1	Offspring size effects vary over fine spatio-temporal scales in Atlantic
2	salmon (Salmo salar)
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25 **Abstract:** Classic offspring-size theory predicts that a single level of investment per offspring 26 maximizes parental reproductive success in a given environment. Yet, substantial variation in 27 offspring size is often observed among females within populations. Variation at this scale may 28 occur because spatio-temporal variation in stabilizing selection prevents erosion of genetic 29 variation. We tested if patterns of size-specific offspring survival of Atlantic salmon (Salmo 30 salar) varies across location and season within a short stretch of a natural stream by 31 manipulating the emergence timing of juveniles from 12 families with different mean egg 32 sizes and assessing their performance at two locations. The relationship between egg size and 33 juvenile survival varied temporally and spatially: large eggs were advantageous for early 34 emergers in one location, whereas egg size had no effect in the other. Furthermore, the 35 performance of later emerging juveniles did not depend on egg size in either location, 36 possibly because the early emergers had grown or established territories. Thus, selection on 37 offspring size can be complex and vary across short periods of time and small geographic 38 distances, thereby preventing the erosion of genetic variation expected under consistent 39 stabilising selection. 40 41 42 43 44 Key words: breeding timing, density-dependent selection, maternal effects, natural selection

46 Introduction

47 Parental fecundity and investment per offspring tend to be positively correlated with parental 48 fitness. However, these traits trade off, such that parents experience stabilizing selection on 49 investment per offspring. Smith and Fretwell (1974) proposed that, in a given environment, one level of per offspring investment will optimize the trade-off between size and number of 50 51 offspring, thereby maximizing parental reproductive success. Yet, despite this prediction, a 52 common observation is that offspring size varies among parents in a population. Identifying 53 mechanisms responsible for maintaining this variation has received much attention in 54 evolutionary biology. Theoretical models predicting such variation are primarily based on the 55 assumption that different mothers provide different environmental conditions for their 56 offspring, and that this influences size optima (Parker and Begon 1986; McGinley et al. 1987; 57 Hendry et al. 2001). Indeed, large offspring size has been found to be particularly 58 advantageous under adverse conditions (e.g. Fox 2000; Wilson et al. 2009; Monro et al. 2010) 59 or under high population densities (Coltman et al. 1999; Marshall et al. 2006; Svanbäck and 60 Persson 2009). Thus, spatial heterogeneity in environmental conditions within populations is a good candidate mechanism (Levene 1953; Bulmer 1971; Gillespie and Turelli 1989). 61 62 Temporal variation in environmental conditions is also expected to have the potential to 63 conserve within-population variation in fitness-related traits (Ellner and Hairston 1994; Ellner and Sasaki 1996). Even though there is empirical support that both temporal and spatial 64 65 variation in selection pressures can contribute to maintain genetic variation in fitness-related traits (Kittelson and Maron 2001; Siepielski et al. 2009) few studies address how these can 66 67 work in concert to shape selection for offspring size. This is particularly relevant for 68 organisms that live in seasonal environments where important environmental characteristics 69 can vary across small spatial scales as well as within one year. Under such circumstances the

location and timing of reproduction could be expected to interact and shape the pattern ofoffspring size selection.

72 Stream-rearing salmonid fishes are one group of organisms that are well suited for 73 studying spatial and temporal variation in selection on offspring size. First, abiotic and biotic 74 environmental conditions in nursery streams can vary considerably across space within 75 populations (Arnekleiv et al. 2006; Finstad et al. 2009). This includes variation in juvenile 76 densities and competitive intensity over small spatial scales (Einum et al. 2011) caused by 77 patchy egg distribution (Finstad et al. 2010) and limited juvenile movements (Einum et al. 78 2008*a*; Foldvik et al. 2010). Second, the breeding season typically stretches over a long 79 period of time (up to 10 weeks within populations of Atlantic salmon (Salmo salar L., 80 Fleming 1996), which results in considerable variation in timing of emergence from nests 81 (Garcia de Leaniz et al. 2007). Depending on their timing of emergence the environmental 82 conditions experienced by the juveniles can vary over the season (Brännäs 1995). Juvenile 83 density is, for instance, expected to increase across the season as more juveniles hatch and 84 enter the population. Finally, even though the evolution of offspring size has been the subject 85 of much interest in stream-rearing salmonid fishes (e.g. Hutchings 1991; Kinnison et al. 2001; 86 Rollinson and Hutchings 2010), interactive effects of egg size, distribution of breeding and 87 timing of emergence remain unknown.

In a recent field study the performance of Atlantic salmon juveniles from groups varying in emergence timing (experimentally manipulated) differed between locations within the same stream (Skoglund et al. 2011*a*). Spatial variation in intraspecific competition was identified as the most important factor responsible for this pattern (Skoglund et al. 2011*a*). The present study is partly based on the same data as Skoglund et al. (2011*a*), but utilising additional information on the juvenile's family origin (based on microsatellite genotyping), and hence egg size. The combined use of egg size data and experimental manipulation of both

spatial distribution of eggs in the river and emergence timing allowed us to study how
distribution and timing of breeding together shape the patterns of selection on offspring size.

98 Methods

99 Experimental fish

100 Twenty-four hatchery reared Atlantic salmon from the Imsa population (12 females and 12 101 males) with a mean weight of 2400 g were used to create 12 full-sib family groups at the 102 Norwegian Institute for Nature Research (NINA) Research Station, Ims, Norway on 22 103 November 2007. At the same time, samples of the parents' adipose fins were taken and stored 104 in ethanol for later genetic analyses (see electronic supplementary material, Genotyping and 105 parental allocation). Fifteen unfertilized eggs were sampled from each family and fixed in a 106 4% formalin buffer and weighed to the nearest 0.1 mg. Egg mass differed significantly among 107 the families (ANOVA: $F_{12,168} = 5917.8$, P < 0.005, range in mean values: 0.12 - 0.17 g). To 108 create offspring with normal, early and late emergence relative to the wild River Imsa 109 population, the fertilized eggs were divided into three groups, consisting of the same number 110 of eggs from each family, which were incubated at different temperatures. The group with 111 normal emergence timing was incubated in ambient temperature water (average incubation 112 temperature 4.4 °C) drawn from Lake Liavatn (source of the River Imsa), whereas the early 113 and the late emerging groups were incubated in heated (7 $^{\circ}$ C) or cooled water (3.7 $^{\circ}$ C), 114 respectively. According to predictions based on a development model (Crisp 1981, 1988) and 115 daily incubation temperatures, this resulted in median dates of emergence for the early, normal and late groups at 4th April, 7th May and 29th May, respectively. The development 116 117 model has previously been shown to accurately predict dates at emergence across a wide 118 range of temperature regimes for this population (Skoglund et al. 2011b).

119 At the eyed stage the resulting embryos had their otoliths marked with a fluorescent dye by immersion of the eggs for 8 h in a solution of 175 mg L⁻¹ alizarin red S (Baer and 120 121 Rösch 2008). During the immersion procedure a water pump and two air pumps ensured that 122 the water was adequately mixed and aerated. Depending on emergence timing group the 123 alevins received one, two or three fluorescent marks (by repeating the procedure at intervals 124 of 16-23 days). This marking procedure is commonly used to mark large numbers of fish 125 (Wright et al. 2002) and is not known to influence embryonic survival or growth (Baer and 126 Rösch 2008).

127

128 Study site and experimental design

129 The experiment was carried out in the Stream Osalandsbekken (Fig. 1a), a tributary to the 130 River Imsa (the origin of the fish used in this study). This stream has no natural salmon 131 population due to a migration barrier, but has a natural population of resident brown trout (S. 132 trutta, see Einum et al. 2006 for details). Alevins from all timing groups were outplanted 133 across two stream reaches (locations) in well aerated artificial nests (Vibert®, Federation of 134 Fly Fishermen, Bozeman, MT, USA) at different times (early 13 March; normal 22 April; late 135 21 May) but at similar developmental stages. At outplanting the alevins had undergone 90.7 136 % (early), 84.1 % (normal) and 87.6 % (late) of the development from fertilization to emergence (based on development model of Crisp 1981, 1988). To ensure variation in 137 138 environmental conditions between these two locations in addition to the potential effects of 139 non-measured variables (e.g. food availability, habitat suitability) we stocked out 3600 alevins 140 in either of two spatial configurations known to have different effects on competitive regimes, 141 food availability and mortality rates (see Einum & Nislow 2005; Einum et al. 2008a). First, 142 we stocked out 72 individuals (2 individuals \times 12 families \times 3 timing groups) in 25 nests that 143 were spaced 12.5 m apart (Fig 1b). The even distribution of alevins across this relatively long

stream reach should lead to overall low juvenile densities, good growth conditions and high survival. Approximately 750 m downstream from the lowermost nest in the upper location we planted out 1800 alevins (50 individuals × 12 families × 3 timing groups) in a single nest (Fig. 147 1*b*). Due to limited dispersal of salmon fry the high number of alevins located at this location should result in depleted food resources and elevated mortality rates (e.g. Einum et al. 2008*a*). In the following these two reaches will be referred to as the low density and the high density location, respectively.

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152 Sampling and lab analyses

153 During 7-8 July 2008 the stream was divided in 26 sections of about 50 m which were 154 electrofished one to four times depending on the number of salmon caught. Salmon juveniles captured were killed by an overdose of anaesthetics (Clove Oil Extract, $>150 \text{ mg L}^{-1}$) and 155 156 frozen for later processing, whereas brown trout were released back into the stream. In the lab, 157 the salmon juveniles were weighed $(\pm 0.1 \text{ mg})$, sampled for subsequent genetic analyses (see 158 electronic supplementary material, Genotyping and parental allocation), and identified to 159 emergence timing groups. The timing group identity of most fish from the late timing group 160 (<1.1 g) and the largest fish from the early group (>3.9 g) was obvious from the size 161 distributions. Fish in the size range where there was overlap between the timing groups (549 162 individuals) were identified by otolith analysis. This was done by first extracting otoliths from 163 the fish and mounting them onto slide glasses with a transparent adhesive (Crystalbond, 164 Buehler, Lake Bluff, IL, USA). They were then polished with grit paper and analysed for dye 165 marks with an epifluorescent microscope (Wright et al. 2002).

166

167 Statistics

168 Apparent survival was calculated as the number of captured individuals of each family from 169 each timing group in each of the two experimental locations. Because specific growth rate 170 depends on body size we used the standardized mass-specific growth rate (Ω %) (Ostrovsky 171 1995):

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173
$$\Omega = \frac{M_t^b - M_0^b}{b \times t} \times 100,$$

174

175 where M_t is final weight, M_0 is the estimated initial weight at emergence for each family (see 176 below), b is the allometric mass exponent for the relationship between specific growth rate 177 and body mass (estimated to be 0.31 for Atlantic salmon juveniles, Elliott and Hurley 1997) 178 and t is the time between date of emergence (estimated using Crisp 1981, 1988) and sampling. 179 To calculate mean fry weight at emergence for each family we used mean family egg size 180 (adjusted for 16.4 % increase in mass due to water absorption, Einum and Fleming 2000) and 181 the relationship between egg mass and mass at emergence (mass at emergence [g] = 0.880 +182 $0.881 \times \ln \text{ egg mass}$, Einum 2003).

183 All statistical analyses were conducted in R, v. 2.11.1. (R Development Core Team184 2010).

185 The effect of mean family egg mass on three performance proxies (apparent survival,186 final mass and growth) was tested in separate models.

187 To test for an effect of mean family egg mass on survival we used a generalized linear 188 mixed model (GLMM) with a poisson error structure (using the function *lmer* from the *lme4* 189 package, Bates and Maechler 2010) and family as a random factor (intercept). The initial 190 fixed factors included the main effects of mean family egg mass, timing group, location and 191 all interaction effects. To test the effect of mean family egg mass on final mass we used a 192 linear mixed effects model (using the function *lme* from the *nlme* package, Pinheiro et al.

193 2009) that included family as a random factor (intercept). The fixed factors in the initial 194 model were the main effects of mean family egg mass, timing group, location and their 195 interactions. Both final mass and egg mass were ln-transformed prior to analysis. Variation in 196 growth rate was analysed using a similar model, but where none of the variables were 197 transformed. To account for heterogeneity of residuals among the timing groups in the latter 198 model we applied the function *varIdent* from the *nlme* package. This led to a decrease in AIC 199 $(\Delta AIC = 179.0, \text{ comparison done with models fitted using Restricted Maximum Likelihood}).$ 200 For the three initial models, the fixed effects were assessed using a backwards 201 selection procedure (Zuur et al. 2009). Thus, starting with a global model (fitted with 202 Maximum Likelihood), fixed factors were sequentially removed and the resulting simpler 203 models were compared to preceding models using likelihood ratio tests. This removal of fixed 204 factors was done consecutively until the removal of any further terms resulted in a significant 205 (p < 0.05) decrease in log-likelihoods.

To obtain a general measure of the upper limits of the selective forces in each location (i.e. not depending on our choice of phenotypic character, egg size), we calculated the opportunity for selection (*I*, Arnold and Wade 1984) as:

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$$210 \qquad I = \frac{SD^2}{\bar{X}^2},$$

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where SD^2 is the variance in apparent offspring survival (i.e. number of juveniles recaptured) among families under the different treatments (location and emergence timing) and \bar{X}^2 is the squared mean family survival within each treatment group. Confidence intervals (*CI*, 95%) for the values of opportunity for selection were obtained by using the 2.5% and 97.5%

quantiles from 1000 resamplings of the distribution of family-specific apparent survivals
using the function *bootstrap* (from the package *bootstrap*, Leich 2007).

The statistical significance of variation in the relationship between egg mass and offspring survival among groups is best obtained through the GLMM approach described above. However, to complement these analyses we also present estimates for directional (unstandardized, β , and standardized, β) and non-linear selection gradients (unstandardized, γ , and standardized, γ) relating relative offspring survival to egg size in the different locations and timing groups based on regression coefficients (Lande and Arnold 1983).

224

225 **Results**

226 Of the 715 juvenile Atlantic salmon recaptured, 475 came from the low density and 240 from the high density location ($\chi^2 = 85.03$, p < 0.0001). In the low density location a migration 227 228 barrier prevented upstream migration and all juveniles were captured within 100 m 229 downstream of the lowermost nest site. In the high density location, 74 % of the captured 230 juveniles were found within 150 m above and below the nest site. None were caught further 231 up than 160 m above, or further down than 270 m below, the nest site. It is therefore very 232 unlikely that individuals had moved between the localities. In addition 2435 young of the year 233 and 839 older brown trout were caught. Of the salmon juveniles captured 666 (440 from the 234 low density location and 226 from the high density location) were successfully genotyped and 235 allocated to their respective family (electronic supplementary material, Genotyping and 236 parental allocation). The remaining 7 % of the juveniles that were not allocated had 237 insufficient marker amplification due to low quality DNA in their samples. All seven loci 238 were in Hardy-Weinberg equilibrium.

239

240 Apparent survival

241 We found no significant correlation between mean family egg mass and apparent survival 242 overall (Pearsons rank order correlation, r = 0.28, T = 0.91, d.f. = 10, p = 0.39). Furthermore, 243 the GLMM that best described variation in apparent survival included only the main effect of 244 timing group and location (comparisons of preceding models p > 0.05, removal of further 245 terms p < 0.001, Table 1). According to this model, apparent survival depended only on 246 locality and timing group. Removal of the three-way interaction (between egg mass, location 247 and timing group) from the initial model, however, lead to an almost significant decrease in 248 log-likelihood (p = 0.051), suggesting that survival effects of egg mass varied between the 249 locations and among the timing groups. Furthermore, low numbers of fish were retrieved 250 from both the normal and late timing groups, causing high uncertainty in survival estimates 251 when individuals were divided among their original families. Since the normal and late timing 252 groups both experienced competition with early emerging conspecifics, we merged them into 253 a single group ('normal/late timing', termed NLT). We then performed a second model 254 selection with an initial model identical to the first one but with the In-transformed number of eggs per family at different locations included as an offset variable (accounting for the 255 256 number of eggs in the NLT group being twice that in the early timing group). The resulting 257 best model included all main effects together with the three-way interaction (decrease in log-258 likelihood for excluding any terms, p = 0.02). Thus, the relationship between egg mass and 259 apparent survival depended on location and emergence timing group. According to this model, 260 the slopes of the relationship between egg mass and apparent survival (Fig. 2) did not differ 261 significantly from zero in either the early or the normal/late emergers at the low density 262 location (slope values \pm SE for the different timing groups [early, ET and normal/late, NLT] 263 given on the log scale: $ET = -0.22 \pm 4.65$, p = 0.96; $NLT = 4.29 \pm 3.64$, p = 0.24). At the high 264 density location, the relationship between egg mass and apparent survival was positive in the 265 early timing group, and not significantly different from zero in the normal/late group (ET =

 12.14 ± 6.0 , p = 0.04; NLT = -6.28 ± 5.44 , p = 0.25). Also according to this model, the slopes 266 267 did not differ significantly between the low density (LD) and high density (HD) location 268 within either the early or normal/late timing group (differences between slopes given on the 269 log scale: HD relative to LD for ET = 12.35 ± 7.58 , p = 0.10; HD relative to LD for NLT = -270 10.57 ± 6.55 , p = 0.11). Neither did they differ significantly between the early and normal/late 271 timing group in the low density location (NLT relative to $ET = 4.51 \pm 5.90$, p = 0.44). 272 However, in the high density location, the effect of egg mass was significantly more positive 273 in the early compared to in the normal/late timing group (NLT relative to $ET = -18.41 \pm 8.09$, 274 p = 0.02).

275

276 Final mass and growth

Variation in final body mass was best explained by a linear mixed model including all main effects (egg mass, location and timing of emergence) together with the two-way interaction between location and timing of emergence (comparison of log-likelihoods of preceding models, $p \ge 0.1$, removal of further terms, $p \le 0.045$). According to this model there was an overall positive effect of larger egg mass and early emergence on final mass (Table 2). In addition, the relative final mass of juveniles in the low density and high density location varied slightly depending on timing group (Table 2).

The linear mixed model that best explained variation in growth included only the main effect of emergence group (comparisons of log-likelihoods of preceding models, $p \ge 0.06$; exclusion of further terms, p < 0.0001). Thus, the model selection did not reveal any correlation between growth and egg mass or differences in growth between the locations (Parameter estimates \pm *SE* for the fixed effects were: Intercept [early timing] = 3.12 ± 0.05 , p< 0.001; normal relative to early timing = 1.43, ± 0.05 , p < 0.001; late relative to early timing = 2.08 ± 0.08 , p < 0.001).

291

Opportunity for selection

Overall, the opportunity for selection (*I*) was substantially lower in the low density (0.09, 95 % CI = 0.049, 0.126) than in the high density (0.26, 95 % CI = 0.138, 0.375) location. Calculations of separate values of *I* for the different timing groups show that this difference

297 normal and late emergers than for the early ones in the low density location, the opposite was

was particularly pronounced for the early emergers. Whereas I tended to be higher for the

true for the high density location (Fig. 3).

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300 Selection gradients

301 Directional selection gradients relating relative offspring survival to egg size in the different 302 locations and timing groups were highest in the early timing group at the high density locality 303 (Table 3). This is consistent with the patterns revealed in the GLMM where there was a 304 significant positive effect of large egg size in the early timing group at the high density 305 location. The estimates for non-linear selection gradients were of a positive sign for all groups 306 (Table 3). This suggests disruptive rather than stabilizing selection, which seems unlikely to 307 reflect real biological processes.

308

309 **Discussion**

By manipulating timing of emergence within family groups and assessing family-specific performance at locations within the same stream we were able to reveal spatio-temporal complexity in the selection on offspring size (egg mass) in Atlantic salmon. In a previous analysis we showed how the effect of timing of emergence on the success of the offspring varied among locations (Skoglund et al. 2011*a*). The results presented here provide additional support for the hypothesis that selection pressures can vary over small spatial scales (100s of 316 meters), as well as throughout the season. More specifically, large egg size led to survival 317 advantages for early emergers in one location, whereas no such effect was found in the other 318 location. Furthermore, performance of juveniles that emerged later in the season did not 319 depend on egg size.

320 Even though the underlying causative agents shaping this selection pattern remain 321 unknown in the present study, variation in competitive intensity and differences in body sizes 322 among the different emergence timing groups represent potential candidates. Two findings in 323 this study provide support for this statement. First, the spatial configurations of nests 324 employed in this study have previously been shown to generate differences in competitive 325 regimes and lead to different survival patterns (Einum and Nislow 2005; Skoglund et al. 326 2011a). Indeed, juveniles in the location with a spatial configuration of nests expected to 327 result in high levels of competition (high density) had significantly lower survival than those 328 in the locality where levels of competition were expected to be low (low density). Thus, the 329 advantage of large egg size for early emergers in the high density locality could be due to 330 higher levels of competition, and the lack of an effect in the low density locality due to low 331 levels of competition. This observation is consistent with density-dependent selection theory 332 (reviewed in e.g. Reznick et al. 2002; Einum et al. 2008b) and also with lab experiments on 333 salmonids, showing that the competitive advantages provided by large egg size (Hutchings 334 1991) are only important in the presence of competition (Einum and Fleming 1999). 335 Second, the lack of egg size effects among later emergers in the high density location can be 336 attributed to asymmetric competition, where earlier emerging juveniles systematically won 337 contests over later emerging ones irrespective of the relative body size of these later ones. 338 This could be either due to the relatively large body size of early emergers achieved through 339 growth, their prior residency (e.g. Cutts et al. 1999; Kvingedal and Einum 2011) or a 340 combination. A similar effect of older cohorts on offspring size-performance has been found

in the side-blotched lizards *Uta stansburiana* (Svensson and Sinervo 2000). Our results also
show how the opportunity for selection can vary over space and time. Consistent with the
findings for egg size, the opportunity for selection was highest for the early timing group and
at the high density location.

Even though there is considerable residual variation (e.g. Fig. 2) and restricted sample sizes (12 families) in this study, we find it unlikely that the pattern revealed is caused by low statistical power for two reasons. First, the finding that selection is strong at high density in the absence of older competitors, is consistent with previous studies (see above). Second, the lack of significant relationships between egg mass and survival in the other treatments (LD, ET; LD, NLT; HD, NLT) are convincing based on visual inspection of the variation, which does not suggest selection patterns to be obscured by outliers (Fig. 2).

352 According to previous experiments performed in Stream Osalandsbekken, the low 353 apparent survival in the high density location relative to that in the low density location was 354 unlikely to have been caused by habitat differences between the sites. In a previous 355 experiment (Einum and Kvingedal 2011), equal numbers of juveniles (two size classes, mean 356 body mass 0.35 and 2.51 g) were released at nine different sites situated 150 m apart along the 357 stream within the stretch used in the current study. One of these sites was identical to our 358 high-density location, and four others were within the low-density location. The estimated 359 survival rate for the smallest size class, which is most relevant for the present study, at the 360 location corresponding to our high density location was 46 %. At the four sites situated in our 361 low density location the mean \pm SD survival rate was 29 \pm 10 %. Thus, there was no 362 indication that the high density location in the present study had a habitat providing lower 363 survival rate than in the low density location. If anything, the trend was in the opposite 364 direction, suggesting that the habitat in the high density site was better than that in the low 365 density site.

366 Our study is likely to be highly relevant for wild salmonid populations since timing of 367 juvenile emergence varies considerably within populations (Garcia de Leaniz et al. 2007). 368 Furthermore, if the difference in apparent survival between the locations was induced by 369 differences in juvenile densities, among-year variation in adult population size (e.g. Myers et 370 al. 1998; Krkosek et al. 2011) could be predicted to shape selection patterns on a coarser 371 temporal scale. Finally, the females distribute their eggs in patches (Geist and Dauble 1998; 372 Moir et al. 1998) which affects the distribution of the resulting juveniles (Foldvik et al. 2010; 373 Teichert et al. 2011). This may result in heterogeneous densities over small spatial scales (e.g. 374 Einum et al. 2008a). Accordingly, it seems likely that egg size selection patterns can vary 375 both temporally and spatially in natural salmon populations, even within one season. Such 376 variation in selection patterns may, together with other processes such as variation in pre-377 emergence selection (Hendry and Day 2003) and straying among populations, contribute to maintenance of the observed additive genetic variation in salmonid egg size ($h^2 = 0.3 - 0.6$, 378 379 Gall and Huang 1988; Su et al. 1997).

380 Spatio-temporal variation in selection pressures caused by differences in local 381 environmental conditions and presence of older juveniles is presumably to a large extent 382 unpredictable. Under such conditions, bet-hedging (reducing temporal variance in fitness) 383 may be an adaptive offspring provisioning strategy. This may involve the production of fewer 384 and larger offspring (conservative), or variable offspring sizes within clutches (diversified) 385 (Philippi and Seger 1989), with the optimal alternative depending on the shape of the egg 386 size-offspring fitness curve (McGinley et al. 1987). In the most extreme case, where egg size-387 offspring fitness curves are dome shaped, diversified bet-hedging may increase maternal 388 fitness in a variable environment (Marshall et al. 2008). However, a previous study on 389 Atlantic salmon found the relationship between offspring size and fitness to be positive and 390 asymptotic, not dome shaped (Einum and Fleming 2000). Simulations that utilized this

391 positive, asymptotic fitness surface suggest that conservative bet hedging almost always 392 yields the greatest maternal fitness, and that diversifying bet-hedging will be advantageous 393 only in extremely variable environments (Einum and Fleming 2004). In the present study, the 394 linear relationships between offspring size and fitness were either positive or non-existent (e.g. 395 Fig 2). Our estimates of non-linear selection (γ ') were of a positive sign, suggesting 396 diversifying rather than stabilizing selection on offspring size, which is a pattern that seems 397 highly unlikely to reflect real biological processes. Thus, given current estimates of non-398 linearity in selection gradients it is becoming increasingly clear that diversified bet-hedging 399 (i.e. intra-clutch variation in egg size) is unlikely to represent an adaptive strategy for 400 salmonids. If the variation in selection patterns is to some extent predictable and in 401 accordance with the findings in the present study, fish breeding early in the season could be 402 expected to benefit from having larger eggs than the ones breeding later in the season. To our 403 knowledge it is not known whether egg size and breeding time are phenotypically and/or 404 genetically correlated within salmonid populations, and future studies assessing this may yield 405 insights into the possible role of correlational selection on these two traits.

406 We found an overall positive relationship between mean family egg mass and final 407 mass across the emergence timing groups and locations, implying that the size advantages of 408 hatching from a large egg is maintained for at least 78 days after hatching for the early, 46 409 days for the normal and 22 days for the late emerging juveniles. Thus, even though large egg 410 size only provided survival advantages in the early group at one location, it may still have 411 positive fitness effects among later emergers due to survival benefits of a large body size at 412 later stages (e.g. increased over-winter survival, Schultz et al. 1998; Biro et al. 2004, but see 413 Hendry et al. 2003). Growth rate was, however, not influenced by egg size but only by 414 emergence timing.

415 In conclusion, we show that the relationship between egg size and juvenile survival in 416 Atlantic salmon may depend strongly on the joint effect of two other maternal traits: the 417 spatial distribution of eggs and the timing of breeding. Accordingly, the optimal solution to 418 the trade-off between egg size and number may vary spatially over small scales and 419 temporally within one season as well as among seasons. These results highlight the 420 complexities shaping selection in general and selection on offspring size in particular. Such 421 complexities will have pronounced impacts on the evolutionary dynamics of this trait, and 422 may be important in maintaining genetic variation within populations.

423

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Tables

Table 1. Parameter estimates from the generalized linear mixed model (with family as a random intercept) best explaining variation in apparent survival of Atlantic salmon juveniles given at the log scale and as treatment contrasts*.

	Estimate ± SE	Ζ	р	
Intercept (LD, ET)	2.69 ± 0.68	39.43	< 0.001	_
HD	-0.67 ± 0.08	-8.14	< 0.001	
NT	$\textbf{-0.2}\pm0.09$	-2.17	0.03	
LT†	-0.39 ± 0.09	-4.09	< 0.001	

*Abbreviations are given for all parameters: Low density location (LD), High density location (HD), early timing (ET), normal timing (NT) and late timing (LT).

+LT relative to NT: estimated value = -0.2 ± 0.1 , Z = -1.95, p = 0.051.

Table 2. Parameter estimates from the general linear mixed model with family as a random

 effect (intercept) that best explain variation in final body mass (ln-transformed) of Atlantic

 salmon juveniles given as treatment contrasts*.

	Estimate $\pm SE$	Т	Р
Intercept (Egg mass, LD, ET)	2.43 ± 0.35	6.9	< 0.0001
HD	-0.05 ± 0.03	-1.85	0.07
Slope (Egg mass)	0.75 ± 0.19	3.95	0.003
NT	$\textbf{-0.29} \pm 0.02$	-12.07	< 0.0001
LT	-1.13 ± 0.03	-42.31	< 0.0001
LL:NT	$0.1\pm\ 0.04$	2.5	0.01
LL:LT	0.04 ± 0.04	1.0	0.31

* LD is the low density location, HD is the high density location, ET is early timing, NT is normal timing, LT is late timing and egg mass is the ln-transformed mean family egg mass (g).

Table 3. Selection gradients relating the relative proportions of juveniles recaptured to family mean egg mass (g) under the different locations (Low density, LD; High density, HD) and emergence timing groups (Early timing, ET; normal/late timing, NLT). Directional selection gradients (unstandardized, β ; standardized, β `) are estimated from linear regression coefficients, and non-linear (stabilizing) selection gradients (unstandardized, γ ; standardized, γ ') are estimated from regression coefficients of squared deviations from the mean. Standard errors are given in parentheses.

	β	β`	γ	γ`
LD, ET	-0.06 (0.97)	-0.004 (0.06)	11.53 (58.27)	0.01 (0.06)
LD, NLT	0.98 (0.76)	0.07 (0.06)	46.73 (42.91)	0.06 (0.06)
HD, ET	2.02 (1.35)	0.22 (0.14)	114.96 (71.42)	0.21 (0.13)
HD, NLT	-0.66 (0.55)	-0.11 (0.09)	25.26 (32.04)	0.07 (0.09)

Figure legends

Fig. 1. (**a**) Map of the Stream Osalandsbekken and its location within Norway. Open and filled circles represent experimental Atlantic salmon nests in the low and high density reach, respectively. Arrows indicate the direction of water flow. (**b**) Experimental design giving the number of alevins from 12 Atlantic salmon families and three emergence timing groups that were distributed among 25 nests in the low density reach and one nest in the high density reach of Stream Osalandsbekken.

Fig. 2. The observed proportions of recaptured Atlantic salmon juveniles originating from families with different mean sized eggs that were outplanted in the (**a**) low density and (**b**) high density location in Stream Osalandsbekken. Filled and open circle(s) indicate the early and normal/late timing groups respectively. Solid and dashed lines are the estimated relationships (from a poisson generalized linear mixed model where differences in initial numbers are taken into account using *offset*) between juvenile recaptures and mean family egg mass within the early and normal/late timing groups, respectively.

Fig. 3. The opportunity for selection (SD^2/\overline{X}^2) and corresponding 95 % confidence intervals *(CI)* for Atlantic salmon juveniles (12 families) from the early, normal and late timing group recaptured at the low density (open circles) and high density (closed circles) location in Stream Osalandsbekken.









Fig. 3.



Emergence timing group