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1	Context-dependent reduction in
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3	salmon infested with sea lice
4	
5	Running page head: Sea louse effect on salmon condition
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20 ABSTRACT: Wild Atlantic salmon stocks have globally declined over recent decades. On their migratory 21 return to coastal waters individuals typically are infested by ectoparasitic caligid sea lice 22 (Lepeophtheirus salmonis Krøyer, Caligus elongatus Nordmann). Infestation in laboratory trials can exert non-lethal impacts on the host fish, including increased stress levels and reduced growth, 23 24 physical condition and swimming performance. However, to date no evidence exists for non-lethal 25 effects of sea lice on wild adult Atlantic salmon. Using observations for >6000 return-migrant adult 26 salmon captured from the coastline at Strathy Point (SP), North Scotland, in the estuary of the River 27 North Esk (NE), East Scotland, and the lower reach of the River Tamar (TA), Southwest England, we 28 show that the somatic condition (weight at length) of wild salmon is associated with mobile sea lice 29 density. This putative sea lice-mediated reduction of condition varied with year and seasonal date of 30 freshwater return, and increased with the proportion of adult female parasites on a given fish. 31 Influence of host sex, sea-age and smolt age was negligible. The estimated impact differed between 32 the three sampling sites likely due to underestimation of infestation levels at NE and TA – largely 33 attributable to negative influences of reduced salinity on parasite retention prior sampling. Caligid 34 infestation in the present samples explained a small, but discernible, proportion of the variation in host condition. Reductions in somatic condition of Atlantic salmon are associated with 35 disproportionate declines in accreted lipid reserves. As these are critical to up-river migration and the 36 37 provisioning of eggs, sea lice infestation could have implications for Atlantic salmon population 38 dynamics.

39

INTRODUCTION

42 Parasitism is the most common consumer strategy (Lafferty, Dobson & Kuris 2006), and the effect on 43 a host ranges from commensalism – causing negligible, if any, damage – to inevitable host death 44 (parasitoids) (Anderson & May 1978). The indirect effects of parasites are varied from augmenting the 45 flow of energy in ecosystems, altering the strength of species interactions, changing productivity and 46 causing trophic cascades (Lafferty et al. 2008). In a natural setting, the influence of parasitism on the 47 host is typically confounded with other factors, and may be difficult to identify when populations are at equilibrium (Tompkins et al. 2002). Accordingly, parasite-mediated effects have commonly been 48 49 revealed through anthropogenic perturbations such as introduction of non-native species (Britton 50 2013) and spill-over from livestock (Smith et al. 2009).

Ectoparasitic sea lice (Copepoda; Caligidae) are associated with a wide range of marine fish species.
Their occurrence on wild salmon has long been known (Pontoppidan 1755, Calderwood 1905); our
understanding of the effect of sea lice infestations on wild salmonid stocks remains debated.

54 The salmon louse (Lepeophtheirus salmonis (Krøyer)) is a specialist caligid ectoparasite of salmonids 55 in seawater, and infestations of wild Atlantic salmon Salmo salar returning to Scotland's north coast show a prevalence of 100%; similarly, the prevalence of the host generalist caligid *Caligus elongatus* 56 57 (Nordmann) also typically approaches 100% (Todd et al. 2006). Notwithstanding the fact that all return 58 adult wild salmon in Scotland carry caligid infestations, it remains likely that recent changes in the 59 dynamics of caligid-wild salmonid interactions have occurred as a consequence of climate change, 60 pollution (MacKinnon 1998, Lafferty & Kuris 1999), and perhaps especially the development since the 1970s of intensive cultivation of Atlantic salmon in coastal open net cages (e.g. Ugelvik et al. 2017, 61 Fjørtoft et al. 2017, Thorstad & Finstad 2018, Halttunen et al. 2018). In contrast to the strong 62 63 seasonality of return migration of wild adult Atlantic salmon from the open ocean to freshwater, 64 industrial production of salmon now results in high densities of potential host fish resident in coastal 65 waters throughout the year (Torrissen et al. 2013). Caligid infestations continue to present a major 66 husbandry and fish health challenge to the salmon aquaculture industry (Hall & Murray 2018), and 67 fjords and sea lochs in the vicinity of salmon farms tend to be subject to increased densities of the 68 planktonic and infective sea lice life stages (e.g. McKibben & Hay 2004, Penston et al. 2011, Harte et 69 al. 2017)., This can increase infestation pressure both for local farmed (Jansen et al. 2012) and wild 70 salmonid stocks (e.g. Tully & Nolan 2002). Springtime (April-June) is a season of particular concern to 71 managers of wild stocks of Atlantic salmon and sea trout because this is the out-migration period for 72 the juvenile smolts (Thorstad et al. 2015, 2018). The small size of Atlantic salmon and sea trout smolts 73 (9-19cm) renders these fish vulnerable to physiological stress from sea lice infestation whilst also 74 undergoing osmotic adaptation to seawater (Wells et al. 2006). The sea lice infestation pressure in 75 coastal areas is likely elevated in spring, as a result of the build-up of the run of wild adult salmon 76 returning from the open ocean (Jackson et al. 1997). In addition, the metabolic rate and fecundity of 77 gravid female sea lice both are elevated in springtime due to increasing water temperatures (Johnson 78 & Albright 1991). Wild smolts may therefore encounter relatively high sea lice infestation pressures 79 whilst migrating outward into offshore waters (e.g. Revie et al. 2009, Halttunen et al. 2018).

80 The pathophysiological effects of parasites on the host fish largely depend on host size and 81 immunological status, but also will vary according to the infestation intensity, and size and metabolic 82 demand of the parasite (determined by ambient conditions, parasite species, developmental stage, sex and origin (i.e. farm or wild) (Ugelvik et al. 2017)). In the specific case of caligid copepods infesting 83 84 salmonids, the pre-adult and adult stages (which are mobile across the body surface of the host fish) 85 are more virulent than are the sessile larval (chalimus) stages (Finstad et al. 2000). Laboratory infestation with L. salmonis can elicit non-lethal physiological effects in salmonids which – at high 86 87 intensities – can be lethal. Reported responses range from increased physiological stress (Atlantic 88 salmon: Finstad et al. 2000) to reductions in growth rate, survival and body condition (Arctic charr 89 Salvelinus alpinus: Tveiten et al. 2010). Salmon lice infestation also can decrease host swimming 90 performance (Atlantic salmon: Wagner et al. 2008, Bui et al. 2016) and compromise their foraging 91 ability (sockeye salmon Oncorhynchus nerka: Godwin et al. 2015). Unsurprisingly, somatic growth in wild sockeye salmon smolts is associated with sea louse infestation (Godwin et al. 2017). Other
proximate effects include possible compromise of disease resistance (sea trout *Salmo trutta*: Bjørn &
Finstad 1997, Wells et al. 2007) and elevated vulnerability to predation (pink salmon *Oncorhynchus gorbuscha*: Peacock et al. 2015).

96 Population abundance indices of Scottish and Norwegian salmon stocks have declined particularly in 97 areas with high salmon farm densities (and thus potentially high caligid infestation rates); but such 98 correlative observations do not necessarily prove a cause-effect relationship (Vøllestad et al. 2009). A 99 correlation between stress levels and caligid infestation has been reported for wild sea trout in 100 Norway (Bjørn et al. 2001), and smolt survival rate of sea trout increased twofold for fish medicinally 101 treated against sea lice before release to the natural environment (Skaala et al. 2014). In field trials 102 using cultured Atlantic salmon smolts, survival to return as adult was also overall higher for anti-103 parasite treated fish (Krkošek et al. 2012, Vollset et al. 2015). In this natural setting (as opposed to 104 laboratory), sea lice also had non-lethal effects on the host fish: for example, medicinal treatment of 105 released salmon smolts increased the weight of adults returning to freshwater after one winter at sea 106 (so-called one sea-winter (1SW) adults) by 123 g (95%CI: 45-200 g; Vollset et al. 2015). However, 107 neither mean length nor weight of fish returning after multiple sea-winters (MSW) was affected by 108 treatment. Furthermore, untreated control smolts tended to stay longer at sea and to return as MSW 109 adults, indicating the possibility of a sea lice-mediated delay in maturation (Vollset et al. 2014). In this 110 context, it should not be overlooked that hatchery-reared smolts, like those used in these 111 experiments, typically perform poorly in the natural environment in comparison to wild smolts, because of differences in their feeding behaviour, predator avoidance, migratory behaviour (Jonsson 112 113 et al. 2003, Skilbrei & Wennevik 2006), and possibly also their tolerance of sea lice. Furthermore, these 114 treatment-release-recapture studies are restricted to an assessment of sea lice impacts on smolts 115 during their outward migration, because prophylactic treatment against infestation is effective only 116 for one to two months following application.

117 While mortality has classically been regarded as the primary mechanism behind parasite effects, there 118 is growing consensus that trait-mediated effects on the host (direct) (as well as non-host species 119 (indirect)) are similarly or more important for community structure and function (Ohgushi, Schmitz & 120 Holt 2012). Non-lethal consequences of sea lice infestation including physiological stress, and reduced 121 growth rate and somatic condition (Finstad et al. 2000, Tveiten et al. 2010, Øverli et al. 2014) all can 122 lead to shifts in age at maturation (Vollset et al. 2014), reduced adult fecundity (Burton et al. 2013b), 123 and a potential impairment both of progeny size and early freshwater survival (Todd et al. 2012, 124 Burton et al. 2013a). Here, we aimed to assess possible non-lethal effects that might influence 125 individual host growth, somatic condition, and hence ultimately individual quality and reproductive 126 potential of Atlantic salmon. Morphometric body condition (weight at length) of return adult Atlantic 127 salmon is strongly associated with lipid stores accreted during the marine phase (Todd et al. 2008) and 128 presumably individual survival during river ascent and fecundity. The primary aim of the present study 129 was to assess whether sea lice infestation intensity and parasite life stage composition is associated 130 with somatic condition of adult wild Atlantic salmon that had survived to return to coastal waters.

131

MATERIALS AND METHODS

132 Infestation and host trait data were collated for Atlantic salmon caught in commercial net fisheries at 133 Strathy Point (SP; North Scotland, $58^{\circ}60'$ N, $04^{\circ}00'$ W) and River North Esk (NE; East Scotland, $56^{\circ}75'$ 134 N, $02^{\circ}44'$ W), and from monitoring of a trap weir in the River Tamar (TA; S England, $50^{\circ}52'$ N, $-04^{\circ}21'$) 135 (Anon 2004, Murray & Simpson 2006, Todd et al. 2006). The salinity levels in the three catchments 136 ranged from fully saline (ca. 35 ppt, SP) through brackish (ca. 5 – 25 ppt, NE) to fresh water (ca. 0 – 5 137 ppt, TA). All catchments were remote from aquaculture facilities, with SP being almost 40 km in a 138 straight line over land from the nearest farm. 139 Salmon sampled from the coastal fishery at SP were destined for several rivers across Britain (Shearer 140 1986), and possibly at their first migratory landfall. 491 wild maiden 1SW (no MSW sampled) trapped 141 in bag nets set at SP within approximately 100 m of the shoreline were examined between June and 142 July of each year (1999-2007) (Todd et al. 2006). Sex was ascribed from external observation for 251 143 individuals (2003-2007). No information on smolt age was available for SP. At NE, sea lice data were 144 available for 1176 1SW and 922 MSW fish caught using a sweep net and coble between May and 145 August (2001-2003) (Murray & Simpson 2006). Similar data were collated for 2583 1SW and 1150 146 MSW salmon trapped at the Gunnislake weir (TA) between March and November in 2004-2005, 2008-147 2013 and 2015-2016 (see Table S1 (Supplement) for details). The sampling gear at all three field 148 locations is non-selective; all adult salmon that were intercepted would have been captured.

149 For the NE and TA data no distinction was made between the two endemic parasite species 150 (Lepeophtheirus salmonis and Caligus elongatus) during sampling. Thus, to maintain analytical 151 consistency between all three sites, both species were combined for analysis. During sampling, living 152 (TA) or dead (SP and NE) fish were inspected by eye, and only mobile lice (pre-adult and adult stages) 153 enumerated. Because the impact on host physiology from the smaller (and very rarely observed) 154 chalimus stages is low (Finstad et al. 2000), these developmental stages were ignored. At SP, mobile 155 lice were further classified (to species, sex and developmental stage) using a microscope. This enabled 156 us to determine how the effect of infestation on host condition varied according to the proportion of 157 adult female L. salmonis, which – due to their size and metabolic demand for egg production – is 158 potentially the most virulent stage. The number of sea lice a host can accommodate will depend on its surface area, hence its weight. Thus, assuming that the effect from n mobile sea lice on a host is 159 160 best described as a function of host weight, we used sea lice density D (mobile lice kg⁻¹) as a 161 comparative measure of infestation.

Body condition index K

Atlantic salmon routinely sampled by fisheries research staff were used to determine the length (l) – weight (w) relationship (LWR, regression of log₁₀-transformed w and l). Prior to any analysis, fish missing relevant information (i.e. w or l), farm escapees or repeat spawners (both identified by scale reading) were discarded. One individual with a Fulton's condition factor of 0.1 (indicating measurement error) was also removed. This left 491 1SW from SP (1999-2007), 3680 1SW and 2024 MSW from NE (2001-2003) and 3750 1SW and 1707 MSW from TA (2003-2013, 2015, 2016) (Table S2). LWR coefficients (intercept β_0 , slope β_1) were estimated for each sea age class separately.

171 Each LWR was used to derive the expected standard weight w_s (regression line) at any given l for each 172 sample:

$$\log(w_s) = \beta_0 + \beta_1 \log(l) \tag{1}$$

The deviation of log-transformed w_i in any individual *i* from the regression line (w_s for *i*: $w_{s,i}$) – i.e. the residual of the LWR – was used as body condition index *K* in the present work (Figure S1):

176
$$K_i = \log(w_i) - \log(w_{s,i})$$

177 *K* was the dependent variable in all analyses. It was strongly correlated with other morphometric 178 condition indices including Fulton's index ($R^2 > 0.95$) and relative mass index ($R^2 > 0.98$). *K* was not 179 correlated with *l*, and was assumed to provide a reliable proxy for individual condition (Supplement 180 2).

181

Sea-age determination (NE and TA)

Sea age (*a*) may well influence possible effects of sea lice infestation on host *K* (Vollset et al. 2014). For fish of unknown *a*, sea age was estimated using a two-component (1SW and MSW) Gaussian mixture model (R-package flexmix (Grün & Leisch 2008) v.2.3-13); 2SW and 3SW fish from each site were compiled into a single category (MSW). A comparison between modelled and known (scale reading) *a* values validated the models to a precision of >98 % (NE) and >95% (TA) (Supplement 3).

(2)

Model specification

The sampling location (marine coastal (SP) *versus* estuary (NE) *versus* freshwater (TA)), capture methods (static bag nets (SP) *versus* sweep nets (NE) *versus* the fixed trap (TA)) differed between catchments. This could introduce a catchment-specific bias in the data. In particular, it should be noted that sea lice density (*D*) estimates were not comparable between sites (Figure S9) likely because the capture method and location (salinity) affected how many lice were lost in the process. Thus comparisons between catchments were inappropriate, and each catchment was analysed separately.

194 Throughout the analysis, only predictors with strong biological reasoning (based on *a priori* 195 investigations or literature) were included from the outset to prevent overparameterisation 196 (Burnham & Anderson 2002).

197

Strathy Point

Data for the parasite life stage and species were available for all 491 1SW fish at SP. This enabled assessment for this site of a potential effect of predictors *D*, parasite life stage composition (proportion of female adult *L. salmonis*) θ , *y* and *s* (for 251/491 fish). To assess the influence of θ , we used values representative of our sample: 0.115, 0.4 and 0.745 corresponding to the 2.5th, 50th and 97.5th percentiles of θ observed at SP. Both continuous variables *D* and θ were scaled (sd=1, mean=0) prior analysis. We used a linear mixed effects model with *y* as random variable (random intercepts and slopes of θ (slopes of *D* resulted in model singularity)) to determine the effect on *K* (Table S4).

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North Esk

No 1SW fish were captured in April and for the MSW stock component none were captured in April 207 2001. We therefore combined the months April and May in order to utilize the whole sample. We 208 used a linear model to determine the effects of D and factors y, m and s on K (Table S5).

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Tamar

To avoid erroneous pooling of sampling years, the data were restricted to the main run of each sea age group. These comprised the months March to November (MSW) and June to November (1SW) 212 each year. Fish trapped outside this period (151/2734 1SW and 42/1192 MSW) were discarded. Due 213 to unbalanced data we first introduced a new time parameter, "season" (t), to reduce the temporal 214 resolution for each y: these were, respectively, March to end of May (t = 1, MSW only), June and July (t = 2), August and September (t = 3) and October and November (t = 4). These non-overlapping 215 216 seasonal components were chosen because they incorporate the returning cohort within a particular 217 spawning season, and each component comprised sufficient numbers (at least 30) of 1SW or MSW of 218 each sex (Table S2). Overall, 2583 1SW and 1150 MSW salmon from TA with known D and factors y, 219 t, s were analysed separately. Here we treated y as a random variable (random intercepts only, as 220 random slopes caused model singularity) with t being nested within y (Table S6 & S7).

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Model selection and inference

For each catchment, a full model containing all reasonable predictors (SP: *D*, *y* and θ; NE: *D*, *m*, *y* and *s*; TA: *D*, *t*, *y*, *s* and *a*) and reasonable two-way interaction terms permitted by sample size, was fitted (Table S4-S7). An information theoretic (IT) approach was used to identify the best set of competing models via AICc (dredge in R package MuMIn (Barton 2018)). We considered models with an AICc deviating less than 4 from the lowest AICc as our "top model set" for each catchment. Inferences were made based on weighted support from all top set models by model averaging (Grueber et al. 2011).

Mixed effects linear models (SP and TA) were fitted in R-package lme4 (Bates et al. 2015). The merTools package in R (Knowles & Frederick 2016) was used to calculate 95 % prediction intervals (function predictInterval). Median *K* values and 95 % confidence intervals were obtained by means of a parametric bootstrap (function bootMer; 10,000 iterations), which is the gold-standard approach in mixed models (Knowles & Frederick 2016). Conditional R² (interpreted as variance explained by both fixed and random effects) was determined using the function rsquaredGLMM from R-package MuMIn (Barton 2018).

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Sampling bias and stochasticity

239 Measurements of *l* and *w* in TA were recorded to a precision of 1 mm and 10 g respectively, so that 240 measurement error here was considered negligible. In contrast, the SP and NE measurements of *l* 241 were rounded down at 4 mm precision, and *w* was recorded to a precision of 50 g. Here we 242 bootstrapped our data to appraise how this precision error propagates to influence our results. The 243 following procedure was iterated 1000 times for SP and NE each:

for each of *n* individual fish *i* we randomly drew one *l* and *w* pair from uniform distributions $U(l_i, l_i + 0.4 \text{ cm})$ and $U(w_i - 0.025, w_i + 0.025 \text{ kg})$ respectively. This resulted in a new LWR for each catchment:

$$\log(w_{s_1:n}) = \beta_0 + \beta_1 \log(l_{1:n}) \tag{3}$$

and accordingly new *K* values:

An overview of parameters is given in Table 1.

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$$K_{1:n} = \log(w_{1:n}) - \log(w_{S_{1:n}})$$
(4)

The full model for SP (Table S4) and NE (Table S5) was updated using the new set of *K* values as the response variable and the top set of models was determined (AICc difference <4). Weighted model average (see above) was used to assess the parasite-mediated effect on condition by simulating host condition using two infestation scenarios: (i) actual infestation levels $D_{1:n}$ resulting in condition parameter $K_{p_{1:n}}$, and (ii) zero infestation ($D_{1:n} = 0$) resulting in $K_{0_{1:n}}$.

- Given the 1000 iterations this resulted in 1000 K_p and K_0 values for each individual fish.
- For TA (negligible measurement error), 1000 K_p and K_0 values per fish were simply simulated using the same LWR (and thus same top model set).

For each catchment and individual *i* the sea lice-mediated condition effect, $E_{i_{1:1000}}$, was expressed as the difference between $K_{0i_{1:1000}}$ and $K_{p_{i_{1:1000}}}$. Thus, a positive E_i corresponded to an increase in K_i if infestation was zero. However, because *K* (and accordingly *E*) is merely a dimensionless measure of weight at given length, the effect of the observed infestation levels was demonstrable as the percentage gain (termed G_i) in individual w_i if infestation was zero. Expected weight at zero infestation (w_{0i}) was obtained by 'adding' each predicted change in condition $E_{i_{1:1000}}$ (log₁₀-scale) to the observed $w_{i_{1:1000}}$ (linear scale):

$$w_{0_{i_{1}:1000}} = w_{i_{1}:1000} 10^{E_{i_{1}:1000}}$$
⁽⁵⁾

267 $G_{i_{1:1000}}$ was then given as:

268
$$G_{i_{1:1000}} = 100 \frac{w_{0i_{1:1000}}}{w_{i_{1:1000}}} - 100$$
(6)

For each individual fish the median of $G_{i_{1:1000}}$ (termed M_i) was used as point estimate to quantify the lice-mediated effect (percentage mass gain) on K in each catchment.

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RESULTS

Strathy Point

The influence of host sex *s* (allocated for a subset of n=251/491 fish)) on the parasite-mediated condition-effect (i.e. on the slope) was relatively unimportant: the interaction terms *s*: *D* and *s*: θ were retained only in 5/13 and 4/13 models from the top model set (not shown). Thus the main analysis was undertaken without *s* in order to utilise the entire sample of 491 fish. No information on smolt age of SP fish was available for inclusion in the models. The model set used to predict the condition-effect of infestation *D* in salmon from SP contained three

280 models, all of which retained covariate D (Table 2). These models explained about 45 % variance in

the data (conditional R²). No data points were eminently influential (all Cook's distances below 0.05).

Sea lice density, *D*, was associated with host condition *K*, and its effect increased with the proportion of adult female *Lepeophtheirus salmonis*, θ (Figure 1, Supplement 4). Simulating infestation removal (D = 0) for the SP 1SW sample resulted in a predicted mass increase *M* of median 3.7 % (95%CI: 3.5 - 3.9 %) (Figure 2), corresponding to 85 g (95%CI: 80 – 90 g). The SP top model set explained a large proportion of the variation (45 %) in host condition (Table 2).

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North Esk

The estimated effect of infestation on host condition was not influenced by smolt age r (allocated for a subset of 2054/2098 fish). Thus the analysis was carried out without r using all 2098 fish. Furthermore, the effect of sea age a was relatively unimportant (retained only in 1/5 top set models) and thus ignored during analysis to prevent zero or low sample size in certain month-year combinations. The models explained 9.7 % variance in the data (Table 3). No data points were eminently influential (all Cook's distances below 0.05)

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The determined sea louse effect on condition increased from April/May to August, and over the period 2001 to 2003, and was essentially equal for both sexes (Figure 3). In the NE sample, simulating sea lice 2001 removal resulted in a median mass gain *M* of 0.46 % (95%CI: 0.37 – 0.56 %) for male and 0.33 % (95%CI: 2002 0.25 – 0.42 %) for female fish (Figure 4). This corresponded to an increase in weight of 14 g (95%CI: 12 2003 – 17 g) for male and 9 g (95%CI: 7 – 12 g) for female salmon.

300

Tamar

The model set used to predict the effect on somatic condition of infestation *D* in salmon from TA comprised five and three models for 1SW and MSW respectively, all of which retained *D*, *s* and *t* as covariates (Table 4 & 5). These models explained over 20 % variance in the data (conditional R^2). No data points were markedly influential (all Cook's distances below 0.15). The estimated influence of sea louse infestation on condition was largely unaffected by season *t* or host sex, but tended to be higher for MSW adults (Figure 5). For the sampled TA salmon, our model predicted a mass gain *M* of median 0.19 % (95%CI: 0.16 – 0.21 %) for 1SW and 0.23 % (95%CI: 0.16 – 0.30 %) for MSW salmon (Figure 6) if infestation was removed. This corresponded to an increase in predicted weight of 4.6 g (95%CI: 4.0 – 5.3 g) for 1SW fish and 10.7 g (95%CI: 7.8 – 13.7 g) for MSW salmon.

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DISCUSSION

314 All returning adult 1SW salmon sampled at SP carry mobile sea lice (Todd et al. 2006). This demonstrates the exceptional capacity of these parasites to locate and infest their host. Nonetheless, 315 316 the proximate impact of caligid ectoparasitic load on individual salmon remains the subject of debate. 317 Previous work at SP did not find salmon condition (Fulton's K or relative mass index Wr) to be 318 associated with abundance (sea lice fish⁻¹) of the two parasite species pooled, or of Lepeophtheirus 319 salmonis alone (Todd et al. 2006). Murray & Simpson (2006) previously analysed the River North Esk 320 sea lice infestation patterns for the same data set (NE) as in the present work. They compared the 321 weight-length relationships for fish carrying either zero or >10 sea lice and found no evidence for a 322 significant relationship between sea lice abundance and host condition. However, given a wide host 323 size range (as is the case in our samples), parasite abundance is an inaccurate measure for assessing 324 the potential influence of sea lice on a given individual host.

Here we show that sea lice density, rather than abundance, is associated with the somatic condition of wild Atlantic salmon. However, this effect was context-dependent. The observed relationship could indicate either a sea lice preference for low-condition hosts, an effect on host condition of parasitic 328 sea lice load, a negative relationship between infestation pressure and food abundance at sea, or 329 perhaps a combination of these. There is, however, no evidence to suggest that wild Atlantic salmon 330 of relatively poor somatic condition (i.e. weight at length) are more susceptible to further sea lice 331 infestation, or that infestation pressure decreases with food abundance. Numerous laboratory 332 investigations and field trials using cultured smolts, on the other hand, have shown that sea lice do 333 cause lethal and non-lethal effects on host salmonids (Finstad et al. 2000, Tveiten et al. 2010, Vollset 334 et al. 2015, Godwin et al. 2017). Thus the likelihood is that the presently observed relationships 335 describe a complex sea lice-mediated condition effect, which is dependent largely upon seasonal date 336 and year of freshwater entry, as well as parasite life stage composition. Taking host mass into 337 consideration, we parameterised host infestation in terms of the mobile sea lice density (lice kg⁻¹) 338 rather than numbers of lice per individual fish. This approach provided a means of assessing whether 339 or not the cost of parasite load is host context-dependent. As a result, and from independent data for 340 three sites in the UK, we likely could show an overall detrimental effect of mobile sea lice 341 (Lepeophtheirus salmonis and Caligus elongatus pooled) on host condition in wild Atlantic salmon 342 returning to Scotland and England.

343 The determined effect of infestation D on K was clearer at SP, whereby there was lower variation in 344 K and relatively evenly dispersed values observed for D (Figures 1, 3 and 5). The SP final model 345 explained 45 % variation in the data (Table 2) and predicted a median mass increase of 3.7 % (95%CI: 346 3.5 – 3.9 %) for 1SW fish (Figure 2) if sea lice were removed. This corresponded to a mass gain of 85 g 347 (95%CI: 80 – 90 g), which concurs well with the reported 123 g (45 – 200 g) mass loss in 1SW salmon 348 attributable to sea lice infection during outward migration in Norway (Vollset et al. 2015). As expected, 349 an increasing proportion of the relatively large, and more virulent, adult female *L. salmonis* life stage 350 elevated the impact on condition (Figure 1). One constraint for the SP data is that sex of host fish was available only for a sub-sample of the 1SW adults, and no data were available on smolt age of the 351 sampled fish. Thus, despite the SP observations being more limited, they are important in the present 352 353 context because they are likely to best represent the true infestation levels of caligid ectoparasites on

return-migrant adult Atlantic salmon: the fish were captured in fully marine conditions and remained free-swimming in the capture bag nets. Thus, the inevitable loss of parasites (to unknown extent) as a result of abrasion during capture (e.g. by sweep-netting at NE), or exposure to reduced salinity in the estuary or subsequently in freshwater (i.e. the in-river weir trap at TA) do not extend to the sample data for SP.

359 For NE, only 9.7 % of the variation in host condition was explained by the final model. The predicted 360 median mass increase was only 0.46 % (95%CI: 0.37 – 0.56 %) for male and 0.33 % (95%CI: 0.25 – 0.42 361 %) for female fish (Figure 4), corresponding to increases of 14 g (95%CI: 12 – 17 g) and 9 g (95%CI: 7 – 362 12 g) respectively. This impact increased from 2001 to 2003, and from May to August of each year 363 (Figure 3). Accordingly, only fish returning in July and August tended to be impacted by infestation. 364 Differences in infestation or susceptibility among the differing sea age stock components at NE could 365 explain this variation, or this could reflect the higher summer seawater temperatures and their 366 influences on caligid metabolism (Tully 1992, Heuch et al. 2000). Thus, individuals returning late in the 367 summer season (August) may have been exposed for longer to conditions of increased parasite 368 mobility and virulence, and this may be manifest as an increased impact on host somatic condition.

369

The final models for TA explained 23 and 21 % of data variation (conditional R² considering both fixed and random effects (Nakagawa & Schielzeth 2013)) for 1SW and MSW adults respectively (Table 4 & 5). The predicted median mass gain was 0.19 % (95%CI: 0.16 – 0.21 %) for 1SW and 0.23 % (95%CI: 0.16 – 0.30 %) for MSW salmon (Figure 6), corresponding to increases of 4.6 g (95%CI: 4.0 – 5.3 g) and 10.7 g (95%CI: 7.8 – 13.7 g) respectively. By contrast to NE, the parasite-mediated effect on somatic condition (i.e. slope) tended to be higher in MSW and unaffected by season *t* (Figure 5).

376

In the present context, direct comparisons between the three sampling sites are difficult to draw
because of the different capture methods (bag net, SP; beach seine net, NE; fixed weir trap, TA) and

379 sampling environments (marine, SP; brackish, NE; freshwater, TA). Indeed overall observed sea lice 380 density differed between the three sites accordingly (Figure S9). Furthermore, the influence of sea lice 381 impacts on the host fish will quite probably depend upon other interdependent factors prevailing at 382 sea (e.g. marine prey and predator abundance, which will vary spatially and temporally) (Vollset et al. 383 2015) which may therefore differentially impact upon populations and sea-age groups. However, our 384 multi-site analysis does provide an insight into the relative dynamics of sea louse effects on somatic 385 condition of individual salmon. Whilst the results obtained for NE and TA salmon should be interpreted 386 tentatively, they can be viewed as complementing the analysis of SP 1SW salmon by providing insight 387 in the potential influence of additional host parameters including the seasonal timing of migratory 388 return, sea age, smolt age and sex.

389

390 The overall impact of sea lice on host condition varied with time (month, season or year) and parasite 391 life stage composition (θ), but was independent of host sea age or sex. Smolt age at river emigration 392 was not associated with condition. None the less, the described impact engendered by mobile sea lice 393 stages on wild salmonid condition is in accord with previous laboratory findings (e.g. Arctic charr: 394 Tveiten et al. 2010). Even though the modelled lice density effect showed similarity across all three 395 catchments, there were important discrepancies in the variance explained by each effect model (45% 396 (SP), 9.7% (NE) and >20% (TA)) in addition to differences in the predicted mass gain for the sampled 397 fish in the absence of caligids (median of 3.7% (SP) vs <1% (NE or TA)). Such discrepancies are likely 398 attributable to the differing infestation levels recorded between sites (median D of 13.9 (SP), 1.4 (NE) 399 and 0.4 (TA)) (Figure S9), which themselves are most likely accountable by the sampling locale (marine, 400 estuarine, in-river) and differences in capture method (bag net, seine net, weir) for the three sites. 401 The severity of the introduced bias in observed parasite levels and its effect on the analysis are 402 unknown.

403 We would emphasize that the presently determined putative somatic condition effect on returning 404 adults is likely an underestimate for the population throughout the marine phase, and perhaps 405 especially so for the early post-smolt stage, which is a critical period of high marine mortality in the 406 salmon life cycle (Thorstad et al. 2012). It has to be stressed that, by definition, sampling of only the 407 successfully returning adults precludes investigation of the large majority of each emigrant smolt 408 cohort. The condition of returning adults is not necessarily representative of the entire population, 409 because no comparative information is available for non-survivors earlier in the post-smolt phase. A 410 further concern is posed by a possible bias caused by condition-dependent survival of salmon during 411 the marine phase: highly infested hosts in relatively low or poor somatic condition may be more likely 412 to die at sea and therefore effectively become unavailable for sampling. Thus, over successive years 413 one might record an apparent increase in condition of return adults within a given population, simply 414 reflecting an increasing proportion of (poor condition) individuals within the population dying 415 unobserved at an earlier stage in the marine phase. In addition, it will be important for future studies 416 to determine whether the reported contrasting sea lice density effect on condition as salmon travel 417 north to south (e.g. SP to NE and TA) is in fact masking a survival effect; salmon with lowered condition 418 being more likely to die along the coastline journey.

419 Irrespective of sea lice-induced mortality of post-smolt Atlantic salmon prior to their migratory return 420 to coastal waters (and therefore their availability to be sampled), the magnitude of the observed 421 parasite effect on host condition that we report for SP does present some cause for concern in the 422 management of wild salmon populations (Susdorf et al. 2018). The 3.7% reduction in adult body mass attributable to sea lice that we found for SP might appear superficially to be rather trivial, but this 423 424 should be more fully considered in relation to the corresponding and disproportionate reduction in 425 the accreted lipid reserves of individual adults. Previously, Todd et al. (2008) showed marked declines 426 in somatic condition factor for salmon captured at SP and NE between 1997 and 2006. Their conclusion was that the primary driver of the observed time-series decline in somatic condition 427 428 (weight at length) of 1SW salmon was climate-associated changes in oceanic prey availability. In the

429 present context, however, it is notable that they also found a markedly sigmoid relationship between 430 individual condition factor and lipid reserves. The poorest condition adults they observed were ~30% 431 under-weight for their length and yet their lipid reserves were reduced by as much as ~80%. The 432 implications for salmon population regulation and the management of wild stocks are clear, in that 433 such compromise of individuals' lipid reserves will have marked consequences for egg number and 434 quality (lipoprotein provisioning) of females at spawning (Todd et al. 2008; Burton et al. 2013b). 435 Furthermore, it is important to emphasize that the present study areas are relatively remote from 436 salmon aquaculture sites in Scotland: the nearest salmon farm to a sampling site (SP) has a Euclidean 437 distance overland of almost 40 km not accounting for hydrodynamics or seaways distance, which is 438 greater than the distance reported for sea lice transport from salmon farms in a Scottish system 439 (Adams 2012; Salama et al. 2016). Thus, the sea lice-condition effect that we report here might be 440 predicted to be yet higher for wild stocks of out-migrating smolts passing through areas with dense 441 salmon farming activity (McKibben & Hay 2004, Penston et al. 2011, Harte et al. 2017).

442

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Abbreviation	Parameter
147	Whole fish mass in kg
1	Fork length in cm
<u></u>	Sev of fish
3	Standard or expected fish mass in kg at any l given $l \sim w$ relationship (LWP)
W _S	Body condition index applied in analysis (residuals from LWR)
ĸ	Soo are (15W or MSW): number of years spont at soo
u r	See age (15W of W5W). Humber of years spent at sea
l d	Smolt age. Humber of years spent as juvenile in neshwater
<u> </u>	
t	Season parameter (IA only) covering the salmon run (March-November): March- May ($t = 1$, MSW only), June-July ($t = 2$), August-September ($t = 3$), October- November ($t = 4$)
т	Month (NE only): April-May (4/5), June (6), July (7) and August (8)
у	Year
D	Individual infestation density, i.e. mobile sea lice kg ⁻¹ host
θ	Proportion of female adult <i>L. salmonis</i> in relation to total abundance of mobile sea lice on a host (SP only)
β _o	Intercept of log-log transformed LWR: -5.02 (SP 1SW); -5.36 & -5.48 (NE 1SW & MSW resp); -4.47 & -4.69 (TA 1SW & MSW resp)
β1	Slope of log-log transformed LWR: 3.02 (SP 1SW); 3.23 & 3.28 (NE 1SW & MSW resp); 2.72 & 2.85 (TA 1SW & MSW resp).
М	Effect of sea lice on individual w in percent

625 Table 1: Abbreviation and explanation of parameters used in the present work

629	Table 2: Top model set (Δ AlCc<4, n=3) with potential fixed (D , θ and D : θ) and random (y) effects used
630	to predict the influence of sea lice density D on condition in 1SW Atlantic salmon from Strathy Point

631 (SP).

Intercept	D	θ	<i>D:</i> Ө	df	logLik	AICc	delta	weight	R ²
0.003776	-0.011160	0.002961	-0.003436	8	912.96	-1809.6	0	0.5	0.459
0.003942	-0.009786			6	910.41	-1808.7	0.98	0.31	0.446
0.003890	-0.009745	0.003236		7	910.98	-1807.7	1.90	0.19	0.455

636Table 3: Top model set (Δ AlCc<4, n=2) with potential covariates (*D*, *m*, *s*, *y*, *D*:*m*, *D*:*s*, *D*:*y* and *m*:*y*) used637to predict the influence of sea lice density *D* on condition in Atlantic salmon from River North Esk (NE).

Intercept	D	т	S	у	D:m	D:s	D:y	m:y	df	logLik	AICc	delta	weight	R ²
-0.00834	0.00166	+	+	+	+		+	+	20	4066.4	-8092.5	0	0.714	0.097
-0.00794	0.00148	+	+	+	+	+	+	+	21	4066.5	-8090.7	1.83	0.286	0.097

- 639
- 640
- 641
- Table 4: Top model set (ΔAICc<4, n=5) with potential fixed (*D*, *s*, *t*, *D*:*s*, *D*:*t* and *s*:*t*) and random (*y*)
- 643 effects used to predict the influence of sea lice density *D* on condition in 1SW Atlantic salmon from 644 River Tamar (TA).

Intercept	D	s	t	D:s	D:t	s:t	df	logLik	AICc	delta	weight	R ²
-0.00407	-0.00128	+	+				8	4735.103	-9454.1	0	0.451	0.227
-0.00102	-0.00127	+	+			+	10	4736.43	-9452.8	1.37	0.227	0.228
-0.00413	-0.00124	+	+	+			9	4735.107	-9452.1	2	0.166	0.227
-0.00103	-0.00127	+	+	+		+	11	4736.431	-9450.8	3.39	0.083	0.228
-0.0035	-0.00165	+	+		+		10	4735.298	-9450.5	3.64	0.073	0.227

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Table 5: Top model set (Δ AICc<4, n=3) with potential fixed (*D*, *s*, *t*, *D*:*s*, *D*:*t* and *s*:*t*) and random (*y*) effects used to predict the influence of sea lice density *D* on condition in MSW Atlantic salmon from River Tamar (TA).

Intercept	D	s	t	D:s	D:t	s:t	df	logLik	AICc	delta	weight	R ²
0.01361	-0.00579	+	+		+		12	2194.057	-4363.8	0	0.607	0.209
0.01362	-0.00581	+	+	+	+		13	2194.057	-4361.8	2.05	0.218	0.208
0.01119	-0.00316	+	+				9	2189.753	-4361.3	2.49	0.175	0.211





Figure 1: Relationship between individual condition index K (y-axis) and infestation density D (x-axis) for Strathy Point 1SW salmon given three representative values (0.115, 0.4 and 0.745) of female adult *L* salmonis proportion θ . Predicted K is depicted as solid black regression lines with 95%CI (purple area), and 95 % prediction intervals (blue area).



Figure 2: Predicted increase in whole mass in percent *M* for sampled 1SW at Strathy Point if sea lice infestation is analytically removed (i.e. D = 0). The boxes cover the inter-quartile range (IQR) (25th to 75th percentile), and the whiskers extend to 1.5*IQR below or above the 25th or 75th percentile respectively (default in R).



Figure 3: Relationship between condition K (y-axis) and lice density D (x-axis) for each sex, month, and year-class for North Esk male (left) and female (right) salmon, with predicted K (black lines),
95%CI (purple area), and 95 % prediction intervals (blue area). Details as for Figure 1.



677 Figure 4: Predicted mass gain percentage 678 (M) for male (top) and female (bottom) salmon sampled at North Esk in April/May-679 August 2001-2003 if sea lice infestation is analytically removed (i.e. D = 0). Box and 680 whisker range as for Figure 2.

Tamar



Figure 5: Relationship between condition factor K (y-axis) and lice density D (x-axis) for male (top) and female (bottom) 1SW (left) and MSW (right) Tamar salmon in relation to season t (months abbreviated to letter). Details as for Figure 1.



Predicted mass gain M

Figure 6: Predicted mass gain percentage (M) for 1SW (left) and MSW (right) male (top) or female (bottom) salmon from River Tamar if sea lice infestation is analytically removed (i.e. D = 0). Each column represents a sampling season t (x-axis). Box and whisker range as for Figure 2.