical taxonomic level (usually  
200 species), counted and measured. We used macroinvertebrate measurements to estimate  
201 biomass of each taxon (Table S1) and multiplied the mean biomass by the abundance of each  
202 taxon to generate a measure of prey biomass. We summed the biomass of each taxon and  
203 calculated the average biomass of measured individuals within each Surber sample to  
204 produce measures of total prey biomass and mean prey size, respectively. After  
205 macroinvertebrate specimens had been picked from the sample, the remaining organic  
206 material was dried to constant mass (70°C, 72 h) and weighed, prior to being combusted  
207 (550°C, 4 h) and re-weighed. The ash-free dry mass of organic material was calculated by  
208 subtracting the combusted weight from the dried weight, producing a measure of basal  
209 resources. Measures of total prey biomass, mean prey size, and basal resources were averaged  
210 across the three Surber samples to create a site-level variable of each.

211 Using data from the summer months in which *Ranunculus* had a positive relationship with  
212 both salmon abundance and growth, led to the use of August and September data to  
213 investigate salmon abundance ( $n = 36$ , i.e., abundance measures in three sites in each of the  
214 three blocks at two months for two years, treated as independent data points), and June and  
215 August data to investigate salmon growth ( $n = 71$ ). Explanatory variables used in the growth  
216 analysis were averaged across June and August to represent average conditions during the  
217 growth period.

#### 218 Data analyses

219 To investigate whether and how the effects of *Ranunculus* cover on salmon abundance and  
220 growth were mediated by changes in the physical habitat, prey resources and/or competition,  
221 we applied confirmatory path analysis (Grace, 2006) in a piecewise structural equation  
222 modelling (SEM) framework (Lefcheck, 2016). SEM is a quantitative framework able to

223 quantify and partition the relative direct and indirect conditional associations of key habitat  
224 characteristics on natural populations (Grace, 2006). This approach has been commonly  
225 applied to investigating complex theory-supported hypotheses about ecosystems and  
226 ecological scenarios. Piecewise SEM fits multiple component sub-models within a global  
227 model to allow for testing of direct and indirect effects within a single causal network  
228 (Shipley, 2009; Lefcheck, 2016). The component models are solved separately and are  
229 therefore appropriate for conventional requirements of generalised mixed models (Shipley,  
230 2000; Shipley, 2009). We constructed the global path model for each main response variable  
231 (abundance and growth) based on previously determined relationships (Marsh et al., 2021a),  
232 and other hypothesised drivers of salmon abundance and growth (Table 1).

233 Firstly, we tested whether the observed positive influence of *Ranunculus* cover on salmon  
234 abundance was best represented as a direct effect of cover, or an indirect effect operating  
235 through changes to the physical environment and prey resources associated with *Ranunculus*  
236 cover (Figure 2a). The path model consisted of a set of linear regressions (identity-link  
237 function), describing the variation in variables representing physical environment or prey  
238 sources, and a negative binomial regression (log-link function), describing the variation in  
239 salmon abundance. A negative binomial distribution was used to account for potential  
240 overdispersion in the abundance data with log site area (m<sup>2</sup>) included in the model as an  
241 offset to account for variation in abundance that might be due to differences in habitat size  
242 (O'Hara & Kotze, 2010). Model residuals were inspected to ensure assumptions of  
243 homogeneity, normality and independence were not violated.

244 Secondly, we investigated whether the observed positive effect of *Ranunculus* cover on  
245 salmon growth in weight was best represented as a direct effect of cover, or an indirect effect  
246 operating through changes to the physical environment, prey resources and/or abundance of  
247 competitors (Figure 2b). This path model consisted of a set of linear and negative binomial

248 regressions to describe physical habitat, prey resource and salmonid abundance variables as  
249 described above, and a linear mixed effect model (identity link function) to describe variation  
250 in salmon growth in weight. The growth model included an offset term of log initial weight to  
251 account for the influence of initial weight on growth potential and a random intercept of site  
252 to account for multiple measures (individuals) at each site.

253 Path models for both salmon abundance and growth were fitted and model performance was  
254 evaluated using R package piecewiseSEM (Lefcheck et al., 2018), which applies Shipley's  
255 (2000) directed-separation test to determine if all variables are conditionally independent and  
256 identify any important missing links in the path model. Any missing links that were indicated  
257 by the analyses and ecologically justifiable were incorporated into the model. Non-significant  
258 links were dropped to ensure they did not have an undue influence on the remaining paths in  
259 the system before refitting the model (Tables S2 – 3 outline the specific process of removal  
260 and addition of links). The goodness-of-fit of models fitted to individual response variables  
261 were summarized using marginal  $R^2$  (and conditional  $R^2$  where random effects were  
262 included), and overall SEM adequacy was evaluated with the approximately distributed Chi-  
263 squared Fisher's C-statistic (Shipley, 2009). Prior to model fitting: (1) collinearity of  
264 explanatory variables was assessed using Pearson's correlation, none of which were highly  
265 correlated ( $r \geq |0.7|$ , Dormann *et al.*, 2013, Figures S1 –2); and (2) all numerical explanatory  
266 variables were standardised prior to analyses by subtracting their mean and dividing by their  
267 standard deviation. All statistical analyses were performed in R version 4.0.5 (R  
268 Development Core Team, 2021), using R packages lmerTest (Kuznetsova, Brockhoff &  
269 Christensen, 2017) and MASS (Venables & Ripley, 2002), in addition to those previously  
270 mentioned.

271 We of course recognize that scientific inferences based on quantitative models, as applied in  
272 this study, depend on a number of mechanistic assumptions, as recently described by Grace

273 and Irvine (2020). In this study, the most general and vital assumptions associated with  
274 network structure that we have included in our models are consistent with both known and  
275 plausible mechanisms whereby *Ranunculus* can influence physical and other biological  
276 system components. The use of experimental data in this study is a great asset in supporting  
277 interpretations. As with all ecological studies, estimates are assumed to be approximate and  
278 the confirmation of conclusions dependent on subsequent studies.

279

280

### 281 3. Results

#### 282 3.1. Salmon abundance

283 Results indicate a strong and positive direct influence of *Ranunculus* cover on salmon  
284 abundances ( $\gamma = 0.52, p < 0.001$ ) and an overall positive influence of *Ranunculus* (Table 2a),  
285 which was mediated through water velocities and depths (Figure 4a). *Ranunculus* cover was  
286 positively associated with increased velocity heterogeneity ( $\gamma = 0.46, p < 0.01$ ), which had a  
287 weakly supported, negative association with the proportion of fast velocities ( $\beta = -0.30, p =$   
288  $0.08$ ). Proportion of fast velocities in turn had a strongly supported, positive influence on  
289 salmon abundances ( $\beta = 0.34, p < 0.01$ ). *Ranunculus* cover was also positively associated  
290 with increased water depths ( $\gamma = 0.40, p = 0.02$ ), which was negatively related to salmon  
291 abundance ( $\beta = -0.29, p = 0.02$ ). There was no significant influence of velocity heterogeneity  
292 or prey biomass on salmon abundance, and *Ranunculus* cover was not related to basal  
293 resources or prey biomass. Consequently, these pathways were not retained in the final  
294 model, though there was a positive relationship between basal resources and prey biomass ( $\beta$   
295  $= 0.31, p = 0.06$ ). The model was an adequate representation of the data (i.e. all paths were  
296 judged to be independent: Fisher's  $C_8 = 6.97, p \text{ value} = 0.54$ ), and explained a high amount of

297 variation in salmon abundance (marginal  $R^2 = 0.70$ ). Model residuals did not violate  
298 assumptions of homogeneity, normality or independence.

### 299 3.2. Salmon growth

300 The direct positive influence of *Ranunculus* cover on salmon growth was not retained in the  
301 path model. Instead, there were multiple indirect influences of *Ranunculus* cover on growth  
302 via physical habitat and prey resource variables (Figure 4b). *Ranunculus* cover was positively  
303 directly related to prey biomass ( $\gamma = 0.84, p < 0.001$ ) and velocity heterogeneity ( $\gamma = 0.53, p$   
304  $< 0.001$ ). Prey biomass had a strong, positive influence on salmon growth ( $\beta = 0.81, p <$   
305  $0.001$ ). Velocity heterogeneity was negatively related to proportion of fast velocities ( $\beta = -$   
306  $0.59, p = < 0.001$ ), which in turn was strongly, positively related to prey biomass ( $\beta = 0.75, p$   
307  $= < 0.001$ ). This latter relationship between proportion of fast velocities and prey biomass  
308 was identified as an important missing link in the initial model. The influences of prey size,  
309 salmon abundance and trout abundance on salmon growth were not significant, and so these  
310 variables were not retained in the final model. Velocity heterogeneity and proportion of fast  
311 velocities did not directly affect salmon growth, so these pathways were also removed. The  
312 overall effect of *Ranunculus* cover on growth was positive (Table 2b), and the data were  
313 judged to be consistent with the model (Fisher's  $C_{10} = 10.27, p = 0.42$ ). The model explained  
314 a high proportion of the variance in salmon growth in weight (marginal  $R^2 = 0.56$ , conditional  
315  $R^2 = 0.66$ ). Model residuals did not violate assumptions of homogeneity, normality or  
316 independence.

317

### 318 4. Discussion

319 This study uncovered evidence that the positive influences of *Ranunculus* cover on salmon  
320 abundance and growth during summer (Marsh *et al.*, 2021a), are likely driven by a direct

321 effect of cover, and indirect effects operating through multiple pathways, respectively. These  
322 findings provide mechanistic understanding of the key influence that macrophytes have on  
323 salmon communities in lowland rivers. Salmon abundance was strongly influenced by a  
324 direct effect of *Ranunculus* and indirectly, through influences on water depths and velocities.  
325 By contrast, salmon growth appears to not be substantially affected by a direct effect of  
326 *Ranunculus*, but instead predominantly influenced indirectly via impacts of *Ranunculus* on  
327 prey resources and water velocities. Multiple indirect pathways were identified in each  
328 analysis, via physical habitat and food resource variables, highlighting the potential of  
329 *Ranunculus* to influence multiple ecosystem components in lowland rivers.

330 Higher *Ranunculus* cover was directly associated with higher salmon abundance, supporting  
331 the hypothesis that densities of juvenile salmonids are determined by availability of shelter,  
332 not just food resources (Riley *et al.*, 2009). Greater plant cover increases the area of shelter  
333 afforded by both the floating canopy on the water surface, to protect salmon from aerial  
334 predators (Johnsson, Rydeborg & Sundstrom, 2004), and a complex vertical structure  
335 throughout the water column to shield or aid evasion from aquatic predators (Dionne & Folt,  
336 1991; Diehl & Kornijow, 1998). Such structural complexity could result in plant cover being  
337 preferable habitat, attracting more individuals and resulting in reduced mortality rates from  
338 predation. Higher salmon abundance in patches with high *Ranunculus* cover might also result  
339 from visual isolation among competitors, which can reduce aggressive behaviour, and  
340 consequently the territory size defended by an individual (Venter *et al.*, 2008), increasing the  
341 effective carrying capacity of a site (Venter *et al.*, 2008). The presence and use of shelters by  
342 salmon (e.g. boulders, tree roots, aquatic vegetation) can also lead to lower metabolic rates  
343 (Millidine, Armstrong & Metcalfe, 2006), sheltering them from extreme daytime summer  
344 temperatures (Gries & Juanes, 1998) and allowing rest out of fast water flows during night  
345 (Riley *et al.*, 2006).

346 We found evidence of a strong, positive effect of fast velocities on salmon abundance, which  
347 is consistent with suggestions that water velocity is a key determinant of juvenile salmon  
348 habitat selection (Morantz *et al.*, 1987; Armstrong *et al.*, 2003; Girard, Grant &  
349 Steingrimsson, 2004). Body morphology can influence habitat choice by fish (Armstrong *et*  
350 *al.*, 2003), and the large pectoral fins and streamlined body assist juvenile salmon to maintain  
351 position near the riverbed in fast flowing water (Arnold, Webb & Holford, 1991). Fast  
352 velocities might be preferred by individual fish due to the continuous supply of drifting prey.  
353 In the study stream, salmon prey heavily on three taxonomic groups, *Baetis* sp. and *Serratella*  
354 *ignita* (Ephemeroptera), and Simuliidae (Diptera), which are characteristic of drift taxa  
355 (Waters, 1972; Marsh *et al.*, 2020b). Faster water velocity might therefore support a greater  
356 number of salmon by providing an adequate supply of drifting invertebrates, to ensure  
357 sufficient resources for greater number of individuals. *Ranunculus* cover mediates this effect  
358 by its strong influence on velocity heterogeneity, that is, the steepness of adjacent velocity  
359 gradients in and around plant stands (Wharton *et al.*, 2006). *Ranunculus* cover thus can be  
360 particularly beneficial in habitats with fast water velocities as it offers a low velocity refuge.

361 The negative effect of water depth on salmon abundance suggests that shallower habitats are  
362 preferred during the summer feeding period. Ontogenetic shifts in habitat selection occur in  
363 salmonid species, and older and larger fish tend to shift to deeper habitats (Morantz *et al.*,  
364 1987; Armstrong *et al.*, 2003). Subsequently, increased water depths are associated with  
365 greater abundance of older, and potentially more successful, competitors and predators  
366 (Gibson, 1993; Keeley & Grant, 1995). Therefore, juvenile salmon individuals might either  
367 not select for, or suffer higher mortality rates in deeper water, and so are recorded in lower  
368 abundance in these habitats. In contrast, juvenile salmonids in upland rivers avoid shallow  
369 water in favour for intermediate depths (Morantz *et al.*, 1987; Girard, Grant & Steingrimsson,  
370 2004), as depth benefits include increased prey encounter rates for drift-feeding salmonids



371 and protection from aerial predators and high flows (Gregory, 1993; Piccolo *et al.*, 2007).  
372 *Ranunculus* can provide comparable habitat conditions, i.e. greater foraging opportunities and  
373 shelter from high velocities and predators, in lowland rivers, which might diminish the  
374 advantages of occupying deeper water over shallow habitats.

375 Velocity heterogeneity and prey biomass were not identified as important drivers of salmon  
376 abundance, suggesting that shelter has a greater influence than access to prey resources. The  
377 strong relationships between *Ranunculus* and velocity heterogeneity and water depth  
378 illustrate how macrophytes can influence the physical environment, predominately through  
379 altering water flow dynamics. *Ranunculus* had little influence on basal resources, contrasting  
380 with existing knowledge on the ability of *Ranunculus* stands to retain substantial amounts of  
381 organic material (Gregg & Rose, 1982; Cotton *et al.*, 2006). However, this finding should be  
382 considered with caution as it could also indicate a limitation of the benthic sampling  
383 procedure, i.e. not enough samples were collected to capture average levels of organic matter  
384 associated with site-level *Ranunculus* cover accurately, due to patchy distributions (Downes,  
385 Lake & Schreiber, 1993).

386 Salmon growth was not directly affected by *Ranunculus* cover in our models, suggesting that  
387 the previously observed positive influence of cover on salmon growth (Marsh *et al.*, 2021a)  
388 was mediated by the physical habitat and food resources associated with *Ranunculus* cover.  
389 The influence of *Ranunculus* operated through positive impacts on velocity heterogeneity and  
390 prey biomass, the latter of which in turn was related to greater growth. This finding supports  
391 the expectation that higher *Ranunculus* cover allows for greater abundance and biomass of  
392 macroinvertebrate prey. The strong association between *Ranunculus* cover and  
393 macroinvertebrate prey in the growth model was not retained in the abundance model that  
394 was parameterised on data collected later in the year when macroinvertebrate abundance  
395 declines and *Ranunculus* plants begin to die back (Marsh *et al.* 2021a). Macrophytes provide

396 microhabitats that invertebrates can occupy (Hutchens *et al.*, 2004), and plants with complex  
397 architectures are purported to support greater macroinvertebrate biomass due to increased  
398 surface area for colonisation and protection from predators (Heck and Crowder 1991; Diehl  
399 & Kornijow, 1998). Structural complexity from highly dissected *Ranunculus* leaves enhances  
400 ability to trap detritus and support epiphytic biomass (Gregg and Rose 1982), that form food  
401 resources for macroinvertebrates. These increased resources might therefore support greater  
402 numbers of individuals and explain why prey biomass was greater in higher *Ranunculus*  
403 cover.

404 High velocities were expected to be detrimental to salmon growth due to the increased  
405 energetic demands of inhabiting fast flowing water (Wankowski & Thorpe, 1979), but a  
406 direct influence of the proportion of fast velocities was not found to be important. Although  
407 drift feeding in salmonids can be considered as energetically expensive (Puckett & Dill,  
408 1985), behavioural responses to foraging in habitats with increased velocities, such as  
409 reducing attack distances, can negate any additional energy costs (Godin & Rangeley, 1989).  
410 The proportion of fast velocities indirectly influenced salmon growth through a positive  
411 association with prey biomass, a relationship that was not originally hypothesised but  
412 identified from the model fit. This could represent the habitat preferences of  
413 macroinvertebrate taxa that were highly abundant in the Surber samples, particularly  
414 rheophilic fauna, such as Baetidae, Simuliidae, Ephemerellidae, Elmidae and  
415 Hydropsychidae (Marsh *et al.*, 2020b). High velocity conditions can afford benefits through  
416 predator evasion and increasing oxygen supply, particularly for larger macroinvertebrate  
417 individuals with greater resource requirements (Sagnes, Merigoux & Peru, 2008).

418 There was no significant influence of salmon or trout abundance on salmon growth detected,  
419 suggesting that growth (and competition for resources) was not negatively impacted by the  
420 abundance of con- or hetero-specifics in this experiment. This finding is likely context

421 specific, however, as other studies have shown negative effects of competitor densities  
422 (salmon or trout) on juvenile salmon growth, linked to individuals competing for similar and  
423 limited resources (e.g. Bal *et al.*, 2011; Gregory *et al.*, 2017). Additionally, due to the  
424 recruitment crash in the river Frome catchment in 2016 (Marsh *et al.*, 2021b), there might  
425 have been insufficient densities of con- or hetero-specifics to detect any influence of  
426 competition on growth. In line with previous works that observed the territory sizes of  
427 juvenile salmon to be within a few square metres (Armstrong *et al.* 1994; Venter *et al.* 2008),  
428 in this study we assumed that the home ranges of juvenile salmon were mostly limited to the  
429 individual 20 m study sites and thus, exposure to the *Ranunculus* treatments. Extending the  
430 scale of the study, both temporally and spatially, would be required to discern whether the  
431 relationships between *Ranunculus* and salmon abundance and growth sustained across larger  
432 scales and, ultimately, resulted in a change at the population level. This work would be an  
433 interesting future research direction.

434 Direct and indirect influences of *Ranunculus* underlie greater abundance of faster growing  
435 salmon at a reach-scale. Even in these relatively simple path analyses, the important links  
436 retained for the two salmon responses demonstrate the complex interactions, which govern  
437 the impact of *Ranunculus* cover on multiple abiotic and biotic variables. This highlights the  
438 role of *Ranunculus* as a key aquatic structure that mediates environmental conditions in  
439 lowland streams promoting abundance, and growth conditions in individuals of species at  
440 multiple trophic levels by creating suitable physical habitat with abundant resources. The  
441 analytical approach taken in the study allowed us to test for salmon responses to varying level  
442 of macrophyte cover as well as associated changes in habitat heterogeneity, providing new  
443 mechanistic insights into how *Ranunculus* benefits juvenile salmon in lowland rivers. The  
444 impact of macrophytes on habitat and ecosystem productivity could generalise to native  
445 macrophytes and fish communities in river systems where large substrates that provide

446 structural complexity are infrequent. Only by elucidating the mechanisms driving change in  
447 population dynamics, can we design and undertake effective restoration of ecosystems  
448 degraded by anthropogenic activity. This study suggests that strategies to maintain or  
449 enhance naturally occurring *Ranunculus* will likely return wide ranging ecosystem benefits,  
450 including for species of high conservation value, such as salmon.

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456 Author contribution statement

457 Conceptualisation: JM, JIJ, RL, PK. Developing methods; data interpretation; writing: JM, JIJ,  
458 RL, JG, PK. Conducting the research, data analysis, preparation of figures and tables: JM.

459 Conflict of interest statement

460 The authors declare that they have no conflict of interest.

461 Data availability statement

462 The data supporting the original experiment and used in this study are available via the Dryad  
463 Digital Repository <https://doi.org/10.5061/dryad.612jm643t> (Marsh et al., 2021c); additional  
464 data are available from the authors upon reasonable request.

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470

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**Table 1.** Summary of the variables used in path analysis of salmon abundance ( $n = 36$ ) and growth rates ( $n = 71$ ) with mean and standard error (prior to standardising).

<b>Model</b>	<b>Variable (unit)</b>	<b>Type</b>	<b>Error</b>	<b>Mean (<math>\pm</math> SE)</b>
Abundance	Salmon abundance (N)	Response	Negative binomial	14.47 (2.07)
	Proportion of fast velocities	Response/Explanatory	Gaussian	0.62 (0.04)
	Velocity heterogeneity (Index)	Response/Explanatory	Gaussian	0.80 (0.06)
	Water depth (cm)	Response/Explanatory	Gaussian	31.58 (1.82)
	Total biomass of prey (mg)	Response/Explanatory	Gaussian	221.27 (17.31)
	Basal resources (mg)	Response/Explanatory	Gaussian	0.83 (0.07)
	<i>Ranunculus</i> cover (%)	Explanatory	-	34.00 (4.28)
Growth	Salmon growth in weight (g)	Response	Gaussian	4.55 (0.20)
	Average salmon abundance (N)	Response/Explanatory	Negative binomial	33.97 (1.47)
	Average trout abundance (N)	Response/Explanatory	Negative binomial	13.23 (1.06)
	Average proportion of fast velocities	Response/Explanatory	Gaussian	0.72 (0.01)
	Average velocity heterogeneity (Index)	Response/Explanatory	Gaussian	0.95 (0.05)
	Average total biomass of prey (mg)	Response/Explanatory	Gaussian	191.62 (11.73)

Average size of prey (mg)	Response/Explanatory	Gaussian	0.32 (0.01)
Average <i>Ranunculus</i> cover (%)	Explanatory	-	53.50 (3.57)

**Table 2.** Standardised effects of the paths retained in the best fitting model between *Ranunculus* cover (R) and salmon abundance (A) and growth in weight (G), shown in section (a) and (b), respectively. Variables represented as VH – velocity heterogeneity, FV – fast velocity, WD – water depth, P – total biomass of prey. Indirect effects are calculated by multiplying the partial path coefficients and represent the effect of *Ranunculus* on salmon abundance and growth directly and mediated by the physical habitat/food resource variables. The overall effect of *Ranunculus* on salmon abundance and growth is calculated by summing all direct and indirect effects.

<b>Response variable</b>	<b>Pathways</b>	<b>Direct effects</b>	<b>Indirect effects</b>	<b>Overall effect</b>
<b>(a)</b>				<b>0.35</b>
Salmon abundance	R → A	0.52	-	
	R → VH → FV → A	-	-0.05	
	R → WD → A	-	-0.12	
<b>(b)</b>				<b>0.64</b>
Salmon growth in weight	R → P → G	-	0.68	
	R → VH → FV → P → G	-	-0.04	



## Figure Captions

**Figure 1.** Partial effects showing the influence of *Ranunculus* through time on (a) juvenile (0+) salmon abundance during August and September, and on (b) 0+ salmon growth of marked individuals caught in June and recaptured in August. These partial effects were predicted from separate models including other explanatory variables that together described the effects of a *Ranunculus* manipulation experiment on salmon abundance and growth (for full details see Marsh et al. 2021a). The solid line and shaded area represent the mean estimate and 95% confidence interval of the marginal effect of *Ranunculus* cover and the solid circles show the raw data points. These data points form the response variables of this study.

**Figure 2.** Hypothesised relationships of the piecewise SEMs to describe variation in (a) 0+ salmon abundance, and (b) 0+ salmon growth. The path diagrams represent hypothesis of direct and indirect effects of *Ranunculus* cover on salmon abundance and growth,  $\gamma$  represents links between an exogenous and endogenous variable,  $\beta$  represents links between two endogenous variables, numbers correspond to the variable number, e.g.  $x_1$ ,  $y_2$ , that are linked. Black arrows denote positive links and red arrows denote negative links, and the strength of effect is highlighted by arrow thickness.

**Figure 3.** Schematic illustration of the study locations and experimental manipulations: (a) location of the River Frome in Dorset, UK; (b) locations of the three experimental blocks on the North stream (black triangles); and (c) a schematic representation showing that each experimental block consisted of three sites (20 m in length), each manipulated to maintain contrasting *Ranunculus* cover (high / medium / low) and bounded by *Ranunculus* denuded buffer strips (5 m in length), from Marsh et al. 2021a.

**Figure 4.** Outcomes of the final piecewise SEMs testing direct and indirect effects of *Ranunculus* cover on (a) 0+ salmon abundance and (b) 0+ salmon growth. Non-significant pathways were omitted from the final model, and missing paths were identified using d-separation tests and incorporated. Coefficient estimates are shown for each pathway with significance levels are denoted as \*\*\*  $p < 0.001$ , \*\*  $p < 0.01$ , \*  $p < 0.05$ , ·  $p < 0.1$ , marginal  $R^2$  values for each component response variable are shown under the response variable name. Black arrows denote positive links and red arrows denote negative links, and the strength of effect is highlighted by arrow thickness.

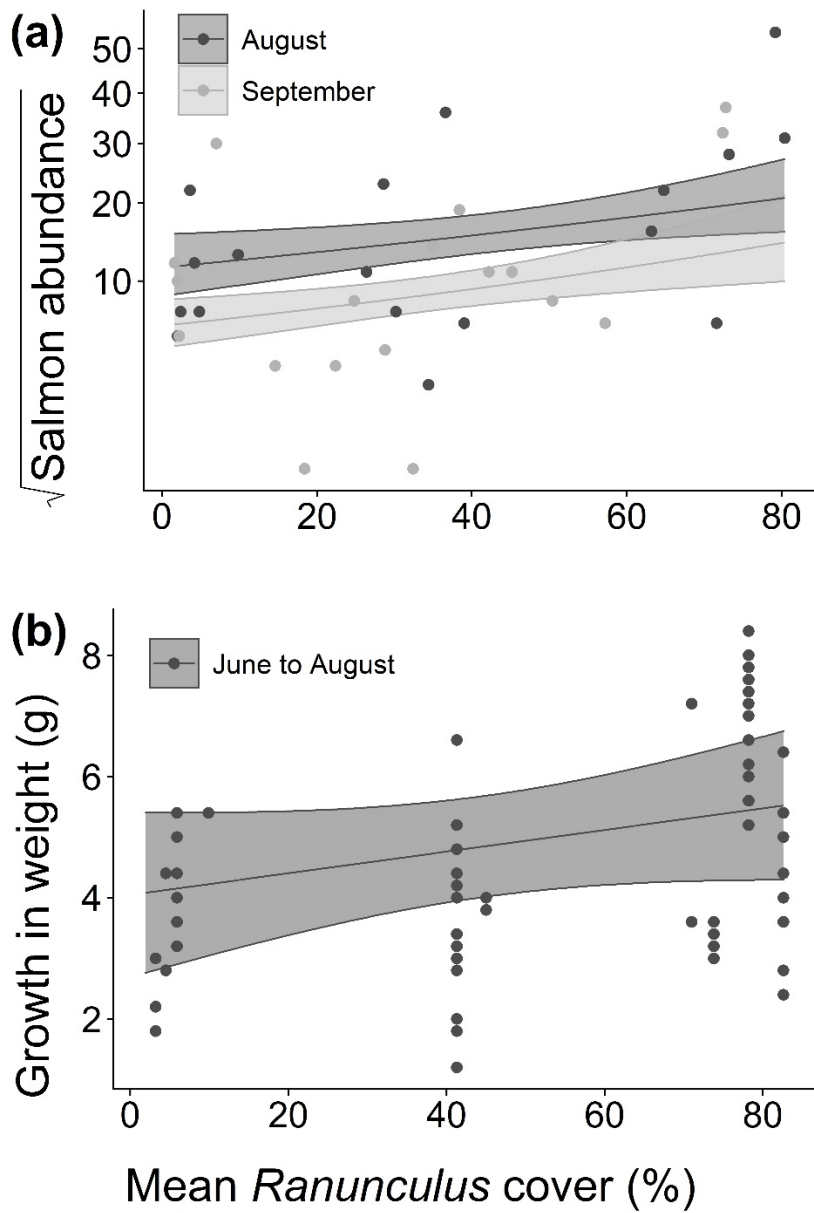


Figure 1.

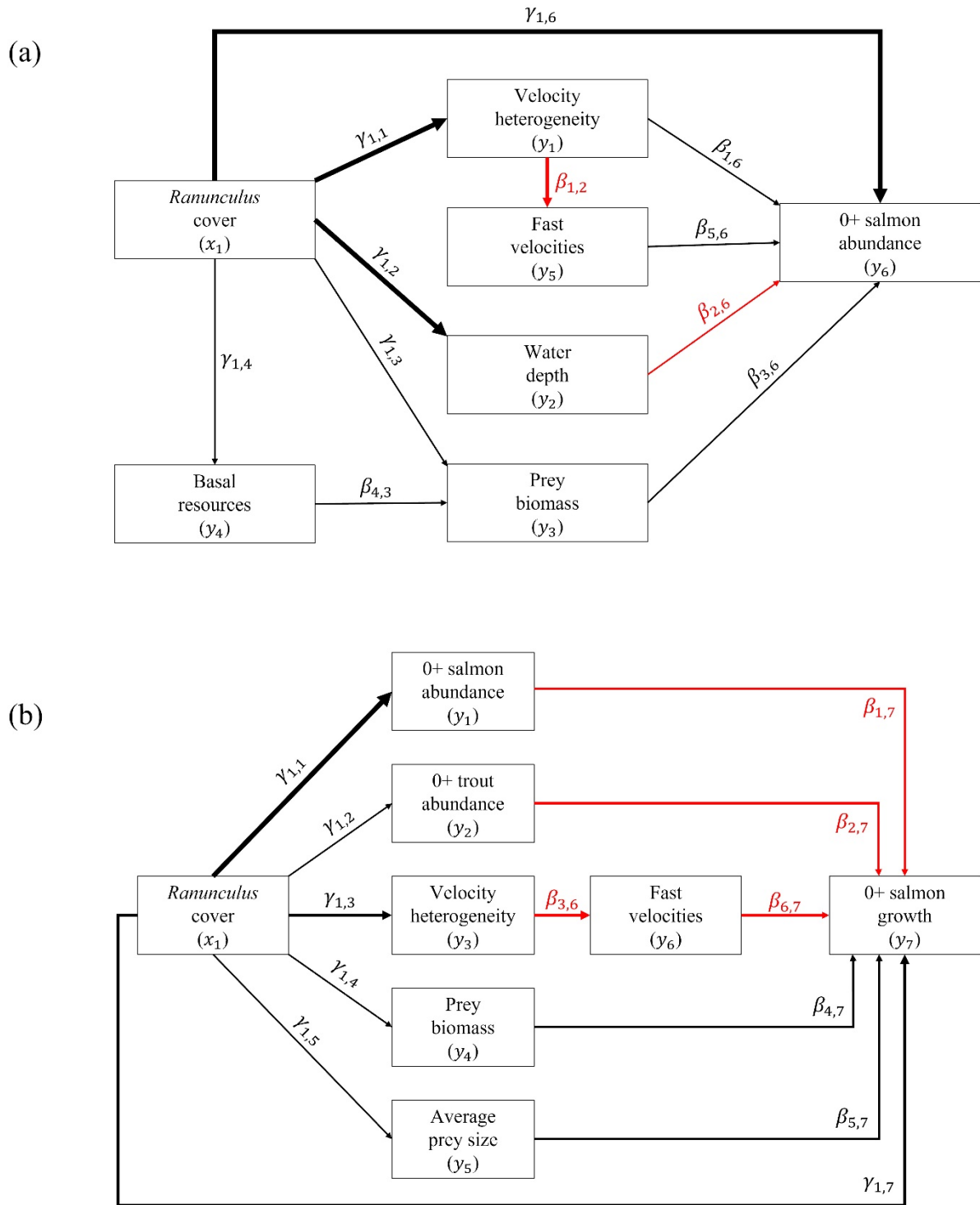


Figure 2.

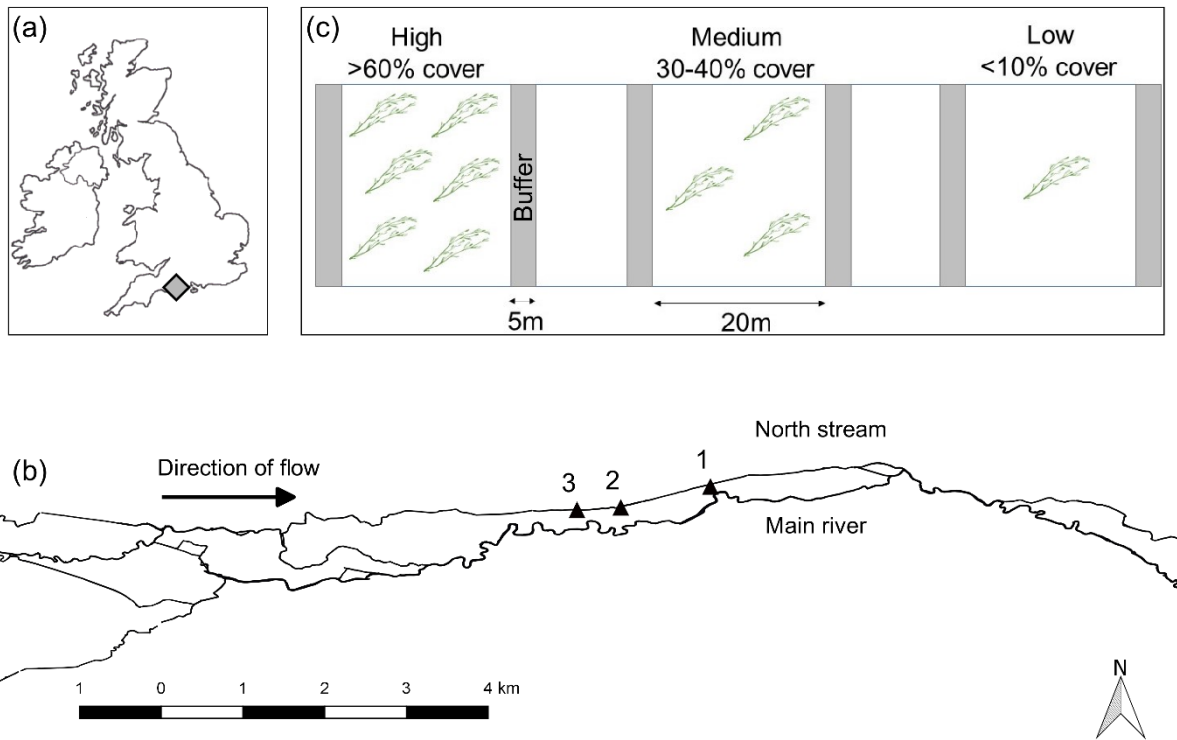


Figure 3.

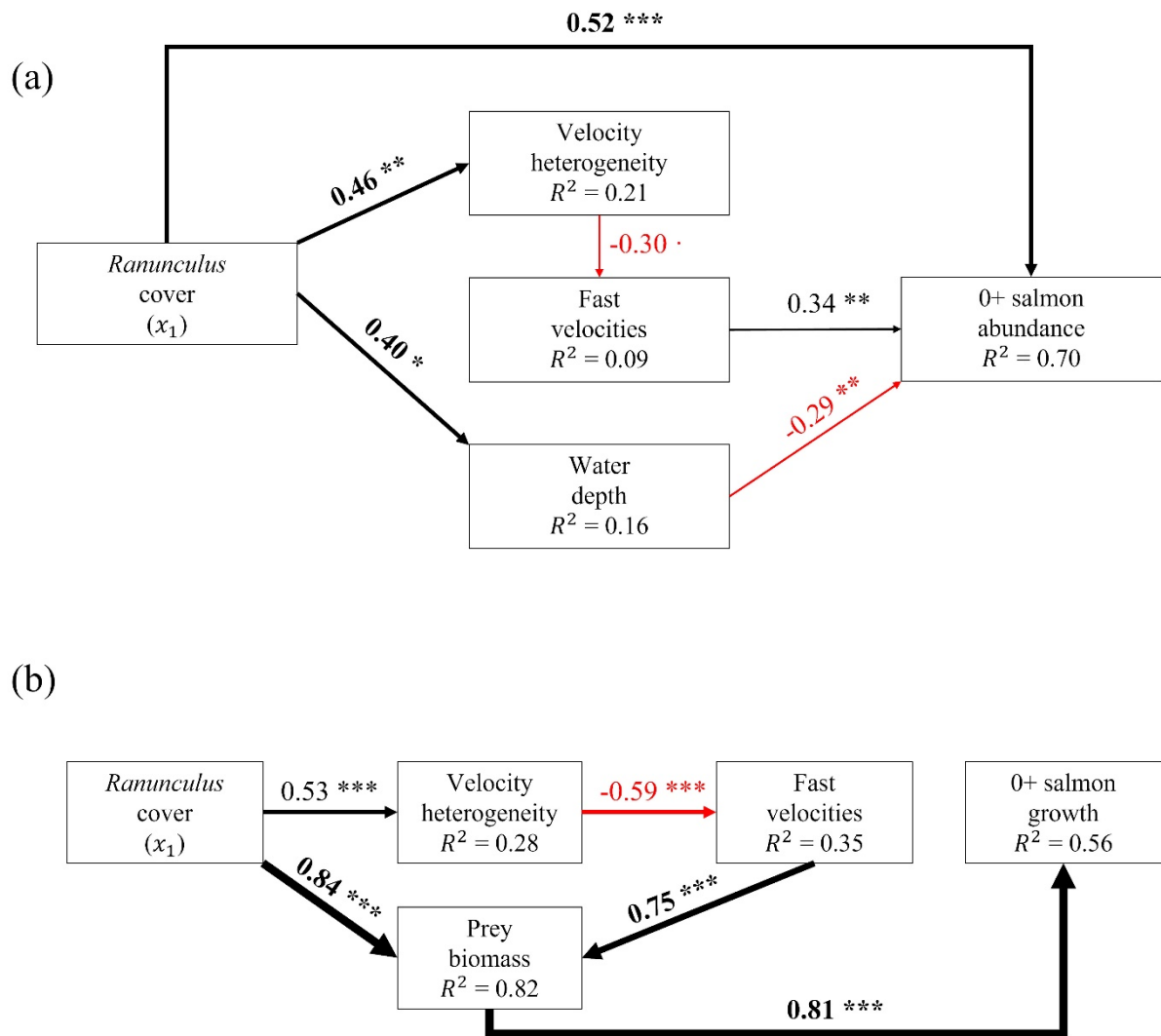


Figure 4.

**Supplementary information in support of: Marsh, J. E., Jones, J. I., Lauridsen, R. B., Grace, J. B & Kratina, P. Direct and indirect influences of macrophyte cover on abundance and growth of juvenile Atlantic salmon. Submission to Freshwater Biology**

**Supplementary Tables**

**Table S1.** Mass-length relationships used to calculate macroinvertebrate biomass. All macroinvertebrate specimens were identified to species, except Oligochaeta (sub-class), Hydrachnidae (family), Simuliidae (family) and Chironomidae (tribe), and counted to determine total abundances per sample. Measures of parasitic Nematoda and Nematomorpha were omitted from these calculations, as these were not considered to represent prey for salmon. We recorded body-size measurements of up to 30 randomly selected specimens of each taxon in each sample and used published length-mass relationships to estimate the mean biomass of each taxon. We then multiplied the mean biomass by the abundance of each taxa to estimate a total biomass of each taxon per sample.

Taxa (Class/Sub-Class)	Reference
Hydrozoa, Insecta, Turbellaria	Benke, A.C., Huryn, A.D., Smock, L.A. & Wallace, J.B. (1999). Length-mass relationships for freshwater macroinvertebrates in North America with particular reference to the southeastern United States. <i>Journal of the North American Benthological Society</i> , 18, 308-343.
Insecta	Burgherr, P. & Meyer, E.I. (1997). Regression analysis of linear body dimensions vs. dry mass in stream macroinvertebrates. <i>Archiv für Hydrobiologie</i> , 139, 101-112.
Copepoda	Dumont, H.J., Van de Velde, I. & Dumont, S. (1975). The dry weight estimate of biomass in a selection of Cladocera, Copepoda and Rotifera from the plankton, periphyton and benthos of continental waters. <i>Oecologia</i> , 19, 75-97.
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Gastropoda	Hawkins, J.W., Lankester, M.W., Lautenschlager, R.A. & Bell, F.W. (1997). Length-biomass and energy relationships of terrestrial gastropods in northern forest ecosystems. <i>Canadian Journal of Zoology</i> , 75, 501-505.
Branchiopods	Herzig, A. (1974). Some population characteristics of planktonic crustaceans in Neusiedlersee. <i>Oecologia</i> , 15, 127-141.
Insecta	Hildrew, A.G. & Townsend, C.R. (1982). Predators and prey in a patchy environment: A freshwater study. <i>Journal of Animal Ecology</i> , 51, 797-815.
Insecta	Johnston, T.A. & Cunjak, R.A. (2001). Dry mass-length relationships for benthic insects: a review with new data from Catamaran Brook, New Brunswick, Canada. <i>Freshwater Biology</i> , 41, 653-674.
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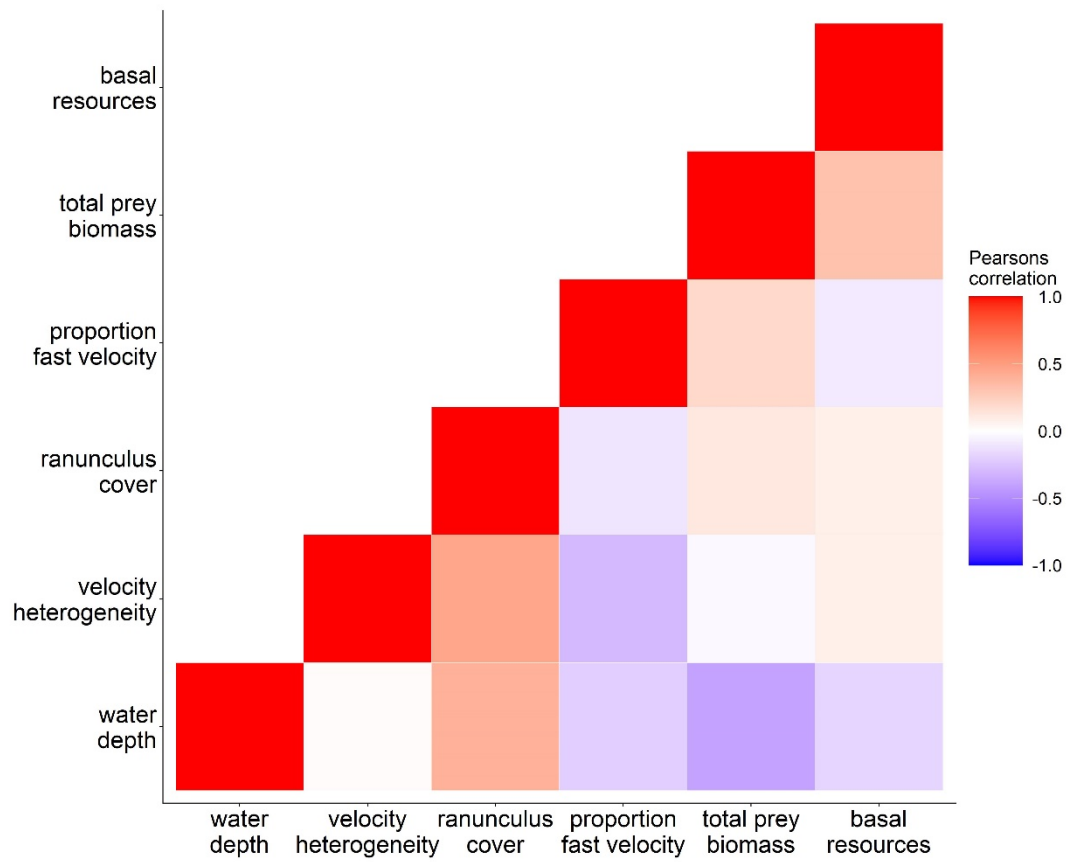
**Table S2.** Model fitting procedure undertaken to identify the final model fit to describe salmon abundance. Non-significant pathways were removed iteratively, and model adequacy was assessed after each removal by calculating Fisher's C test. A – Abundance, R – Ranunculus, FV – Fast velocities, D – water depth, VH – velocity heterogeneity, P – biomass of prey, BR – basal resources.

Model	Model terms	Model term change	Fisher's C	df	P value
1	A ~ R + FV + D + VH + P D ~ R FV ~ R + VH VH ~ R P ~ R + BR BR ~ R	Full model	28.11	18	0.06
2	A ~ R + FV + D + P D ~ R FV ~ R + VH VH ~ R P ~ R + BR BR ~ R	Removed velocity heterogeneity (VH) from abundance (A) model	28.26	20	0.10
3	A ~ R + FV + D + P D ~ R FV ~ VH VH ~ R P ~ R + BR BR ~ R	Removed Ranunculus (R) from fast velocities (FV) model	28.55	22	0.16
4	A ~ R + FV + D + P D ~ R FV ~ VH VH ~ R P ~ R	Removed basal resources (BR) model and BR term in prey (P) model	22.60	14	0.07
5	A ~ R + FV + D D ~ R FV ~ VH VH ~ R	Removed prey (P) model and P term in abundance (A) model	6.97	8	0.54

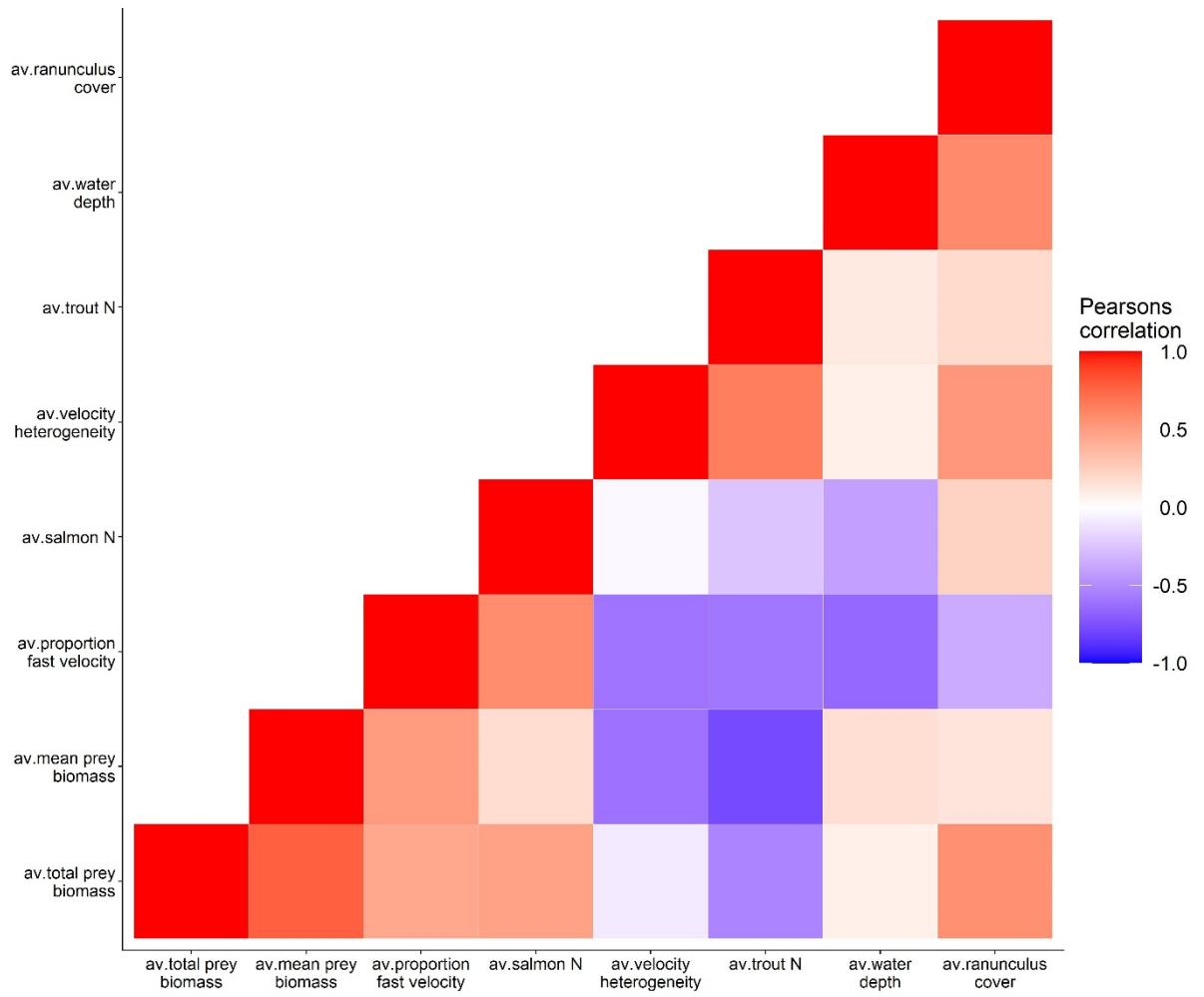
**Table S3.** Model fitting procedure undertaken to identify the final model fit to describe growth in weight of salmon individuals. Non-significant pathways, and important missing pathways were identified using Shipley's d-sep test and were removed/added iteratively. Model adequacy was assessed after each removal/addition by calculating Fisher's C test.

Model	Model terms	Model term change	Fisher's C	df	P value
1	G ~ R + VH + FV + PS + P + SA + TA PS ~ R + P P ~ R FV ~ VH VH ~ R SA ~ R TA ~ R	Full model	558.51	28	0
2	G ~ R + VH + FV + PS + P + TA PS ~ R + P P ~ R FV ~ VH VH ~ R TA ~ R	Removed salmon abundance (SA) model, and SA term in growth (G) model	455.04	18	0
3	G ~ VH + FV + PS + P + TA PS ~ R + P P ~ R FV ~ VH VH ~ R TA ~ R	Removed Ranunculus (R) from growth (G) model	455.01	20	0
4	G ~ VH + FV + PS + P PS ~ R + P P ~ R FV ~ VH VH ~ R	Removed trout abundance (TA) model, and TA term in growth (G) model	254.59	12	0
5	G ~ FV + PS + P PS ~ R + P P ~ R FV ~ VH VH ~ R	Removed velocity heterogeneity (VH) from growth (G) model	256.2	14	0
6	G ~ FV + P P ~ R FV ~ VH VH ~ R	Removed prey size (PS) from growth (G) model	101.19	10	0
7	G ~ P P ~ R FV ~ VH VH ~ R	Removed fast velocity (FV) from growth (G) model	102.40	12	0
8	G ~ P P ~ R + FV FV ~ VH VH ~ R	Included fast velocity (FV) in prey biomass (P) model after identified as an important missing pathway	10.27	10	0.42

Supplementary Figures



**Figure S1.** Correlation matrix for explanatory variables included in salmon abundance SEM.



**Figure S2.** Correlation matrix for explanatory variables included in salmon growth SEM.