Environmental change influences the life history of salmon Salmo salar in the North

## Atlantic

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Together, climate and food availability are main drivers of change in the ecology of species. The recent changes in size and age at maturity of Atlantic salmon Salmo salar of the River Imsa appear to be an example of that. Annual mean total length $\left(L_{\mathrm{T}}\right)$ of wild one-sea-winter (1SW) S. salar decreased from 63 cm to 54 cm , and there was a corresponding decrease in condition factor (CF) for cohorts migrating to sea from 1976 to 2010. The reduction in $L_{T}$ is associated with a $40 \%$ decline in mean individual mass, from 2 kg to 1.2 kg . Released hatchery fish reared from parental fish of the same population, exhibited similar changes from 1981 onwards. The decrease in $L_{T}$ correlated negatively with near-surface temperatures in the Eastern Norwegian Sea, the main feeding area of the present stock. Furthermore, S. salar exhibited significant variations in the proportion of cohorts attaining maturity after only one winter in the ocean. The proportion of S. salar spawning as 1 SW fish was lower both in the 1970s and after 2000 than in the 1980s and 1990s associated with a gradual decline in postsmolt growth and smaller amounts of reserve energy in the fish. In wild S. salar, there was a positive association between post-smolt growth and the sea survival back to the River Imsa for spawning. In addition, among smolt year-classes there were significant positive correlations between wild and released hatchery $S$. salar in $L_{\mathrm{T}}, C F$ and age at maturity. The present changes may be caused by ecosystem changes following the collapse and rebuilding of the Norwegian spring spawning herring Clupea harengus population, a gradually decrease in zooplankton abundance and climate change with increasing surface temperature in the Norwegian Sea. Thus, the observed variation in life history traits of S. salar appears primarily associated with major changes in the pelagic foodweb in the ocean.

Keywords: climate, condition factor, environmental change, growth, sea survival, temperature.

## INTRODUCTION

Climate is a governing variable of the biosphere, and the ongoing climate change has serious effects on Earth's ecosystems. With a nearly $1^{\circ} \mathrm{C}$ rise in global warming experienced during the past century, significant shifts in distribution, phenology, behaviour and life history of organisms are already evident (Parmesan \& Yohe, 2003). With an expected increase up to $7^{\circ} \mathrm{C}$ by 2100 , the earth's biota will be even more seriously affected (Sheridan \& Bickford, 2011).

Increasing temperature accelerates biochemical and physiological processes of poikilothermic organisms and affects their body size (Clarke, 2003; Pörtner \& Farrell, 2008). The 'temperature - size rule' states that there is tendency for poikilotherms to grow faster, but reach adulthood earlier, at a smaller body size in a warmer climate (Atkinson, 1994; Zou et al., 2012). Consistent with this rule, several recent studies have demonstrated smaller sizes of poikilotherms associated with climate warming (e.g. Daufresne et al., 2009; Ohlberger et al., 2011; Sheridan \& Bickford, 2011). This contrasts the effect when developmental rate is stimulated by increased food quality, where organisms will mature larger, not smaller, for their age (Berrigan \& Charnov, 1994). This differing effect between higher temperature and better food quality on body size, called 'the Berrigan \& Charnov’s life-history puzzle’, has received much attention in ecological literature (e.g. Perrin, 1995; Jonsson et al., 2013, Trip et al., 2014).

Growth rates of fishes fed to satiation increase with temperature up to a maximum point denoted 'the optimal temperature for growth', after which growth-rate decreases gradually with a further temperature increase (Brett, 1956; Elliott, 1994). Ambient
temperatures of fish living in temperate waters are often encountering water colder than their optimal temperature for growth (Forseth et al., 2009). For instance, in the North Atlantic, S. salar typically exploit habitats that are colder than their optimal temperature (Gudjonsson et al., 2015; Minke-Martin et al., 2015), which for post-smolts is about $14^{\circ} \mathrm{C}$ (Handeland et al., 2008). Thus, growth usually increases with increasing ambient temperature given that the fish find enough high quality food. However, if resource availability is poor or temperature unusually high, increased temperature can give lower growth because maintenance costs increase with temperature. Furthermore, at temperatures above the optimum, oxygen content in the water becomes a limiting factor for growth (Pörtner, 2010). For instance, S. salar exhibit negative growth at temperatures above $25^{\circ} \mathrm{C}$ because of this (Jonsson et al., 2001).

Fast-growing individuals typically mature younger than conspecifics that grow more slowly (Alm, 1959), except when growth is exceptionally high. In the latter case, maturity can be delayed (Jonsson et al., 1984, 1999). Early maturation is stimulated if the somatic energy density is high, or impeded if the amount of reserved energy is critically low (Mangel \& Satterthwaite, 2008). In addition, the effect of increased temperature varies depending on the energy assimilation of the fish. If the energy assimilation is high, higher temperature increases, not decreases size at maturity. The synergistic effect of high temperature and energy food quality was demonstrated in experiments with S. salar (Jonsson et al., 2012, 2013). Furthermore, increased growth rate before the onset of maturation, whether this is owing to enhanced lipid content in food or increased water temperature, will decrease age and therefore potentially reduced size at maturity. Thus, within populations of poikilotherms, year-to-year variation in age at maturity can be associated with both ambient temperature and food availability.

The study population, S. salar of the Norwegian River Imsa, typically migrate to sea at a body length between 12 and 30 cm, 1-3 years of age (Jonsson \& Jonsson, 2014b). They are called smolts at the time of seaward migration, and post-smolts during the first summer and autumn at sea. Most smolts enter seawater during spring and early summer to pursue oceanic feeding. Juveniles entering seawater at other times of the year survive poorly (Hansen \& Jonsson, 1989), and when returning to fresh water for spawning, their homing precision is low (Hansen \& Jonsson, 1991). The present fish spend 1-2 (seldom 3) years in the ocean until they attain sexual maturity (Jonsson et al., 1991, 2003). When spawning after 1 winter in the ocean, they are called one-sea-winter (1SW) fish, after two or more winters they are called multi-sea-winter (MSW) fish. They return for spawning during autumn freshets between August and October (Jonsson et al., 2007). The majority ascend the river during a period of 4 to 6 weeks. In a warmer climate, the proportion of MSW S. salar is expected to increase because of poorer growth, or decrease if feeding opportunities are sufficiently good (Jonsson et al., 2013).

In the present study, we assess life history data on S. salar of the River Imsa, monitored from 1976 onwards. Age at maturity is given as proportion of the cohorts that attained maturity as 1SW fish. The mass-length relationship, post-smolt growth and proportion of the seaward migrating fish that return to the home river for spawning were estimated. Based on this, regressions between life history variables and surface temperature in the Eastern Norwegian Sea were tested. Sea surface temperature during the first period in the ocean is important for growth and survival of S. salar (Friedland et al., 2009, 2014). Furthermore, each year from 1981 onwards, groups of S. salar reared to smolts in a hatchery, wereb released in the river. As brood stock, adults returning to the River Imsa were used. The
groups were released to test if environmental conditions influenced wild and released hatchery fish similarly.

Based on life history theory (e.g. Roff, 1992), it may be expected that fish length at maturity, mass-length relationship and survival at sea would increase with near-surface temperature during the first year the fish spent at sea. If so, a higher proportion of the cohorts would attain maturity as 1SW fish. However, a complicating factor is that availability of zooplankton has decreased and competition from planktivorous fish, e.g. the Norwegian spring spawning herring Clupea harengus, have gradually increased gradually during the study period (Beaugrand \& Reid, 2012; Huse et al., 2012; Utne et al., 2012). Thus, growth may have decreased and age at maturity increased instead.

## MATERIAL AND METHODS

## RIVER AND HATCHERY

The River Imsa, South-Western Norway empties into the Boknafjorden near the city of Stavanger (Fig. 1). The river, which is 1 km long, have a restricted spawning area, is characterized by a temperature ranging from 2 to $3^{\circ} \mathrm{C}$ in winter and summer maxima of approximately $20^{\circ} \mathrm{C}$. The water discharge ranges from less than $2-3 \mathrm{~m}^{3} \mathrm{~s}^{-1}$ in summer to autumn maxima of about $10 \mathrm{~m}^{3} \mathrm{~s}^{-1}$ or more (Jonsson et al., 1989).

In the river, a Wolf trap (Wolf, 1951; apertures 10 mm , inclination 1:10) situated 150 m above the river outlet, catches all descending $S$. salar longer than ca .10 cm . All ascending fish were captured in a fixed box trap located besides the Wolf trap on top of a three steps fish ladder (Jonsson \& Jonsson, 2011). The traps were emptied twice every day during the study period.

The River Imsa hatchery uses river water at slightly elevated temperature that ranges from 2 to $8{ }^{\circ} \mathrm{C}$ in winter to approximately $20^{\circ} \mathrm{C}$ in summer. Eggs were incubated in Heath vertical stack incubators with a water flow of $4 \mathrm{~L} \mathrm{~min}^{-1}$ and about 1.5 L of eggs per tray. At the onset of feeding, the alevins were transferred to $1-\mathrm{m}^{2}$ pools with water depth of 20 cm and water flow of $5.5 \mathrm{~L} \mathrm{~min}^{-1}$. The young fish were fed EWOS food pellets every 10 minutes during daylight until they reach 10 cm in length, after which they were fed every 20 minutes during daylight. In early June, the hatchery fish were transferred to $4 \mathrm{~m}^{2}$ pools with a water depth of 50 cm and a flow of about $40 \mathrm{~L} \mathrm{~min}^{-1}$ where they grew for the remainder of the first year. About $50 \%$ or more of the fish smolted 1 year old. The remainder were held at the hatchery in $12 \mathrm{~m}^{2}$ pools with a water flow of $120 \mathrm{~L} \mathrm{~min}^{-1}$ until they smolted 2 year old.

THE FISH

The wild fish originated from adults spawning naturally in the River Imsa during November - December after having completed the entire life cycle in the wild. The embryos incubated through the winter in gravel nests and emerged during the following April. The young (parr) used the river as a nursery before smolting and seaward migration. On average,
the wild fish smolted at 1 (14\%), 2 ( $78 \%$ ) or $\geq 3$ ( $8 \%$ ) years of age and migrated into the ocean where most of their growth occurred (Jonsson et al., 1998; Jonsson \& Jonsson, 2003). From 1976 to 2010, 36201 wild smolts migrated from the river to the sea, and tagging studies indicate that they fed in the North Norwegian Sea north of $62^{\circ} \mathrm{N}$ (Hansen et al., 1993; Jonsson et al., 1993). Of these, $2.9 \%$ returned to the trap as adults.

The hatchery fish were reared to smolts from S. salar captured in the trap of the River Imsa. Usually more than 10 parents of each sex were used. In early May, the smolts were released at the time when the majority of the wild smolts migrated to sea (Jonsson \& Jonsson, 2014b). The release site was located ca. 100 m above the river outlet, just beneath the fish traps in the River Imsa (Jonsson \& Jonsson, 2011). Thus, the hatchery fish completed the remainder of their growth to adulthood naturally. During 1981 to 2010, 182740 1- and 2-year-old hatchery-reared smolts were released. Of these, $1.5 \%$ returned to the trap as adults. All adults had similar genetical background within year, as the parental fish were tagged adults returning to the River Imsa, but they may have differ somewhat across years. However, the river is short and the spawning area very restricted and there are therefore only one spawning population in the river.

The released hatchery fish followed the same migratory route and experienced the same marine resources in the North Atlantic as wild S. salar (Jonsson et al., 1993). On average, approximately $80 \%$ of the adults attained maturity after one winter and two summers in the ocean (1SW), the rest matured as MSW fish, i.e. chiefly two winters at sea (Jonsson et al., 1991). As maturing adults, hatchery and wild S. salar returned to the Norwegian coast
concurrently, but the hatchery fish tended to enter the River Imsa later in the fall than the wild fish (Jonsson et al., 1990).

TREATMENT OF MATERIAL

Before leaving the river, all wild and hatchery S. salar were individually tagged with numbered Carlin tags (Carlin, 1955). Natural tip lengths (L, mm; i.e. total length of fish with naturally distended caudal fin, Ricker (1979)) and body mass ( $W$, g) were recorded. The fish were anaesthetized with chlorobutanol before being treated, and after treatment, they rested until any visual effect of the anaesthetization had disappear. Then, the fish were returned to the river.

## ESTIMATIONS

Post-smolt growth $(G)$ was back-calculated from measurements of the first marine annulus in scales of returning adults caught in the trap ( $L_{\mathrm{T} 1} \mathrm{~mm}$ ) (Jonsson \& Stenseth 1976; Jonsson \& Jonsson, 2011). From this estimated length we subtracted the smolt length (L $\mathrm{L}_{\mathrm{T}}$ $\mathrm{mm})$. In total, 668 and 658 scale samples from wild and hatchery-reared fish, respectively, were used: $G=\ln L_{T 1}-\ln L_{T 0}$.

Fulton's condition factor (CF) of the returning adults was calculated as $C F=100 \cdot W / L^{3}$, where $W$ is mass (g) and $L$ is natural tip length (cm). This mass-length relationship was used because the individuals were recaptured within a short period in the autumn, were in the same maturity stage and originated from the same population (Wootton, 1998). Bacon at al. (2009) evaluated various mass - length relationships and found that for $S$. salar, this simple index gives virtually the same biological interpretation as relationships that are more complex. Furthermore, CF of adult S. salar caught in the box trap in the River Imsa correlated significantly with their somatic energy density ( $E \mathrm{KJ} 100 \mathrm{~g}^{-1}$ wet mass): $E=8.87+553.0 \mathrm{CF}$, $r^{2}=0.44, \mathrm{df}=1,24, P<0.001$. The energy density was measured by summing the energy in proteins, lipids and carbohydrates in the somatic tissue (Craig et al., 1978). Protein was determined by analysis of Kjeldahl. Total lipid content was determined by hydrolysis of a 5-g sample tissue in hydrochloric acid and extraction with diethylether. The carbohydrate content (i.e. the total amount of glucose in the sample) was determined according to Mason (1983). These analyses were carried out in triplicate and mean values used. The caloric coefficients for making energy estimates from proximate composition data were 17 kJ in both protein and carbohydrate and 38 kJ in lipid. For more details, see Jonsson \& Jonsson (2003).

The majority of the fish attained maturity after 1 winter in the ocean (Jonsson et al., 2003), and MSW fish were excluded from the analyses because they were too few for their findings to be representative except for the last few years. In the tests, mean values of the cohorts were used. When appropriate, similar relationships of individual fish within the cohorts were also tested.

Proportion of the tagged S. salar within cohorts that returned to the River Imsa was used as an index of annual sea survival. This was estimated as number of returning adults divided by number of smolts forming the cohort. The figures were not adjusted for any possible mortality effect of tags or tagging, because it was no intention of estimating the absolute survival rate of untagged fish. However, Hansen (1988) found total recapture rates of adult S. salar at $7.7 \%$ for unmarked, $4.1 \%$ for adipose fin-clipped and $3.1 \%$ for Carlin tagged smolts, indicating that the survival rate would be more than two times higher for unmarked than Carlin-tagged smolts.

## SEA SURFACE TEMPERATURE

Near-surface temperatures (0-10m depth) were retrieved from a several decade’s long realization of a hydro-dynamical model resolving the entire Nordic Seas (Lien et al., 2014). A transect consisting of 17 stations (ranging from $62^{\circ} 4^{\prime} \mathrm{N}, 5^{\circ} 2^{\prime} \mathrm{E}$ off the Norwegian west-coast continuing offshore to $64^{\circ} 7^{\prime} \mathrm{N}, 0^{\circ} \mathrm{E}$ ), in the feeding area of the River Imsa S. salar, was applied to represent temperature means in the Norwegian Sea (Fig. 1). This transect was monitored regularly 3-4 times per year by the Norwegian Institute of Marine Research, and the observed temperatures were used to ensure that the model results were unbiased and realistically resolving inter-annual variability. Mean water temperature was estimated for two periods, summer (April through September) and winter (October through March). From April through September is the main growing period of the post-smolts, whereas October through March is a period when growth is low and the fish typically lose weight. In both seasons, sea surface temperature increased significantly during the study period (Fig. 2). Correlations between life history variables and temperature during three months periods were also tested, but these gave no extra information and therefore omitted from the results.

## STATISTSICS

The data were analyzed by use of SPSS version 22. Proportional data were not arcsine square root transformed, contrasting the advised by Sokal \& Rohlf (1981), because this transformation may produces effects that are difficult to interpret [see Warton \& Hui (2011) for a comprehensive argumentation]. However, there was no difference in the general trends and the coefficient of determination of the models were approximately the same, whether or not this transformation was used. Simple linear regression analysis were generally used because the purpose of the study was not to elaborate formal models to explain variability in S. salar life history, but to provide simple associations among vital descriptors and time trends. The linear regressions were tested for deviation from normality by use of the Kolmogorov-Smirnov test. Relationships that were significantly correlated, are given in the results. Differences in slopes of the linear regression lines between hatchery produced and wild S. salar were tested by use of analysis of covariance (Snedecor \& Cochran, 1973). The data series were differenced by one year when comparing life history variables of hatchery and wild cohorts to stabilize the series, because of their monotonic trends. Furthermore, in no case was use of a second independent predictor variable (temperature or life history character) significant in a multiple regression model ( $P>0.05$ ). For the relationship between proportion of 1 SW fish and year of seaward migration, the 4 parameter Weibull equation (Wahed et al., 2009) used, estimated by Python programming (www.Python.org).

## RESULTS

The length ( $L_{T}$ ) of 1SW S. salar of the River Imsa decreased gradually during the study period. This holds for hatchery produced fish released in the river as smolts as well as wild S. salar from the river [Fig. 3(a,b)]. For instance, mean length of wild fish decreased from 63 cm to 54 cm , equivalent to a decrease in mass from approximately 2 kg to 1.2 kg (40\%). The slopes of the lines differed significantly (comparison of slope; $F_{1,59}=61.9, P<$ 0.001). Performing similar analyses for individual fish showed that the variability was large, but the trends were similar and highly significant (Table I).

The condition factor (CF) of both released hatchery and wild S. salar decreased by $8 \%$ [Fig. 3(c,d), comparison of slopes: $F_{1,59}=2.08, P>0.05$ ], whereas the post-smolt growth in length decreased faster for hatchery than wild fish [Fig. 3(e,f); comparison of slopes: $F_{1,44}=$ 13.02, $P<0.001$ ]. Furthermore, $S$. salar exhibited an increase in proportion of 1SW S. salar for cohorts migrating to sea before the mid-1990s and a decrease between 2000 and 2010. Similar trends were observed in both wild and hatchery produced fish [Figure 3(g,h)]. For 1SW fish, there was no significant temporal trend in annual survival at sea, in smolt size or growth during the second summer, i.e. the year they returned for spawning $(P>0.05)$. Age at maturity as illustrated by the proportion of 1SW fish, increased from the 1970s to the 1990s and decreased again after 2000 to a minimum of about 30\% ([Fig. 3(i)]. From the 1980s onwards, the hatchery fish exhibited a similar maturity pattern [Fig. 3(j)]. Similar significant changes in life history traits were observed when individual fish from each cohort was used (Table I).

There were some significant relationships between life history variables and ocean temperatures (Table II), but the coefficient of determination was low exhibiting that only a
small proportion of the total variation was explained by the temperature. $L_{T}$ of 1SW fish decreased with increasing ocean temperature. The slope of the regression was steeper for hatchery than wild S. salar [Table II(a), Summer temperature: $F_{1,57}=8.8, P=0.005$ ]. The similar regression between $L_{\mathrm{T}}$ and winter temperature differed in that wild fish exhibited the steepest slope (Winter temperature: $F_{1,57}=9.1, P=0.004$ ). Furthermore, body mass relative to length (CF) tended to be higher in years when near-surface water in the Norwegian Sea was relatively cold. The slopes of the corresponding regressions did not differ significantly between hatchery and wild fish [Table II (a): Summer temperature, $F_{1,58}=3.10, P>0.05$, Winter temperature $\left.F_{1,59}=3.64, P>0.05\right]$. In addition, a higher proportion of the cohorts of hatchery S. salar attained maturity as 1SW fish when the ocean was relatively cold. This holds for both winter and summer temperature when tested separately. The corresponding relationships for wild $S$. salar, however, were not significant ( $P>0.05$ ). There was no significant correlation between sea survival and temperature in the Norwegian Sea.

In both released hatchery and wild S. salar, mean $L_{T}$ increased with growth during the first summer at sea with a steeper slope in wild than hatchery fish ([Table II(b), $F_{1,44}=7.48, P$ $=0.01]$. Furthermore, the proportion that attained maturity as 1 SW fish increased with $C F$. The slope of the regression was steeper in hatchery than in wild fish ( $F_{1,55}=7.2, P=0.01$ ). In released hatchery, but not in wild fish, the proportion of 1SW fish also increased with growth during the first summer at sea. On the other hand, there was also a positive relationship between the sea survival of the cohort and growth during the first year at sea in wild S. salar, but the corresponding relationship was not significant in hatchery S. salar.

Among years, there was significant positive correlation between the lengths of wild 1SW fish on corresponding cohorts of released hatchery S. salar [Fig. 4(a)]. However, differencing the two time series by one year, gave no significant relationship ( $P>0.05$ ). Furthermore, CF of wild on hatchery S. salar were highly correlated (Fig. 4(b)], but lower for hatchery than wild $S$. salar, particularly in years when the condition factor was low.

Differencing the two time series reduced the coefficient of determination $\left(R^{2}=0.35, P=\right.$ 0.001 ). Also, the proportion of wild on hatchery S. salar returning to the River Imsa as 1 SW fish were highly significant [Fig. 4(c)]. The proportion of 1SW S. salar was higher in wild S. salar in years when relatively few fish attained maturity as 1SW fish, but lower in years when the proportions were high. Differencing the two time series with one year reduced the coefficient of determination $\left(R^{2}=0.40, P=0.002\right)$. There was no significant correlation between estimated growth rate or survival at sea of hatchery and wild River Imsa S. salar ( $P>$ $0.05)$.

## DISCUSSION

The time-series show that $L_{\mathrm{T}}, C F$ and post-smolt growth of $S$. salar have decreased since the 1970s. Similar trends are evident in both wild fish and artificially reared fish released as smolts in the river. The consistency of the patterns indicates that the changes are real, and the decrease in size appears associated with reduced post-smolt growth and not with growth as parr in freshwater or during the second summer at sea. Furthermore, this appears not to be an epigenetic effect of thermal conditions during embryonic development (Finstad \& Jonsson, 2012; review in Jonsson \& Jonsson, 2014a). Jonsson et al. (2014) did not find any effect of increased temperature during early ontogeny on either post-smolt growth or age at
maturity in experimental tests with $S$. salar of the River Imsa. Selection or genetic drift was probably not important for changes in these ecological characters, as the decrease in body mass has been as large as $40 \%$ over only a few generations, and similar trends have been observed elsewhere, as discussed below. Growth and size in fishes are very plastic (cf. Wootton, 1998), and phenotypic responses to environmental change are rapid (Jonsson \& Jonsson, 2011).

Many species exhibit smaller sizes because of climate change (Sheridan \& Bickford, 2011), but the present case may not only be an effect of the rising temperature, as temperature explained little of the variation in these ecological characters. Although significant, the temperature increase has been only approximately $1^{\circ} \mathrm{C}$, which is little, and should not reduce the mass of 1SW salmon by nearly as much as $40 \%$. Instead, reduced feeding can cause such a dramatic decrease, which may be associated with lower primary and secondary production in the pelagic foodweb as argued by Beaugrand \& Reid (2012). During the first year at sea, $S$. salar feed largely on pelagic crustaceans (Amphipoda, Euphausiacea) (Jacobsen \& Hansen, 2001; Haugland et al., 2006). The abundance of these organisms have decreasing concurrently with an increased abundance of competing pelagic fish (Huse et al., 2012), among which the Norwegian spring spawning $C$. harengus may be the keenest post-smolt competitor. This assumption is indicated by negative correlations between $C$. harengus abundance in the Norwegian Sea and S. salar survival (evidence reviewed in Mills, 2003). After the collapse of the Norwegian spring spawning C. harengus in 1970, post-smolt growth was particularly good in the 1970s. But as the Norwegian spring spawning C. harengus population recovered gradually (Huse et al., 2012; Nøttestad et al., 2015), post-smolt growth has decreased as it has in other European rivers (McCarty et al., 2008; Peyronnet et al., 2008; Friedland et al., 2009).

An additional reason for the decrease in growth and $C F$ with time may be the recent increase in abundance of sea lice Lepeophtheirus salmonis Burmeister 1834, which may have affected post-smolt growth negatively (Stien et al., 2005). The increased abundance of $L$. salmonis is probably a consequence of the growth of the S. salar farming industry since the 1970s (Ford \& Myers, 2008). For instance, Gargan et al. (2012) reported that L. salmonis can induce considerable mortality on S. salar. Moreover, Skilbrei et al. (2013) found 6 \% larger mass of 1SW hatchery-produced S. salar treated as smolts either with prophylactic substances emamectin benzoate or EX (Pharmac), against $L$. salmonis prior to release in nature, compared with untreated control groups. A similar effect of $L$. salmonis may have reduced growth of the present $S$. salar. However, the reported effect of the $L$. salmonis treatment was much smaller than the $40 \%$ decrease in mass found in the present study, and no parallel increase in mortality was observed. Thus, L. salmonis may not be a major contributor to the decrease in fish size in the present population.

Todd et al. (2008) reported a similar decrease in CF of 1SW S. salar from Scotland towards the end of the twentieth and beginning of the twenty-first century. They related reduced CF and lipid density of the fish to increased sea surface temperature, associated with a recent climate-driven change in the eastern North Atlantic pelagic ecosystem. Furthermore, they see parallels with this decrease to effects on other top consumers, such as Atlantic bluefin tuna Thunnus thynnus (L. 1758) and seabirds (Todd et al., 2008). The decreased CF of the present $S$. salar indicates a similar reduction in lipid deposits, as the $C F$ correlates significantly with the lipid density of the returning adults in the River Imsa. A similar correlation was found for S. salar spawning in the Norwegian River Drammen (Jonsson et al., 1997). Thus, the climate-driven ecosystem change may have had a similar effect on the
present Norwegian S. salar as reported from Scotland, indicating that this is a general trend in the Northeast Atlantic.

Although reduced energy assimilation of the post-smolts may be the prime reason for the decrease in growth, temperature may still have had a contributing effect because of associated metabolic costs. Friedland \& Todd (2012) reported that S. salar growth was associated with temperature during winter and spring. Furthermore, Hokkaido chum salmon Oncorhynchus keta (Walbaum 1792) have had elevated growth and survival rates since the 1980s (opposite to the present results). They assumed that this was a direct effect of higher surface temperature because of global warming (Kaeriyama et al., 2014). In addition, the growth of Chinook salmon Oncorhynchus tshawytscha (Walbaum 1792) in Alaska appears strongly influenced by oceanic winter temperatures (Wells et al., 2008). Thus, the possibility that increased temperature may have had an additional effect on the ongoing change in size and growth of $S$ salar, cannot be rejected.

Age at maturity varied during the study period. First, the proportion of 1SW fish increased, but after 2000, it decreased. Age at maturity is highly influenced by growth and size of the fish (Alm, 1959; Jonsson et al., 1984) and for S. salar, Friedland \& Haas (1996) indicated that late summer growth of the post-smolts was higher in S. salar maturing as 1SW than 2SW fish. Furthermore, Salminen (1997) reported that good feeding opportunities gave relatively more 1SW S. salar than do poorer feeding opportunities in the Baltic Sea. For released hatchery S. salar, they found a significant relationship between early maturation and post-smolt growth, as also found in rearing experiments with the River Imsa salmon (Jonsson et al., 2012, 2013). But in the present study, the relatinship was nonlinear.

A hypothesis that may explaining this nonlinearity is that the relatively high proportion of MSW fish in the 1970s was related to the eminent S. salar growth during this decade (Jonsson \& Jonsson, 2004). Salmonids can delay maturation if growth is unusually good. Ultimately, this may be an inherited response because of an "expected" increase in reproductive success by postponing maturation (because of the increase in size), more than compensating for the expected decrease in survival, as explained by Jonsson \& Jonsson (1993). For instance, when juvenile brown trout Salmo trutta L. 1758 switch from zoo-benthic to fish feeding, growth rate increases and maturation is delayed relative to those that still feed on zoobenthos (Jonsson et al., 1999). Similarly, very fast-growing S. salar can delay maturation relative to conspecifics with more average growth rate (Jonsson \& Jonsson, 2004). This relationship between maturation and growth seems to hold across populations (Jonsson et al., 1991) and in other salmonid species (Larsson et al., 2005; Forseth et al., 2009). Thus, extremely high post-smolt growth may have contributed to the elevated proportion of MSW fish in the 1970s.

However, maturation can also be delayed if growth rate is poor (Alm, 1959), which may be the situation after 2000. In this period, post-smolt growth was low and a gradually increasing proportion of the cohorts may have lacked the needed energy resources to carry out the return migration and spawn successfully as 1SW fish, relative to the requirements of their genetic programme. Furthermore, maturation can be delayed if lipid densities are low (Rowe \& Thorpe, 1991; Duston \& Saunders, 1999). This holds also for a number of other species, such as S. trutta L. 1758 (Bohlin et al., 1994), O. tshawytscha (Silverstein et al., 1998) and North Sea Plaice Pleuronectes platessa L. 1758 (Grift et al., 2007). Hence, both extra high
and extra low growth in the post-smolt period may be associated with maturation as MSW fish.

Otero et al. (2012) reported a similar decrease in proportion of 1SW S. salar after 2000 in other Norwegian S. salar populations emphasizing the generality of this finding. They suggested that the increased age at maturity may be explained by the large-scale ecological changes in the Northeast Atlantic pelagic food web affecting post-smolt growth, which appears reasonable, as discussed above.

Salmo salar may have moved farther north for feeding during recent years. Jensen et al. (2014) reported that considerable numbers of S. salar were observed as far north as the Svalbard Islands, between $78^{\circ}$ and $79^{\circ} \mathrm{N}$. This is north of the earlier known area of the species. The individuals sampled did not belong to the River Imsa population, but other rivers chiefly in Northern Norway. It is known that feeding areas of salmonids have changed because of large-scale climate-driven ecosystem shifts in the ocean (Nielsen et al., 2013), and this may hold for many other species (cf. Pörtner \& Peck, 2010). For instance, Potts et al. (2014) reported that Argrosstomus coronus De la Pylaie 1835 on the African coast changed their distribution area because of global warming. Thus, the possibility that the feeding area may have shifted northwards in association with climate warming cannot be ruled out, and if the area has changed, this may have influenced growth opportunities of S. salar.

Marine survival of S. salar in the River Imsa decreased with time before 2000 (Jonsson \& Jonsson, 2004), but there is no such trend in the present material. This may be associated with fisheries regulation and reduced fishing intensity at sea, particularly on MSW
S. salar (cf. Jensen et al., 1999). But the effect of a less intensive fishing on MSW fish probably had minor effect on the size of 1SW fish. Multi-sea-winter S. salar feed mainly on pelagic fish such as C. harengus (Jacobsen \& Hansen, 2001; Haugland et al., 2006), and should not compete keenly with post-smolts for food. So far, there is no evidence of density dependent survival among Atlantic salmon in the ocean (Jonsson et al., 1998). Thus if anything, mean size should have increased with a reduced fishing pressure on relatively large fish, opposite to the observed trend. There is no knowledge about possible genetic effects of reduced fishing in S.salar in the ocean (cf. Marty et al., 2014; Uusi-Heikkilä et al., 2015).

Survival appeared little influenced by water temperature during the post-smolt period. For instance, there was no support for the hypothesis that mortality changed with increasing temperature for either wild or hatchery produced $S$. salar. This seems not to agree with the general view that survival is closely associated with environmental temperature in fishes (Somero, 2004; Griffiths \& Harrod, 2007). In the present case, the effect may be small compared to other sources of mortality, such as predation and parasitism (e.g. Hansen et al., 2003). In addition, variation in age at maturity may have affected their survival. On the other hand, the retrospective growth analysis from scales exhibited a positive correlation between first year growth and survival in wild S. salar. This parallels observations in other populations of S. salar (e.g. McCarty et al., 2008; Peyronnets et al., 2008; Friedland et al., 2009) and in other anadromous species such as S. trutta (Jonsson \& Jonsson, 2009) and Chinook salmon (Woodson et al., 2013). Growth may mediate survival by the functional relationship between post-smolts and their predators as discussed in Jonsson \& Jonsson (2009). The first period of marine life may be particularly demanding for the fish with high mortality when slower growing individuals may be lost because of predation, disease or starvation (Wells et al., 2012). Particularly in years when feeding opportunities are poor, large recruits may survive
better than smaller conspecifics. Saloniemi et al. (2004) and Cross et al. (2008) reported that that relatively large body size gave survival benefits for juvenile $S$. salar in years when the mortality was high.

There were close correlation between wild and hatchery-produced $S$. salar concerning $C F$ and age at maturity. This indicates that environmental conditions at sea influenced these life history variables similarly, and that life history variation in released hatchery S. salar at least partly reflects that of wild fish with similar genetic structure. Thus, hatchery produced $S$. salar may be used as an ecological indicator on production conditions for wild S. salar at sea. From the Pacific Northwest, it was reported that in spite of temporal and spatial variation, the general trend in marine survival is similar for both wild and hatchery coho salmon Oncorhynchus kisutch (Walbaum 1792) (Coronado \& Hilborn, 1998) indicating that this has wider application than for this population.

In all, the present study revealed a significant change in life history of S. salar of the River Imsa strain since the 1970s. The main drivers appear to be the ecosystem change with the collapse in the Norwegian spring spawning $C$. harengus and the subsequent rebuilding of the pelagic fish abundance, together with a decrease in large zooplankton abundance and increase in sea-surface temperature in the Norwegian Sea. Smaller energy stores may render S. salar more susceptible to parasitism and the distribution area may move northwards where growth opportunities are poorer. The pelagic ecosystem is highly dynamic, and as a predator, the life history of S.salar is highly dependent on other species. With large fluctuations in primary and secondary production in the food web, bottom-up effects on S. salar and other predators are expected.

The authors are indebted to the staff of the NINA Research Station at Ims for rearing the fish and daily monitoring the fish trap during the study period. Marius Jonsson, University of Oslo, estimated the nonlinear regressions. Economic support was received from the Norwegian Institute for Nature Research, the Norwegian Environmental Agency, and the Norwegian Reasearch Council’s reseach programme NORKLIMA (project no 225014). This research was approved by the Norwegian Animal Research Experimentation Agency (NARA).

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\(\left.$$
\begin{array}{lcc}\text { Regression } & r^{2} & F-\text { statistics }\end{array}
$$ \begin{array}{l}Degrees of <br>

freedom\end{array}\right\}\)|  |  |  |
| :--- | :--- | :--- |
| $L_{\mathrm{H}}=3680.5( \pm 3.14 .2)-1.55(( \pm 0.16) Y$ | $96.5^{* * *}$ | 1,2255 |
| $L_{\mathrm{W}}=5539.7( \pm 409.1)-2.48( \pm 0.21) Y$ | $146.1^{* * *}$ | 1,830 |
| $C F_{\mathrm{H}}=2.09( \pm 0.50)-0.001 Y$ | $6.93^{* * *}$ | 1,2242 |
| $C F_{\mathrm{W}}=3.58( \pm 0.646)-0.001 Y$ | $18.82^{* * *}$ | 1,830 |
| $G_{\mathrm{H}}=13.06( \pm 1.60)-0.006( \pm 0.001) Y$ | $57.6^{* * *}$ | 1,662 |
| $G_{\mathrm{W}}=5.62( \pm 1.34)-0.002( \pm 0.001) Y$ | $11.5^{* * *}$ | 1,665 |

Table 2. Significant relationships between life history variables of released hatchery (H) and wild $=($ W) S. salar of the River Imsa strain migrating to sea between 1976 and 2010 and (a) Sea surface temperature, and (b) between life history variables. $P M=$ Proportion (\%) of annual cohort that attained maturity as one-sea-winter fish, $C F=$ Fulton's condition factor , $L$ $=$ Total length at maturity $(\mathrm{mm}), G=$ Growth during the first year at sea estimated from scale measurements, $T_{\mathrm{su}}=$ Mean temperature at $0-10 \mathrm{~m}$ depth from April through September in the Norwegian Sea during the first year the fish were at sea, $T_{\mathrm{Wi}}=$ Mean temperature at $0-10 \mathrm{~m}$ depth from October through September in the Norwegian Sea during the first year the fish were at sea, $S=$ Percentage of cohort that returned to the River Imsa after feeding in the ocean. The parameters are given with standard error ( $\pm$ S.E.) and were significant ( $P<0.05$ ). $r^{2}$ $=$ coefficient of determination, $F=F$-statistics, d.f. $=$ degrees of freedom, $*=\mathrm{P}<0.05, * *=\mathrm{P}$ $<0.01, * * *=P<0.001$.

| Regression | $r^{2}$ | $F$ - statistics | Degree <br> freedon |
| :--- | :---: | :---: | :---: |
| (a) With Sea Surface Temperature |  |  |  |
| $L_{\mathrm{H}}=882.0( \pm 111.9)-29.6( \pm 11.6) T_{\mathrm{Su}}$ | 0.20 | $6.51^{*}$ | 1,26 |
| $L_{\mathrm{W}}=857.5( \pm 1.27 .2)-28.6( \pm 13.28) T_{\mathrm{Su}}$ | 0.13 | $4.65^{*}$ | 1,30 |
| $L_{\mathrm{H}}=902.4( \pm 105.8)-43.7( \pm 15.5) T_{\mathrm{Wi}}$ | 0.18 | $8.21^{* *}$ | 1,27 |
| $L_{\mathrm{W}}=967.5( \pm 102.6)-55.6( \pm 14.8) T_{\mathrm{Wi}}$ | 0.32 | $14.0^{* * *}$ | 1,30 |
| $C F_{\mathrm{H}}=1.18( \pm 0.18)-0.042( \pm 0.019) T_{\mathrm{Su}}$ | 0.16 | $5.02^{*}$ | 1,27 |


| $C F_{\mathrm{W}}=1.33( \pm 0.15)-0.055( \pm 0.016) T_{\mathrm{Su}}$ | 0.29 | $12.6^{* * *}$ | 1,31 |
| :--- | :--- | :--- | :--- |
| $C F_{\mathrm{H}}=1.27( \pm 0.15)-0.072( \pm 0.021) T_{\mathrm{Wi}}$ | 0.29 | $11.5^{* *}$ | 1,28 |
| $C F_{\mathrm{W}}=1.18( \pm 0.15)-0.054(0.021) T_{\mathrm{Wi}}$ | 0.18 | $6.59^{*}$ | 1,31 |
| $P M_{\mathrm{H}}=272.60( \pm 82.11)-19.72( \pm 8.49) T_{\mathrm{Su}}$ | 0.18 | $5.39^{*}$ | 1,25 |
| $P M_{\mathrm{H}}=278.31( \pm 63.46)-28.17( \pm 9.13) T_{\mathrm{Wi}}$ | 0.27 | $9.52^{* *}$ | 1,26 |
| (b) Between life history variables |  |  |  |


| $L_{\mathrm{H}}=419.64( \pm 62.06)+188.97( \pm 65.98) G_{\mathrm{H}}$ | 0.20 | $5.03^{*}$ | 1,20 |
| :--- | :--- | :--- | :--- |
| $L_{\mathrm{W}}=355.14( \pm 90.96)+226.85( \pm 87.78)$ | 0.22 | $6.68^{*}$ | 1,24 |
| $G_{\mathrm{W}}$ |  |  |  |

$P M_{\mathrm{H}}=-118.7( \pm 51.7)+260.4( \pm 66.66) \quad 0.37 \quad 15.27 * * * \quad 1,26$
$C F_{\mathrm{H}}$
$P M_{\mathrm{W}}=-67.3( \pm 54.1)+181.6( \pm 68.0) C F_{\mathrm{W}} \quad 0.20 \quad 7.14^{* *} \quad 1,29$
$P M_{\mathrm{H}}=-71.02( \pm 46.00)+161.54( \pm 49.07) \quad 0.35 \quad 10.8^{* *}$
$G_{\mathrm{H}}$
$S_{\mathrm{W}}=-6.15( \pm 4.31)+9.61( \pm 4.16) G_{\mathrm{W}}$
0.18
5.33*
1, 24

## Legend to figures

FIG. 1. Northern North Sea and eastern Norwegian Sea with the location of the River Imsa at the Southwest coast of Norway and the transect further north where near-surface temperature values were retrieved.

FIG. 2. Mean surface temperature ( $0-10 \mathrm{~m}$ depth ) at $62^{\circ} \mathrm{N}$ in the Northeast Atlantic Ocean ( $T$ ${ }^{\circ} \mathrm{C}$ ) during (a) April-September 1976-2010 $\left(Y_{\mathrm{S}}\right): T_{\mathrm{S}}=-51.96( \pm 13.92)+0.03(0.007) Y ; r^{2}=$ $0.39, F_{1,31}=19.5 ; P<0.001$, (b) October-March $1976-2011\left(Y_{\mathrm{W}}\right): T_{\mathrm{W}}=50.49( \pm 9.70)+0.029$ $( \pm 0.005) Y, r^{2}=0.52, F_{1,32}=35.0, P<0.001$.

FIG. 3. Regressions of wild S. salar of the River Imsa cohorts migrating to sea between 1976 and 2010, and hatchery fish produced from the same stock and released when ready for seaward migration between 1981 and 2010, on year of release $(Y)$. (a) Total length at return of 1SW hatchery fish $\left(L_{H} m m\right): L_{H}=5582( \pm 1244)-2.5( \pm 0.62) Y ; r^{2}=0.36, F_{1,28}=16.0, P<$ 0.001 . (b) Total length at return of 1 SW wild fish $\left(L_{\mathrm{w}}\right): L_{\mathrm{W}}=6104.1( \pm 988.0)-2.77( \pm 0.50)$ $Y ; r^{2}=0.51, F_{1,30}=31.2, P<0.001$. (c) Condition factor of hatchery fish $\left(C F_{\mathrm{H}}\right): C F_{\mathrm{H}}=5.95$ $( \pm 2.04)-0.003( \pm 0.001) Y ; r^{2}=0.19, F_{1,28}=6.46, P=0.02$. (d) Condition factor of wild fish $\left(C F_{\mathrm{W}}\right): C F_{\mathrm{W}}=4.67( \pm 1.69)-0.002( \pm 0.001) Y ; r^{2}=0.15, F_{1,31}=5.27, P=0.03$. (e) First-year growth estimated from scales of hatchery fish $\left(G_{\mathrm{H}}\right): G_{\mathrm{H}}=12.43( \pm 2.04)-0.006( \pm 0.001) \mathrm{Y} ; r^{2}$ $=0.37, F_{1,20}=11.9, P=0.002$. (f) First-year growth estimated from scales of wild fish $\left(G_{\mathrm{W}}\right)$ : $G_{\mathrm{W}}=7.37( \pm 2.61)-0.003( \pm 0.001) Y ; r^{2}=0.20, F_{1,24}=5.9, P=0.02$. (g) Percent 1 SW hatchery fish of adults returning to the River Imsa $\left(P M_{H} \%\right): P M_{H}=1.16( \pm 2.04)(\mathrm{Y}-$
1909.4)^1.01( $\pm 1.13) * \exp \left(-\left((\mathrm{Y}-1909.4)^{*} 0.01\right)^{\wedge 18.43( \pm 1.13)}\right), R_{4,24^{2}}=0.71, P<0.01$. (h) Percent 1SW wild fish of adults returning to the River Imsa ( $P M_{\mathrm{W}} \%$ ): $P M_{\mathrm{W}}=1.93( \pm 4.45$ $\left(*(Y-1950.6)^{\wedge 1.09}( \pm 0.57) * \exp \left(-\left((\mathrm{Y}-1950.6)^{*} 0.018( \pm 0.001)\right)^{\wedge} 5.84( \pm 3.04)\right), R_{4,27^{2}}=0.59, P\right.$ $<0.01$.

FIG 4. Regression of (a) total body length of wild ( $L_{\mathrm{w}} \mathrm{mm}$ ) on hatchery produced ( $L_{\mathrm{H}} \mathrm{mm}$ ) River Imsa S. salar migrating to sea from 1981 to 2010 and returning as 1 SW fish: $L_{\mathrm{W}}=$ $142.4( \pm 115.1)+0.73( \pm 0.19) L_{\mathrm{H}} ; r^{2}=0.37, F_{1,25}=14.3, P<0.001$. (b) Mean condition factors of wild $\left(C F_{\mathrm{W}}\right)$ on hatchery produced $\left(C F_{\mathrm{H}}\right)$ River Imsa salmon migrating to sea from 1981 to 2010 and returning as 1 SW fish $\left(C F_{\mathrm{W}}\right): C F_{\mathrm{W}}=0.14( \pm 0.08)+.85( \pm 0.11) C F_{\mathrm{W}} ; R^{2}=$ $0.70, F_{1,27}=62, P<0.001$. (c) Proportion of wild $\left(P_{\mathrm{w}}\right)$ on hatchery produced ( $P_{\mathrm{H}} \mathrm{mm}$ ) River Imsa salmon migrating to sea from 1981 to 2010 and returning as 1 SW fish: $P M_{\mathrm{W}}=18.3$ $( \pm 6.3)+0.57( \pm 0.07) 78 P M_{H} ; r^{2}=0.71, F_{1,24}=58.1, P<0.001$.


Figure 1.


Figure 2.


Figure 3.




Figure 4.

