



# Environmental and endocrine control of gill corticosteroid receptor number and affinity in Atlantic salmon (*Salmo salar*) during smolting

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## Abstract

During smolting, cortisol acts on the gill through intracellular corticosteroid receptors (CR). Regulation of CR concentration ( $B_{\max}$ ) and dissociation constant ( $k_d$ ) by environmental factors, however, has not been investigated. We subjected juvenile Atlantic salmon (*Salmo salar*) to changes in photoperiod and temperature to determine the effect on gill CR  $B_{\max}$  and  $k_d$ . Cortisol, growth hormone (GH), insulin-like growth factor 1 (IGF-1), thyroxine ( $T_4$ ), and triiodothyronine ( $T_3$ ) were measured to determine endocrine factors that correlated with changes in CR  $B_{\max}$  and  $k_d$ . Gill  $\text{Na}^+, \text{K}^+$ -ATPase activity was measured as an indicator of smolting. Control fish were maintained under ambient Connecticut River water temperatures and natural photoperiod (LDN). In the first experiment, fish were also reared at elevated temperature (constant 10 °C), or long day photoperiod (LD 16:8; 16 h light), or a combination of these two treatments. In the second experiment, fish were subjected to an advanced river temperature regime, or short day photoperiod (LD 9:15; 9 h light), or a combination of these two treatments. Seasonal changes in CR  $B_{\max}$  were found to be significantly affected by temperature, but not photoperiod. A decline in CR  $B_{\max}$  occurred when mean daily temperature increase exceeded 1.5 °C per week, preceding the increase in gill  $\text{Na}^+, \text{K}^+$ -ATPase activity. CR  $B_{\max}$  was found to be correlated positively with  $T_4$  and negatively with IGF-1. Gill CR  $k_d$  changed significantly over the spring,

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but manipulation of temperature and photoperiod had little effect. CR  $k_d$  was found to be positively correlated with gill  $\text{Na}^+, \text{K}^+$ -ATPase activity, plasma GH, cortisol, IGF-1 and  $T_4$ . Temperature appears to influence seasonal changes in CR  $B_{\max}$  observed, whereas endocrine factors appear to be more closely related to seasonal changes in CR  $k_d$ .

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## 1. Introduction

Smolting is a preparatory physiological adaptation in which salmon develop tolerance to increased salinity just prior to migration to the sea. This physiological transformation is stimulated by the increase in day length that occurs during the spring (McCormick and Saunders, 1987). Altering the natural photoperiod cycle has been shown to shift the timing of smolting, evidence that photoperiod acts as a zeitgeber for smolting (Clarke et al., 1978). Temperature has also been found to affect smolting. Seasonal increase in water temperature concurrent with an increase in photoperiod is a stronger stimulus to smolting than each alone (Muir et al., 1994). The evidence for temperature directly stimulating smolting is equivocal in the absence of a photoperiod stimulus (see McCormick et al., 1987; Staurnes et al., 1994). Low temperature, however, has been shown to limit the effect of photoperiod on smolting when Atlantic salmon were subjected to long day length (16 h light) in mid-February (McCormick et al., 2000).

The endocrine system is the primary signaling pathway between the external stimuli and the seasonal physiological response. Sensitivity of cells and tissues to a given hormone is dependent on stability of the hormone receptor complex, affinity of the ligand for the receptor, and the number of receptors specific for the hormone in question (Clark and Peck, 1977). Changes in receptor concentration and affinity have been observed during development indicating that sensitivity of tissues changes with life stage. For example, during metamorphosis in amphibians, the increase in circulating thyroxine levels is associated with increased liver expression of the thyroid hormone receptor gene (Yaoita and Brown, 1990) and higher concentrations of the liver thyroid hormone receptor protein (Eliceiri and Brown, 1994). Unlike metamorphosis where a single endocrine pathway is dominant, smolting involves a number of interacting endocrine systems. Cortisol, growth hormone (GH), insulin-like growth factor 1 (IGF-1), and thyroid hormones increase during the spring in response to seasonal changes in photoperiod and temperature and stimulate smolting (Hoar, 1988). For one of these hormones, cortisol, we have observed changes in corticosteroid receptor (CR) concentration ( $B_{\max}$ ) and dissociation constant ( $k_d$ ) when fish smolt during the spring (Shrimpton and McCormick, 1998a; Shrimpton et al., 2000). Given the importance of photoperiod and temperature on the parr–smolt transformation, we varied these variables to determine their influence on gill CR of Atlantic salmon during smolting.

## 2. Materials and methods

### 2.1. Experiment 1—increased temperature and long-day photoperiod

Juvenile Atlantic salmon were obtained as parr from the White River National Fish Hatchery (Bethel VT, USA) and brought to the Conte Anadromous Fish Research Center in mid-October. Atlantic salmon show bimodal growth distribution and by November, there is a clear difference in size between the upper mode fish which smolt the following spring and the lower mode fish that will smolt 1 year later. The fish were graded, and upper mode fish were randomly divided into four isolated photoperiod rooms containing two 1-m diameter tanks supplied with ambient river water at a flow rate of 4 l/min and supplemental aeration. Lighting was supplied by overhead fluorescent lights (500 lx at the water surface), and photoperiod adjusted twice a week. Each tank contained approximately 80 fish. The fish were fed to satiation (Zeigler, Gardners, PA) with automatic feeders.

Initially, all groups were maintained on a natural photoperiod (LDN) and water was maintained at ambient temperature in all tanks. On 7 January, two of the groups were supplemented with heated water to maintain a temperature of 9–10 °C. On 8 February, one group in each of the temperature regimens was subjected to an abrupt increase in day length to 16 h (LD 16:8).

### 2.2. Experiment 2—advanced temperature and short-day photoperiod

Juvenile Atlantic salmon were obtained and treated as described above. Initially, all groups were maintained on a simulated natural photoperiod and water was maintained at ambient temperature in all tanks. On 12 January, two groups remained on short days (LD 9:15; 9 h daylight) while the remaining two groups continued on a light dark natural photoperiod (LDN). On 14 February, one group in each of the photoperiod treatments was supplemented with heated water to mimic the natural temperature increase that occurs during the spring, but advanced by approximately 1 month.

### 2.3. Fish and sampling procedures

At approximately 2-week intervals throughout the spring, six fish were rapidly netted from a tank and transferred to a bucket containing 200 mg l<sup>-1</sup> tricaine methane sulphonate (neutralized and buffered with sodium bicarbonate, pH 7.0). Once the fish were anesthetized, fork length (*L*) and body weight (*W*) were measured. Blood was collected in heparinized syringes from the caudal vasculature. Collection of blood was completed within 5 min of first disturbing the fish to ensure that a stress-associated rise in cortisol did not occur (Sumpter et al., 1986). Blood was stored on ice for less than 30 min, centrifuged at 3000 × *g* for 5 min, plasma removed and frozen on dry ice. A gill biopsy (approximately six to eight primary gill filaments) was taken and placed in 100 µl of SEI (150 mM sucrose, 10 mM Na<sub>2</sub>EDTA, 50 mM imidazole, pH 7.3) on ice for determining Na<sup>+</sup>,K<sup>+</sup>-ATPase activity. Samples were frozen on dry ice within 30 min. The remaining gill tissue was removed and placed in 2 ml of TEMS (10 mM

Tris–HCl, 1 mM Na<sub>2</sub>EDTA, 12 mM monothioglycerol, 20 mM sodium molybdate, 10% v/v glycerol, pH 7.4) and frozen immediately on dry ice for later analysis of corticosteroid receptor concentration and affinity. All samples were stored at –80 °C until analyses. After the first tank from each room was sampled, six fish from the second tank were sampled as described above. Only the first three fish sampled from each tank, however, were included in this data set due to the length of time needed to remove all the tissue for corticosteroid receptor samples. There was no significant difference between tanks for any of the parameters measured. Samples from both tanks were pooled for analysis and sample size (*n*) equals six for each parameter reported in this paper.

#### 2.4. Analysis of gill Na<sup>+</sup>,K<sup>+</sup>-ATPase activity

Gill Na<sup>+</sup>,K<sup>+</sup>-ATPase activity was measured according to the microassay protocol of McCormick (1993). Gill filaments were homogenized in SEI buffer containing 0.1% sodium deoxycholate. Following centrifugation (3000 × *g* for 0.5 min) to remove large debris, Na<sup>+</sup>,K<sup>+</sup>-ATPase activity was determined by linking ATP hydrolysis to the oxidation of nicotinamide adenine dinucleotide (NADH), measured at 340 nm for 10 min at 25 °C, in the presence and absence of 0.5 mM ouabain. Protein content in the gill homogenate was measured using a bicinchoninic acid (BCA) protein assay (Pierce, Rockford, IL, USA). Specific activities were expressed as μmol ADP mg<sup>-1</sup> of protein h<sup>-1</sup>.

#### 2.5. Determination of plasma hormone levels

Plasma GH levels were measured in duplicate samples using a specific double-antibody salmon GH radioimmunoassay outlined by Björnsson et al. (1994). Plasma IGF-1 levels were measured by a homologous radioimmunoassay as outlined by Moriyama et al. (1994). Plasma cortisol levels were quantified using a competitive solid-phase microtitre enzyme immunoassay (EIA) following the protocol of Carey and McCormick (1998).

#### 2.6. Corticosteroid receptor analysis

Corticosteroid receptors were measured on the cytosol fraction of gill tissue using the method of Maule and Schreck (1991) as modified by Shrimpton and McCormick (1998b). Binding studies were conducted with [<sup>3</sup>H]triamcinolone acetonide (TA; 1,4-pregnadien-9α-fluoro-11β,16α,-17α,21-tetrol-3,20-dione-16,17 acetonide) with a specific activity of 43.8 Ci mmol<sup>-1</sup> (Dupont-NEN). Gill cytosol (100 μl) was incubated in aliquots with 100 μl of buffer containing [<sup>3</sup>H]TA with or without a 500-fold excess of cold TA for 2 h on ice. Final concentration of [<sup>3</sup>H]TA in each assay were 0.1, 0.3, 1, 3, and 6 nM. After incubation, unbound steroids were removed by incubation for 10 min with 0.5 ml of TEMS containing 2.5% (w/v) activated charcoal and 0.25% (w/v) dextran and then centrifuged at 3000 × *g* for 15 min. Supernatant (0.5 ml) was added to 3 ml of aqueous counting scintillant and radioactivity counted. Specific binding was determined by subtracting nonspecific bound from the total bound.

The origin of corticosteroid receptors in the gills may be cytosolic or nuclear, but are referred to as cytosolic as they are found in the cytosol fraction following tissue processing (Welshons and Jordon, 1987). The CR concentration measured is comprised of the unbound receptor population. The equilibrium dissociation constant ( $K_d$ ) and the concentration of corticosteroid receptor sites ( $B_{max}$ ) were calculated according to Scatchard (1949).  $B_{max}$  was divided by the homogenate protein concentration, and CR concentration was expressed as fmol  $mg^{-1}$  protein. To estimate cooperativity between CR and ligand, the Hill coefficient was calculated according to Sandor et al. (1984).

Affinities of CR for cortisol measured in this study are similar to those found for the glucocorticoid receptor (GR) in vertebrates, and of lower affinity than for the mineralocorticoid receptor (MR) (Diaz et al., 1998). Competition experiments with other steroids indicated that the two synthetic glucocorticoids TA and dexamethasone showed the highest affinity for the ligand (Shrimpton and McCormick, 1999), also a characteristic of GR (Ducouret et al., 1995), but distinct from the MR homologue recently found in rainbow trout (Colombe et al., 2000). A true mineralocorticoid function of this receptor in fish has yet to be determined. Receptors measured in this study are characteristic of GR, but we refer to the receptors as CR to avoid any confusion concerning their physiological action.

### 2.7. Calculations and statistical analysis

For seasonal changes in gill CR  $B_{max}$ ,  $K_d$ , and gill  $Na^+,K^+$ -ATPase activity, a three-way analysis of variance (ANOVA) was used to determine whether time of sampling, temperature regime or photoperiod regime had a significant effect on these variables. When factors were found to be statistically significant, Tukey's test was used to determine differences between the treatments and time interval. Statistical significance was taken at a level of  $P < 0.05$ . All values are expressed as mean  $\pm$  1 S.E.

To examine the role that temperature plays on seasonal changes in gill  $B_{max}$  and  $k_d$ , changes in these parameters were plotted as a function of the accumulated thermal units (ATU). ATU is calculated as the additive daily temperature in degrees Celsius experienced since 1 January. To determine a relationship between the decline in CR  $B_{max}$  and temperature, we calculated the daily change in temperature between the sample date when the decline was observed and the previous sample intervals (2 weeks). The rate of change in temperature was then expressed as temperature change per week ( $\Delta T$ ,  $^{\circ}C/week$ ). The difference in CR  $B_{max}$  between the two sample dates ( $\Delta B_{max}$ ) was compared to  $\Delta T$  by linear regression analysis for each of the different temperature regimes. Photoperiod was not included in the analysis.

Linear regression analysis was conducted on data from all individual animals for both  $B_{max}$  and  $k_d$  as a function of all endocrine parameters. Correlation analysis was also conducted on gill CR and gill  $Na^+,K^+$ -ATPase activity. To examine the relative explanatory power of endocrine parameters on gill  $B_{max}$  and  $k_d$ , a best subsets regression analysis was used. Endocrine parameters (cortisol, GH, IGF-1, triiodothyronine ( $T_3$ ), and thyroxine ( $T_4$ )) were logarithmically transformed for these analyses.

### 3. Results

#### 3.1. Experiment 1

Gill CR  $B_{\max}$  increased steadily from January to April in all groups (Fig. 1). Highest levels were reached in all groups at the end of March, and subsequently declined. Three-way ANOVA indicated that there was a significant effect of time ( $P < 0.001$ ) and temperature ( $P < 0.005$ ), but not photoperiod ( $P = 0.576$ ). There was also an interaction

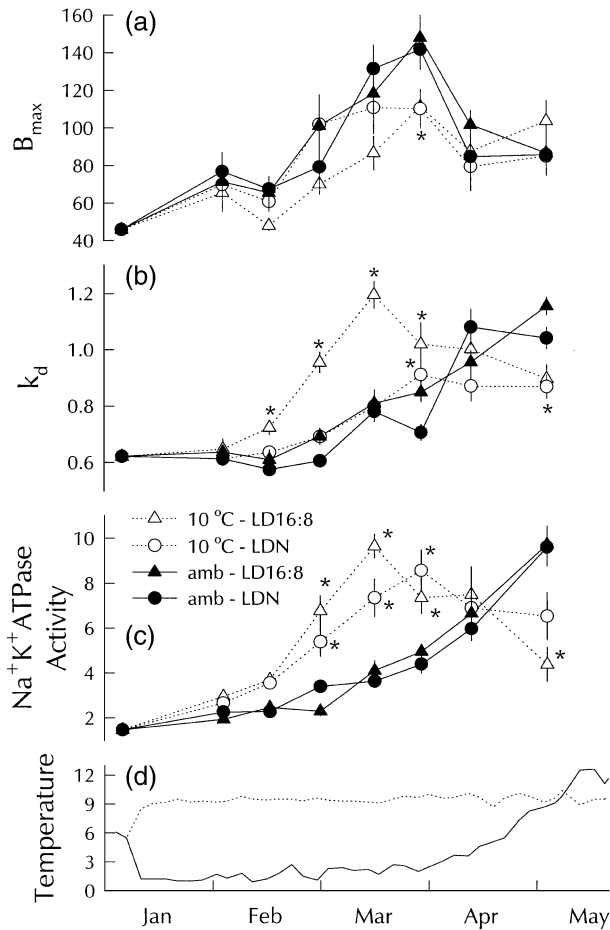


Fig. 1. Seasonal changes in gill corticosteroid receptor (a) concentration ( $B_{\max}$ ; fmol  $\text{mg}^{-1}$  of protein), (b) dissociation constant ( $k_d$ ; nM), (c) gill  $\text{Na}^+, \text{K}^+$ -ATPase activity ( $\mu\text{mol ADP mg}^{-1}$  of protein  $\text{h}^{-1}$ ) for juvenile Atlantic salmon, and (d) water temperature for Experiment 1. 10 °C are groups reared at constant 10 °C temperature, and amb is the natural water temperature of the Connecticut River. LDN is a simulated natural photoperiod and LD 16:8 groups were subjected to 16 h light and 8 h dark photoperiod regime from 8 February to the end of the experiment. \* Indicates value is significantly different from the amb-LDN for the same sampling interval. Values are mean  $\pm$  1 S.E.M.

effect between time and temperature ( $P < 0.05$ ). The 10 °C temperature groups tended to be lower than the ambient temperature groups. Tukey's test found that  $B_{\max}$  was significantly lower (40%) in the 10 °C groups than the ambient groups for the sample point at the end of March.

Gill CR  $k_d$  was significantly affected by time ( $P < 0.001$ ), photoperiod ( $P < 0.001$ ), temperature ( $P < 0.005$ ), and there was an interaction between the three factors ( $P < 0.001$ ). CR  $k_d$  stayed fairly constant until mid-March in three of the groups, but increased steadily after this point. The 10 °C–LD 16:8 group was the exception and showed an earlier rise (1 week after increased day length) (Fig. 1). Tukey's test on all pairwise comparisons indicated that  $k_d$  in the 10 °C–LD 16:8 group was significantly greater than the amb–LDN group at all time points between mid-February and the end of March. There was also a significant difference between the two-photoperiod groups held at 10 °C for the first two March samples. Following the maximum value of  $1.20 \pm 0.05$  nM for the 10 °C–LD 16:8 group,  $k_d$  declined 25% by the end of the experiment. The 10 °C–LDN group showed a smaller increase during the spring than the LD 16:8 group at the same temperature and was also significantly greater than the amb–LDN at the end of March and then declined slightly. The ambient temperature groups continued to increase throughout the study. By the last sample point in mid-May, the amb–LDN group was significantly greater than the 10 °C–LDN group.

Gill  $\text{Na}^+, \text{K}^+$ -ATPase activity increased during the spring. Three-way ANOVA revealed that there was a significant effect of time ( $P < 0.001$ ), and temperature ( $P < 0.001$ ), but not photoperiod ( $P = 0.575$ ). There was also a significant interaction between time and temperature ( $P < 0.001$ ). For fish held in 10 °C water, gill  $\text{Na}^+, \text{K}^+$ -ATPase increased steadily from January to mid-March; however, the increase was advanced in the LD 16:8 group, but the activities did not differ significantly between the two LD 16:8 groups at any time (Fig. 1). Increases in gill  $\text{Na}^+, \text{K}^+$ -ATPase activity were slower in the ambient temperature groups and differed significantly from the 10 °C groups throughout March.

### 3.2. Experiment 2

Gill CR  $B_{\max}$  showed significant changes over the spring (Fig. 2). CR  $B_{\max}$  increased from January until April in the ambient temperature groups, whereas in the advanced temperature groups CR  $B_{\max}$  increased slightly until late February and subsequently declined. There was a significant effect of time ( $P < 0.001$ ), and temperature ( $P < 0.001$ ), but not photoperiod ( $P = 0.418$ ) on CR  $B_{\max}$ . There was also an interaction effect between time and temperature ( $P < 0.001$ ). Pairwise comparisons indicated that  $B_{\max}$  was significantly lower in the advanced temperature groups compared to ambient temperature groups during March and early April.

There was a significant increase in  $k_d$  for all groups over the spring, and a significant drop in  $k_d$  between the last two sample times (Fig. 2). Seasonal changes in gill CR  $k_d$  were less than those seen in the first experiment. ANOVA indicated a significant effect of time ( $P < 0.001$ ), but not photoperiod ( $P = 0.121$ ) or temperature ( $P = 0.311$ ).

Gill  $\text{Na}^+, \text{K}^+$ -ATPase activity increased during the spring. Three-way ANOVA revealed that there was a significant effect of time ( $P < 0.001$ ), and temperature ( $P < 0.001$ ), and

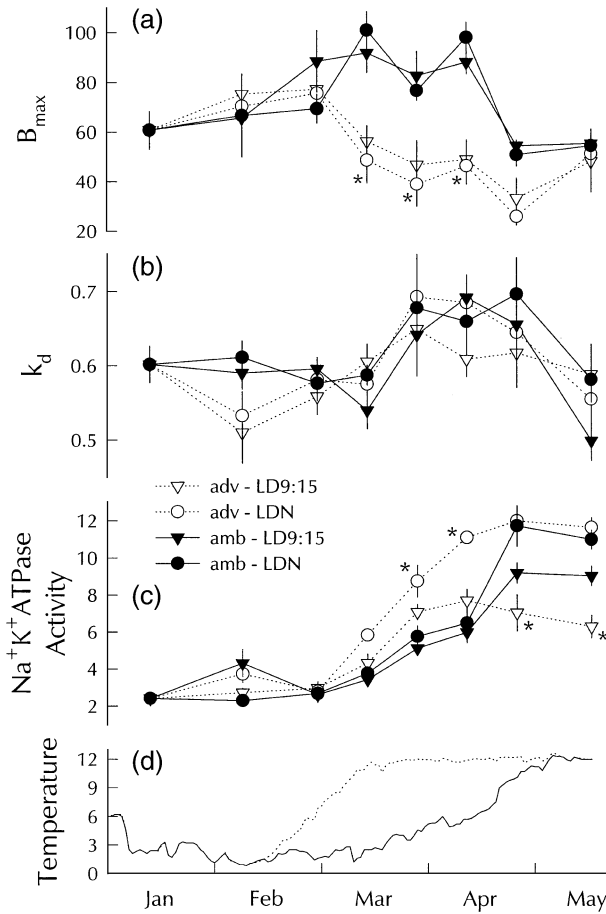


Fig. 2. Seasonal changes in gill corticosteroid receptor (a) concentration ( $B_{max}$ ; fmol  $mg^{-1}$  of protein), (b) dissociation constant ( $k_d$ ; nM), (c) gill  $Na^+,K^+$ -ATPase activity ( $\mu mol ADP mg^{-1}$  of protein  $h^{-1}$ ) for juvenile Atlantic salmon, and (d) water temperature for Experiment 2. adv are groups reared at a temperature regime that was advanced over ambient temperature increases by approximately 6 weeks, and amb is the natural water temperature of the Connecticut River. LDN is a simulated natural photoperiod and LD 9:15 groups were subjected to 9 h light and 15 h dark photoperiod regime from 14 February to the end of the experiment. \* Indicates value is significantly different from the amb-LDN for the same sampling interval. Values are mean  $\pm 1$  S.E.M.

photoperiod ( $P < 0.001$ ). There was also a significant interaction between time, temperature, and photoperiod ( $P < 0.001$ ). In the advanced temperature LDN group, gill  $Na^+,K^+$ -ATPase increased steadily after temperature began to increase but diverged over the spring; activities differed significantly between the two advanced temperature groups in late April and May (Fig. 2). The increase in  $Na^+,K^+$ -ATPase activity was more gradual in the ambient temperature groups, but highest gill  $Na^+,K^+$ -ATPase levels were seen concurrent with the adv-LDN group. There was no significant difference between the two-photoperiod treatments at ambient temperature.



### 3.3. Environmental effects on gill CR

Photoperiod treatments did not appear to influence gill CR  $B_{\max}$  (Figs. 1 and 2). Changes in photoperiod also had little effect on gill CR  $k_d$ , except for the 10 °C group in Experiment 1; transition from LDN to LD 16:8 on 8 February advanced the increase in  $k_d$ .

Temperature, however, did have an affect on gill CR  $B_{\max}$ . This was most noticeable in Experiment 2 when the two temperature regimes are compared. The maximum  $B_{\max}$  and decline in  $B_{\max}$  values were seen earlier in the season in the advanced temperature groups. This trend was not seen in Experiment 1. Maximum gill CR  $B_{\max}$  values were reached at approximately 170 ATU for all the temperature groups, except for constant 10 °C (Table 1). For the seasonally increasing temperature regimes, this corresponded to a difference in time of approximately 1 month. Absolute temperature did not appear to affect the maximum value of  $B_{\max}$  as it varied from 2 to 6.9 °C among the different groups. Maximal  $k_d$  did not appear to be related to the temperature, accumulated thermal units, or date (Table 1).

For the groups that were reared on water that changed seasonally, there was a marked decrease in CR  $B_{\max}$  that occurred after the peak. The interval between the highest  $B_{\max}$  and the decline differed for the different experimental treatments; it ranged from 2 to 6 weeks. The decrease in CR  $B_{\max}$  is temperature dependent as  $B_{\max}$  remained near the seasonal maximum value until the temperature began to increase in the ambient temperature and advanced temperature groups (Figs. 1 and 2). The rate of increase in temperature over the 2-week interval prior to the decline in  $B_{\max}$  was calculated to be greater than 1.5 °C/week. There was less of a decline in the constant 10 °C groups, but this is also reflected in a smaller increase in CR  $B_{\max}$ . A summary of the  $\Delta B_{\max}$  as a function of environmental factors is given in Table 2. An examination of the data for each of the temperature groups indicated that there was a significant relationship between  $\Delta B_{\max}$  and  $\Delta T$  for three of the temperature regimes, but not the constant temperature group. For Experiment 1 ambient and constant temperature groups,  $R^2=0.463$ ,  $P=0.007$ , and  $R^2=0.036$ ,  $P=0.52$ , respectively. For Experiment 2 ambient and advanced temperature groups,  $R^2=0.738$ ,  $P=0.0003$ , and  $R^2=0.408$ ,  $P=0.025$ , respectively.

Table 1

The maximum gill CR  $k_d$  and  $B_{\max}$  for each of the treatment groups for Experiments 1 and 2

	$B_{\max}$	Maximum $B_{\max}$			$k_d$	Maximum $k_d$		
		ATU	Day	°C		ATU	Day	°C
amb-LDN #1	142 ± 11	178	88	2	1.08 ± 0.06	230	102	3.6
amb-LD 16:8	148 ± 13	178	88	2	1.16 ± 0.03	357	122	9.1
amb-LDN #2	101 ± 8	167	73	2.5	0.70 ± 0.05	400	115	10.1
amb-LD 9:15	92 ± 8	167	73	2.5	0.69 ± 0.07	216	87	4.6
adv-LDN	76 ± 5	170	59	6.9	0.69 ± 0.03	629	101	11.9
adv-LD 9:15	77 ± 8	170	59	6.9	0.65 ± 0.06	463	87	12
10 °C-LDN	111 ± 14	674	75	9.1	0.91 ± 0.04	801	88	9.7
10 °C-LD 16:8	111 ± 8	801	88	9.7	1.20 ± 0.05	674	75	9.1

The accumulated thermal units (ATU, °C × day), number of days since 1 January, and the temperature (°C) when maximal values for  $k_d$  and  $B_{\max}$  were reached are listed.

Table 2

Temperature change over the 2-week interval prior to the decline in  $B_{\max}$  for each of the study groups

	Mean $\Delta B_{\max}$	Environmental conditions			
		$\Delta T$	Average $T$	ATU	Day
Ambient #1	-51.6	$1.81 \pm 0.18$	$2.1 \pm 0.1$	227	102
Ambient #2	-40.4	$2.03 \pm 0.13$	$2.3 \pm 0.1$	400	115
Advanced	-24.0	$2.53 \pm 0.05$	$2.9 \pm 0.2$	300	73
10 °C	-27.2	$0.05 \pm 0.06$	$8.9 \pm 0.1$	928	102

The temperature change ( $\Delta T$ ) is expressed as the degree temperature change per week ( $^{\circ}\text{C}/\text{week}$ ) and was calculated over the interval of 2 weeks prior to the decline in  $B_{\max}$ . Average temperature is calculated as the average from 1 January to the maximum value of  $B_{\max}$  in  $^{\circ}\text{C}$  for each of the treatment groups. The accumulated thermal units (ATU,  $^{\circ}\text{C} \times \text{day}$ ) were calculated from 1 January. Day is the number of days since 1 January.

### 3.4. Gill CR correlations with endocrine factors and gill $\text{Na}^+, \text{K}^+$ -ATPase activity

Correlations for gill CR  $B_{\max}$  with individual plasma hormone concentrations are shown in Table 3. Correlations with  $B_{\max}$  were strong for plasma  $\text{T}_4$  and plasma IGF-1, weak for plasma GH and plasma cortisol, and not significant for plasma  $\text{T}_3$ . A best subsets regression model of hormones on gill  $B_{\max}$  found that plasma GH, IGF-1, and  $\text{T}_4$  were significant parameters ( $P < 0.0001$ ,  $R^2 = 0.119$ ); cortisol and  $\text{T}_3$  were excluded from the analysis during the stepwise regression. Coefficients for GH (11.6) and  $\text{T}_4$  (20.6) were positive, but the coefficient for IGF-1 ( $-71.8$ ) was negative.

Correlations for gill CR  $k_d$  and plasma hormone concentrations are shown in Table 3. Strong correlations existed for  $k_d$  and all hormones measured except for  $\text{T}_3$  which was not correlated with  $k_d$ . The best fit for a single hormone on  $k_d$  was for GH; mean gill CR  $k_d$  is plotted against mean plasma GH (Fig. 3;  $R^2 = 0.48$ ). A best subsets regression model of hormones on gill  $k_d$  found that plasma cortisol, GH, and  $\text{T}_4$  had a significant effect on the analysis ( $P < 0.0001$ ,  $R^2 = 0.214$ ); IGF-1 and  $\text{T}_3$  were excluded from the analysis during the stepwise regression. Coefficients for all hormones were positive; 0.153, 0.079, and 0.059 for GH,  $\text{T}_4$ , and cortisol, respectively.

Table 3

Results of correlation analysis for CR  $B_{\max}$  and  $k_d$  when regressed against measured plasma levels of cortisol, growth hormone (GH), insulin-like growth factor 1 (IGF-1), thyroxine ( $\text{T}_4$ ), and triiodothyronine ( $\text{T}_3$ )

	Hormone	Coefficient	$P$
$B_{\max}$	Cortisol	-4.44	0.079
	GH	6.49	0.068
	IGF-1	-39.14	<0.001
	$\text{T}_4$	21.85	<0.001
	$\text{T}_3$	-3.95	0.808
$k_d$	Cortisol	0.072	<0.0001
	GH	0.164	<0.0001
	IGF-1	0.192	0.001
	$\text{T}_4$	0.072	0.048
	$\text{T}_3$	-0.066	0.463

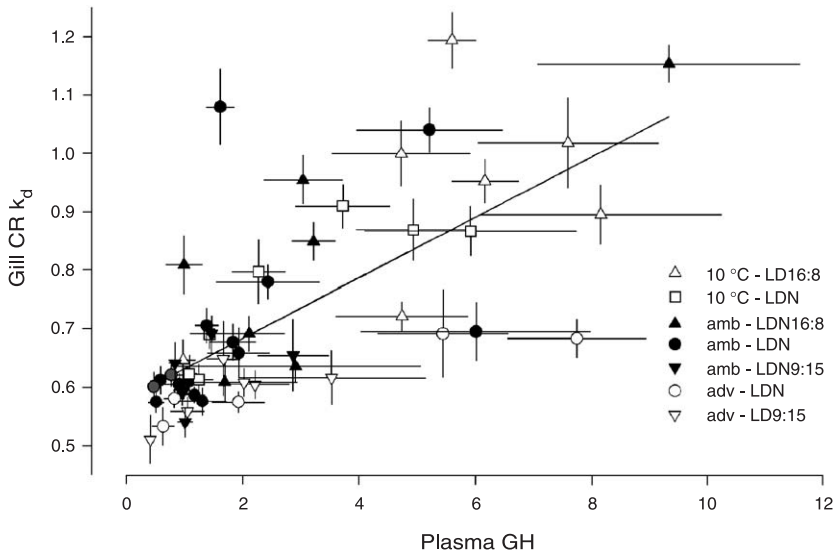


Fig. 3. Regression of plasma growth hormone with gill CR  $k_d$ . Values are means ( $n=6$ ) for each time point from each of the photoperiod and temperature treatments. Groups as described in Figs. 1 and 2.

Gill CR  $B_{max}$  was not correlated with gill  $\text{Na}^+, \text{K}^+$ -ATPase activity. Regression analysis, however, was significant for gill CR  $k_d$  and gill  $\text{Na}^+, \text{K}^+$ -ATPase activity and is plotted in Fig. 4 ( $R^2=0.33$ ).

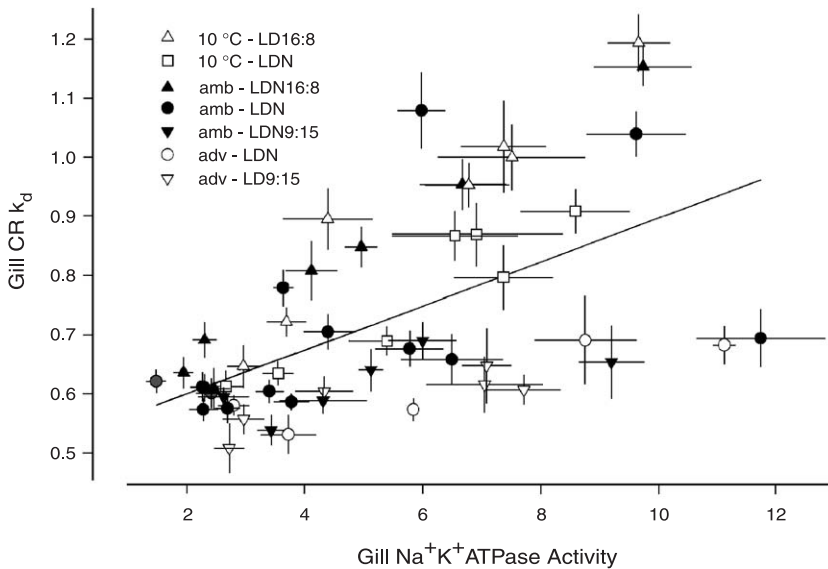


Fig. 4. Regression of gill  $\text{Na}^+, \text{K}^+$ -ATPase activity with gill CR  $k_d$ . Values are means ( $n=6$ ) for each time point from each of the photoperiod and temperature treatments. Groups as described in Figs. 1 and 2.

#### 4. Discussion

Changes in the corticosteroid receptor have been seen during development in fish and are thought to be important to the ontogeny of hormone action. During early embryonic development in tilapia, (*Oreochromis mossambicus*), the highest concentration of cortisol receptor mRNA was found just after fertilization (Tagawa et al., 1997). Gill CR concentration ( $B_{\max}$ ) and dissociation constant ( $k_d$ ) have been observed to change seasonally in Atlantic salmon (Shrimpton and McCormick, 1998a; Shrimpton et al., 2000). The density of GR immunoreactive neurons was greater in olfactory regions of the brain in sexually mature sockeye salmon (*Oncorhynchus nerka*) compared to immature fish (Carruth et al., 2000). These observations indicate that CR dynamics change seasonally with developmental stage in fish and increase when cortisol is functionally important. The present study sought to characterize environmental and endocrine factors that regulate gill CR in Atlantic salmon during the parr–smolt transformation.

A seasonal increase in plasma cortisol is one of the endocrine factors that stimulate smolting in juvenile salmon (Hoar, 1988). The mechanism of cortisol action in the gills is mediated by the CR (Clark and Peck, 1977). A direct relationship between tissue sensitivity to cortisol and CR concentration, and to a lesser extent affinity, has recently been shown by Shrimpton and McCormick (1999). For all study groups in Experiments 1 and 2 (except 10 °C), maximal values of  $B_{\max}$  were reached earlier in the year than gill  $\text{Na}^+, \text{K}^+$ -ATPase activity. The increase in  $B_{\max}$  prior to the peak in smolting will increase sensitivity of the gill to cortisol, before the springtime increase in cortisol. The increase in  $B_{\max}$ , therefore, appears to be preparatory for the seasonal increase in cortisol that plays a role in stimulating smolting. Changes in gill CR  $k_d$  occurred synchronously with changes in gill  $\text{Na}^+, \text{K}^+$ -ATPase activity, and appear to reflect physiological changes associated with smolting.

##### 4.1. Environmental regulation of CR

Photoperiod is the primary stimulus for smolting (McCormick et al., 1987). The lack of a change in CR  $B_{\max}$  following photoperiod manipulation, therefore, is surprising. It appears that the seasonal changes in  $B_{\max}$  are independent of photoperiod. Temperature does have an effect on CR  $B_{\max}$  (Tables 1 and 2). Temperature affects CR  $B_{\max}$  in two ways; first, the seasonal increase is independent of a change in temperature but correlated with ATUs, and second, the decrease in  $B_{\max}$  is dependent on the seasonal increase in temperature.

The maximum values of  $B_{\max}$  corresponded to approximately 170 ATU for the ambient and advanced temperature groups, but were not a function of date. An examination of other studies that have shown changes in CR  $B_{\max}$  over the spring in Atlantic salmon indicates that ATU is also related to maximum  $B_{\max}$ . A calculation of maximum  $B_{\max}$  in the study by Shrimpton and McCormick (1998a) found that the maximum was reached at 247 ATU. This is greater than the ATU calculated in the present study; however, the sampling interval was monthly. Two weeks earlier, the mid-point between the two sampling intervals would correspond to 184 ATU and a value similar to that calculated in the present study. Average temperature was 2.5 °C from January to May, and the temperature was 5.6 °C when the maximum was reached. The relationship between ATU

and maximum  $B_{\max}$  did not hold for another study where water temperature during the winter months was warmer ( $>8$  °C during January and February) (Shrimpton et al., 2000). It appears, therefore, that when water temperatures are below 3 °C,  $B_{\max}$  is closely associated with ATU. The strong correlation between maximum  $B_{\max}$  and ATU is suggestive that the rate of development is dependent on temperature not date, as the timing of the highest values of  $B_{\max}$  differed by approximately 1 month (Table 1).

The decline in  $B_{\max}$  was found to be a function of temperature change for groups where temperature was not held constant. As the rate of increase in temperature exceeded 1.5 °C/week during the 2-week interval before the sample, there was a marked decrease in  $B_{\max}$  (Table 2). Regression analysis of  $\Delta B_{\max}$  on  $\Delta T$  also indicate that the seasonal increase in temperature drives the decline in  $B_{\max}$ . A similar relationship was calculated for the data published by Shrimpton and McCormick (1998a). In the 10 °C constant temperature group, there was less seasonal change in CR  $B_{\max}$ ; values in the ambient temperature groups were significantly greater than the 10 °C groups in early April, the maximum values for  $B_{\max}$  (Fig. 1). Correspondingly, there was a smaller decrease in  $B_{\max}$  following the peak. In the study by Shrimpton et al. (2000), there was a change in temperature over the year, but the temperature regime did not follow a profile characteristic of natural systems. Temperature was warmer in the winter (mean greater than 7 °C in both years) and decreased during the spring. The effect of this thermal regime is not clear, but the seasonal decreases in CR  $B_{\max}$  were certainly protracted.

Although photoperiod is the main environmental factor to stimulate smolting, increases in water temperature can affect the timing of smolting. As assessed by peak gill  $\text{Na}^+, \text{K}^+$ -ATPase activity, smolting occurs several weeks earlier when rearing temperature is increased (McCormick et al., 1997). Smolting has also been linked to changes in temperature as the seasonal increase in gill  $\text{Na}^+, \text{K}^+$ -ATPase activity is more pronounced when temperature increases in conjunction with photoperiod (Muir et al., 1994). Other smolt related characteristics such as time of migration have also been found to be strongly associated with a temperature threshold (Jonsson and Ruud-Haansen, 1985). The increases in  $B_{\max}$  were not affected by absolute temperature changes or a threshold in temperature, but the seasonal increase in temperature is associated with the decline in  $B_{\max}$  at the end of the spring. The elevation of temperature to 10 °C at the start of Experiment 1, however, had no effect on  $B_{\max}$ . Whether the increase in temperature in the first experiment occurred too early in development for  $B_{\max}$  to be affected is not clear.

Seasonal changes in CR  $k_d$  were also observed in the present study consistent with findings of other studies on Atlantic salmon (Shrimpton and McCormick, 1998a; Shrimpton et al., 2000) and coho salmon (Shrimpton et al., 1994), but not in steelhead or rainbow trout (McLeese et al., 1994). Changes in CR  $k_d$  were correlated with gill  $\text{Na}^+, \text{K}^+$ -ATPase activity, unlike  $B_{\max}$  (Fig. 4). A strong correlation between these two variables has been seen in previous studies. Shrimpton and McCormick (1998a) found that highest  $k_d$  values were coincident with maximum gill  $\text{Na}^+, \text{K}^+$ -ATPase activity. Strong seasonal correlation was also seen in the study by Shrimpton et al. (2000). In coho salmon, a similar relationship was also found for fish captured in the wild, but not for hatchery fish reared at a constant temperature (Shrimpton et al., 1994). The seasonal increase in  $k_d$ , therefore, appear to be a function of physiological changes associated with smolting that also alter  $k_d$ .

Advances in photoperiod (Experiment 1) increased gill CR  $k_d$ , but not at low temperature (Fig. 1). The increase in  $k_d$  correlated well with an increase in gill  $\text{Na}^+, \text{K}^+$ -ATPase activity and in turn plasma GH (see McCormick et al., 2000). It is possible that the changes in  $k_d$  that respond to photoperiod are a function of endocrine changes associated with smolting. The short day treatments in Experiment 2 did not affect  $k_d$  (Fig. 2).

#### 4.2. Endocrine control of CR

In the present study, there were a number of correlations between endocrine parameters and  $B_{\max}$ . Individual regression of hormone data showed that there was a significant relationship between CR  $B_{\max}$  and  $T_4$ . This was not seen with  $T_3$ , which would appear contrary to the findings of Shrimpton and McCormick (1999) who showed that exogenous  $T_3$  significantly increased CR  $B_{\max}$ . This may be due to species differences or due to the activity of these two hormones. Seasonal increases in  $T_4$  is a normal feature of smolting, yet there appears to be little increase or change in  $T_3$  (Hoar, 1988). The correlation of gill CR  $B_{\max}$  with  $T_4$  and not  $T_3$ , therefore, may reflect deiodination of  $T_4$  in the gill tissue itself.

There was a weak, but positive correlation between GH and  $B_{\max}$ . Given the evidence from injection studies (Shrimpton et al., 1995; Shrimpton and McCormick, 1998b), it is surprising that the relationship between  $B_{\max}$  and GH is not stronger. GH has been shown to interact with cortisol to increase saltwater tolerance (Madsen, 1990). A number of mechanisms have been proposed to account for this finding. GH may have a direct action on the gill through gill GH receptors (Gray et al., 1990), GH may act to increase gill response to cortisol through an increase in CR  $B_{\max}$  (Shrimpton et al., 1995), or GH cause release of IGF-1 which may have a direct effect (Sakamoto et al., 1993). The hypoosmoregulatory action of IGF-1 on salmonids has been shown by McCormick et al. (1995) and an additive effect with cortisol has been shown by McCormick (1996). If the GH effect is through IGF-1, then we would expect that a strong correlation should exist between CR  $B_{\max}$  and IGF-1. Indeed a strong relationship exists between CR  $B_{\max}$  and IGF-1, but the correlation is negative. The reason for this apparent discrepancy is unclear particularly given the findings of McCormick (1996).

Interestingly, plasma cortisol was not significantly correlated with CR  $B_{\max}$ . A strong negative correlation between plasma cortisol and CR  $B_{\max}$  would be expected and has been demonstrated in several previous studies (Maule and Schreck, 1991; Pottinger et al., 1994; Shrimpton and Randall, 1994). The results of these studies, however, may not contradict the present findings as they were not conducted during the spring when the fish were smolting. During the spring, the seasonal increases in plasma  $T_4$  and GH may function to limit the decline in CR  $B_{\max}$  when cortisol also increases. Indeed, the multi-hormonal control of the gill cortisol receptor may explain why correlations with individual hormones are weak or contradictory.

In the present study, plasma cortisol, GH, IGF-1 and  $T_4$  were positively correlated with gill CR  $k_d$ . There was no relationship with  $T_3$ . Injection experiments with cortisol have shown an increase in CR  $k_d$  in the gills of coho salmon (Maule and Schreck, 1991; Shrimpton and Randall, 1994) and rainbow trout liver (Pottinger et al., 1994). GH injections increase  $k_d$

in Atlantic salmon (Shrimpton and McCormick, 1998a,b), but GH and  $T_3$  were not found to affect CR  $k_d$  in rainbow trout (Shrimpton and McCormick, 1999). Whether the correlation between IGF-1 and  $T_4$  are causal is not known. Further experimentation is required to determine a regulatory role for these two hormones on gill CR  $k_d$ .

Under natural temperature regimes CR  $B_{max}$  increased prior to elevations in gill  $Na^+,K^+$ -ATPase activity, while changes in  $k_d$  correlated with the seasonal increases in gill  $Na^+,K^+$ -ATPase activity. The change in  $B_{max}$  may be preparatory, as the gill will be more responsive to cortisol as part of the endocrine system driving higher activities of gill  $Na^+,K^+$ -ATPase and seawater tolerance. Higher temperatures will result in earlier development of salinity tolerance (McCormick et al., 1997). The present study, however, indicates that advances in the seasonal increase in water temperature will lead to a decrease in CR  $B_{max}$  with little change in CR  $k_d$ , reducing gill sensitivity to cortisol earlier in the year. Higher temperatures also result in more rapid losses of salinity tolerance and decreases in gill  $Na^+,K^+$ -ATPase activity (McCormick et al., 1997). The earlier decline in CR  $B_{max}$  in fish reared under temperature regimes that are advanced or warmer may be a mechanism for the shortened smolt window in fish reared under seasonally advanced or warmer water temperatures.

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