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Top-Down and Bottom-Up Control of Life-History Strategies in Coho Salmon (Oncorhynchus kisutch)

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ABSTRACT: Sexual maturation profoundly affects population dynamics, but the degrees to which genetic, top-down, and bottomup controls affect age at maturity are unclear. Salmonid fishes have plastic age at maturity, and we consider genetic and environmental effects on this trait by developing fitness functions for coho salmon (Oncorhynchus kisutch). The functions are based on size-specific survival and reproductive success, where reproductive success is the product of fecundity and ability to defend nests (females) or the product of sperm volume and ability to mate (males). We model genetic and bottom-up controls (e.g., food availability) with an environmentally explicit growth function and top-down control (predation mortality) with survival functions that consider both sizedependent and size-independent mortality. For females, we predict that early maturation rarely maximizes fitness, but males can maximize fitness by maturing early if they grow well in freshwater. We predict that early maturation is most affected by the bottom-up effects of resource distribution at sea, followed by bottom-up and genotypic effects in freshwater. Top-down processes are predicted to have strong effects on the likelihood of delayed maturation.

Keywords: reproductive fitness, freshwater, saltwater, genotype.

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Age at sexual maturity profoundly affects fitness and population dynamics. The advantages of maturing early (e.g., higher probability of surviving to reproduction) must be traded off with aspects that justify delaying maturation (e.g., maximizing fecundity through increased body size; Stearns 1992; Hendry and Stearns 2004). In species or populations that display variability in age at maturity, it is important to understand the mechanisms that influence it (Pyper et al. 1999). Expected growth, survival, and reproductive success are key factors affecting the evolution of age at maturity, and all three processes are influenced by environmental and genotypic variability (Morris and Ryan 1990; Rose et al. 2002).

In the environment, top-down and bottom-up processes can influence growth, survival, fecundity (Hunter and Price 1992; Quinn et al. 2004), and, hence, age at maturity. Within a food web, effects that propagate through mortality processes and are independent of resource (food) availability are considered top-down controls; when resource availability controls food-web dynamics, the system is considered to be under bottom-up control (Hunter and Price 1992). Elucidating the conditions under which top-down controls outweigh bottom-up controls (and vice versa) is a current topic of interest in the ecological literature (e.g., Menge 2000; Diaz-Pulido and McCook 2003; Munch et al. 2005), and here we focus on how these controls interact to influence age at maturity.

Salmonid fishes are a useful "test taxon" for developing an understanding of how ecosystem-control processes influence life histories. These fishes have amazingly diverse life histories, ranging from obligate freshwater through facultative anadromous to obligate anadromous (Stolz and Schnell 1991; Behnke 2002), which have evolved across a complex environmental background where top-down and bottom-up controls vary in space and time. Salmonids have variable age at maturity, and the plasticity of this trait may allow these fishes to accommodate the range of conditions they encounter in their freshwater and marine environments (Bisbal and McConnahan 1998). For the anadromous salmonids, various correlative studies have demonstrated links between ocean conditions, marine survival, and mean length

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of returning spawners (e.g., Cox and Hinch 1997; Pyper et al. 1999; Cole 2000; Hobday and Boehlert 2001; Mueter et al. 2002). Despite this valuable work, we lack a mechanistic understanding of how ecosystem dynamics, including both top-down and bottom-up effects, influence salmonid life history (Cooney et al. 2001) or age at maturity. This is surprising, given the importance of genotypic and phenotypic diversity to the viability of depleted salmon populations (McElhany et al. 2000; Watters et al. 2003).

In salmonids generally and in coho salmon (Oncorhynchus kisutch) specifically, male age at maturity is linked to reproductive tactic. Males that mature early typically sneak matings; conversely, older males usually fight to gain access to females (Gross 1985; Sandercock 1991). There are also distinct phenotypic differences: early-maturing males are small and cryptically colored and have poorly developed kypes (hooked jaws), while older males are much larger and more brightly colored and have well-developed kypes (Sandercock 1991). Furthermore, individual coho that grow best in freshwater are most likely to mature early and use the sneaking tactic (Garrison 1971; Hager and Noble 1976). One can envision switch points that are related to growth performance in freshwater; on either side of such switch points, fitness might be maximized by different life histories (e.g., maturing early and sneaking matings vs. maturing late and fighting; Gross 1996). For coho, growth performance in freshwater can be described either by length at the smolt transformation (the transformation that occurs to prepare salmonids for the migration from freshwater to saltwater; e.g., Gross 1996) or by a measure of intrinsic growth potential (i.e., the rate at which parr grow toward the maximum smolt length; see Snover et al. 2005). Interestingly, exceptional growth performance in freshwater may decrease growth potential at sea because the behaviors that confer feeding advantages to individuals in freshwater may not be effective in saltwater (Jonsson and Jonsson 1993; Snover et al. 2005). Since the behaviorenvironment interaction is abruptly altered when salmonids migrate from freshwater to saltwater, reproductive tactics and their links to switch points should also be considered in the context of growth potential at sea.

For coho salmon and probably other salmonids as well, age at maturity will probably vary in response to genotypeby-environment interactions that occur in both freshwater and saltwater. Variation in age at maturity occurs both between and within coho populations (Sandercock 1991). Variability between populations is driven by environmental differences that occur over the geographic range of the species and genetic differences that are perpetuated by the homing instinct (Silverstein and Hershberger 1995; Quinn et al. 2001b). Variability within populations is driven by environmental effects on the relative performance, in terms of growth, survival, and fecundity, of different phenotypes (Watters et al. 2003). To our knowledge, longterm longitudinal studies that track the genetic and environmental histories of individual coho and relate these histories to age at maturity and reproductive success are not available (note, however, the relevant longitudinal studies on Atlantic salmon Salmo salar, e.g., Letcher and Gries 2002). Fortunately, however, data on coho salmon are plentiful, and this species is therefore still a useful "test species" for developing a modeling approach that can be used to untangle genotype-by-environment interactions and predict their effects on age at maturity.

Here, we seek to identify how genotype-by-environment interactions are predicted to influence the age at which coho salmon mature. We derive size-specific fitness functions for both males and females, using the growth model of Snover et al. (2005), length-at-age data from Shapovalov and Taft (1954), and parameters from other literature sources. We use the fitness functions to consider the interplay of genetic, top-down, and bottom-up controls in determining the likelihood that coho salmon mature early or delay maturation. We show that the relative frequencies of phenotypes (ages at maturity) that arrive on the spawning grounds are controlled by both bottom-up and topdown processes. The former processes affect early maturation, and the latter affect late maturation. Ultimately, however, an individual's genotype acts as the template on which these processes act, and thus, we posit that gametheoretic modifications to the effective transmission of genotypes between generations may create an interesting feedback loop.

Methods

We define fitness as the expected lifetime reproductive success at a given age at maturity (corresponding to a stationary population), and we model fitness as a function of genetic, bottom-up, and top-down processes. We account for the former two processes in a growth model, and we account for the latter process in a survival model. Note that we do not include negative frequency dependence (i.e., reductions in fitness when the spawning population is dominated by a particular age at maturity) in our fitness equations. Our focus is on how environmental factors, which may be more important to determining reproductive tactic than genetics (Aubin-Horth and Dodson 2004), influence age at maturity; this is a different topic than that typically addressed by game theory.

Individual Growth

The reproductive components of fitness (e.g., for females, egg biomass and ability to defend a nest; for males, gonad size and ability to achieve matings) correlate strongly with adult body size (van den Berghe and Gross 1984, 1986, 1989; Gross 1985), and therefore we model adult length at a given age at maturity using the growth model developed by Snover et al. (2005). This model facilitates consideration of both genetic and bottom-up effects on age at maturity. It is based on the von Bertalanffy growth function (VBGF):

$$\frac{dL}{dt} = E - kL,\tag{1}$$

where L is length, t is time, E is a coefficient of anabolism, and k is a coefficient of catabolism (von Bertalanffy 1938). In applying this growth model to our fitness functions, we assume that E relates to bottom-up factors and that k relates to both the genotypic and phenotypic capacity for growth (we discuss this decomposition later). The solution of equation (1) is

$$L_t = L_{\infty} - (L_{\infty} - L_0) \exp(-kt), \qquad (2)$$

where L_t is length at time t, L_0 is an initial length, and L_{∞} is the asymptotic size at which growth rates are 0. The asymptotic size is a function of E and k (see appendix for further discussion of these parameters):

$$L_{\infty} = \frac{E}{k}.\tag{3}$$

Snover et al. (2005) applied the VBGF to coho salmon by modeling three growth stanzas (one stanza in freshwater and two stanzas in saltwater). Here, we consider four growth stanzas, with the additional stanza being an additional growing season in saltwater (two sea winters). The four growth stanzas are temporally separated by the smolt transformation and the first and second sea winters, and they approximate Sandercock's (1991) description of the predominant life cycle of these fish (see appendix for a detailed description).

The amount of resources available to an individual fish depends on its genotype, its phenotype, and bottom-up environmental forces. The parameter E, or resources available to an individual, is not solely a measure of the total amount of food produced from bottom-up forcing because behavior interacts with resource distribution to determine the amount of food available to an individual. Behavior is associated with k (discussed below), and therefore it seems appropriate to let E be a function of k (Snover et al. 2005; see appendix for further discussion of eq. [4]):

$$E = \gamma \cdot k^{\psi}, \tag{4}$$

where the parameter k is the coefficient of catabolism from equation (1); it also (from eq. [2]) determines the rate at which L_t approaches L_{∞} . We assume that an individual's k is determined both genetically and by environmental conditions that influence an individual's behavioral phenotype and occur early in life (e.g., during the egg and alevin stages). One can envision a partition of these genetic and early environmental effects by assuming that k is a random variable whose expectation—say, k—is genetically determined with individual deviations from this mean that describe the diversity of behavioral phenotypes derived from early environmental experiences (see the last paragraph of this subsection and "Fitness and Age at Maturity"). Snover et al. (2005) provide considerable discussion about possible links between k and behavior, but generally, coho with larger k are considered to be more aggressive (appendix). Hence, we follow their model and make the assumption that *k* is constant throughout life, and thus, the length of an individual at some time after its emergence from the gravel is conditioned on k. The parameters γ and ψ determine the degree to which E depends on bottomup factors: γ is a scalar >0 that is constant across individuals and determines overall resource abundance or density; ψ is a parameter that both describes the distribution of resources and tempers the influence that *k* (effects from genetics and behavior) has on an individual's ability to sequester resources. Following Snover et al. (2005), we constrained ψ to be in the interval [0, 1]. When $\psi = 0$, resources are considered to be homogeneously distributed through the environment, and individuals do not get increased access to resources through aggressive behavior. When $\psi = 1$, resources have a clumped distribution, and aggressive behaviors can be used to increase access to them. Intermediate values of ψ provide intermediate interactions between resource distribution and behavior.

For growth in freshwater, we followed Snover et al. (2005) and considered only $\psi = 1$ in this habitat:

$$E_{\rm FW} = \gamma_{\rm FW} \cdot k, \tag{5}$$

where the subscript FW indicates freshwater. Combining equations (2), (3), and (5) provides an equation for predicting length in freshwater y months after emergence from the gravel, given k:

$$L_{\text{FW}}(y|k) = \gamma_{\text{FW}} - (\gamma_{\text{FW}} - L_f) \exp(-ky), \tag{6}$$

where $L_{\rm f}$ is fry length at emergence from the gravel, and, given equations (3) and (5), the scalar $\gamma_{\rm FW}$ is equal to the asymptotic length at the smolt transformation. We use t to denote time generally (e.g., eq. [2]), y to denote time in freshwater (e.g., eq. [6]), and z to denote time in saltwater (e.g., eq. [8]).

For growth at sea, we also followed Snover et al. (2005) and considered the full diversity of predictions offered by equation (4):

$$E_{\rm sw} = \gamma_{\rm sw} \cdot k^{\psi}. \tag{7}$$

The subscript SW denotes saltwater.

As noted previously, Snover et al. (2005) divided growth in saltwater into two stanzas, with E_{sw} changing after the first winter at sea. Here, we expand this to three saltwater stanzas and include consideration of the two-sea-winter life history (4-year-old spawners; appendix). To observe general patterns in the fitness functions we develop here, we assume that E_{sw} is constant during the entire portion of life spent at sea (appendix). Combining equations (2), (3), and (7) provides an equation for predicting length at z months after the smolt transformation for a fish in saltwater:

$$L_{\text{SW}}(z|k) = \frac{E_{\text{SW}}}{k} - \left[\frac{E_{\text{SW}}}{k} - L_{\text{FW}}(12|k)\right] \exp(-kz),$$
 (8)

where $L_{\text{FW}}(12 \mid k)$, computed from equation (6), is the length at the smolt transformation.

Snover et al. (2005) identified parameter values for the growth model described in equations (5)-(8) by comparing predicted lengths at age to observations presented in Shapovalov and Taft (1954), and we applied those parameter values here. We allowed k to vary among individuals and range from 0.03 to 0.18, which corresponds to smolt lengths ranging from about 7 to 16 cm. We achieved this by drawing random values of individual k from gamma distributions with mean values $k = \{0.06, 0.07, 0.08\}$ and coefficients of variation = 0.30. For growth in freshwater, we set $L_{\rm f}=2$ cm and $\gamma_{\rm FW}=18$ cm. For growth at sea, we explored values from the set $\psi = \{0, 0.15, 0.25, 0.35,$ 0.5, 0.75, 1}. We paired values of γ_{SW} with values of ψ , using the equation $\overline{E}_{\rm SW} = \gamma_{\rm SW} \cdot \vec{k}^{\psi}$, and considered mean levels of k from the set $k = \{0.06, 0.07, 0.08\}$ with mean levels of E_{SW} from the set $\overline{E}_{SW} = \{6.2, 6.4, 6.6\}$. Thus, for a given value of ψ , variation in \overline{E}_{SW} (a summary parameter we use to simplify the presentation of our results) has the same interpretation as γ_{sw} . Note, however, that individual fish realized individual levels of E_{SW} because we used the pairs $\{\psi, \gamma_{SW}\}\$ and individual values of k in equation (7).

Survival

We consider survival to time t, S(t), in both freshwater and saltwater, to be size dependent. The general form of the equation is

$$S_{i}(t) = \prod_{x=1}^{t-1} \exp \left[-m_{0,i} + \frac{m_{1,i}}{L_{i}(x|k)} \right], \tag{9}$$

where *i* indexes habitat (freshwater or saltwater). Time, *t*, goes from 1 to 12 in freshwater and from 1 to 6, 18, or 30 in saltwater. The parameter m_0 is a size-independent mortality term, and the parameter m_1 adjusts the sizedependent component of mortality. Salmon are subjected to size-dependent mortality in the ocean (McGurk 1996), but the proportion of mortality that is attributable to sizedependent versus size-independent sources is unknown. To encompass a range of possibilities, we partition mortality between the size-dependent and size-independent terms. For both freshwater and saltwater, we used 10 pairs of m_0 and m_1 values such that the amount of mortality attributable to size-independent factors varied between about 1% and 100% while overall survival was held constant (table 1). Each pair of m_0 and m_1 values resulted in total survival of approximately 1.8% from emergence to outmigration and about 8% from outmigration to spawning after one winter at sea. We chose 8% because it is intermediate to the values reported in the literature (Shapovalov and Taft 1954; McGurk 1996; Coronado and Hilborn 1998; see table 1). We also considered total survival of 6% and 10% from outmigration to spawning after one

Table 1: Values of size-independent (m_0) and sizedependent (m_1) terms used in the survival equation

			Proportion of		
			mortality		
			attributable to		
			size-independent		
	m_0	m_1	factors		
Freshwater	.001	1.980	.01		
	.010	1.930	.12		
	.020	1.870	.22		
	.030	1.810	.31		
	.040	1.750	.39		
	.060	1.630	.52		
	.075	1.540	.60		
	.100	1.390	.71		
	.150	1.095	.85		
	.334	.001	.00		
Saltwater	.001	5.600	.02		
	.005	5.400	.09		
	.010	5.200	.18		
	.020	4.750	.33		
	.025	4.600	.39		
	.035	4.200	.51		
	.048	3.700	.63		
	.060	3.200	.72		
	.075	2.600	.81		
	.140	.001	1.00		

sea winter to determine the effect of changes in overall survival on age at maturity. These values were based on a smolt length of 11 cm and vary somewhat for smolts of different lengths. To simplify the presentation of our results, we focus on the relative contributions of size-independent and size-dependent mortalities to total mortality (rather than specific values of m_0 and m_1) from here forward. For simplicity, we assumed that the habitat-specific partitioning of size-independent and size-dependent mortalities was constant throughout an individual's entire time in freshwater or at sea.

Sex-Specific Fitness Functions

Females. In female coho salmon, there is a positive relationship between body size and fecundity (van den Berghe and Gross 1989; Quinn et al. 2004). Specifically, van den Berghe and Gross (1989) described the relationship between total egg biomass, g, and body length at maturity, L^* , as

$$g(L^*) = 3.88 \times 10^{-4} (L^*)^{3.41}.$$
 (10)

Van den Berghe and Gross (1989) present data on the relationship between the percentage of nests still intact at the end of the breeding season and female length. In particular, the probability of nest survival (S_{nest}) given female length at maturity, L^* , is

$$p(S_{\text{nest}}|L^*) = \min[(1.47L^* - 28.26)100^{-1}, 1],$$
 (11)

such that $p(S_{\text{nest}}|L^*) = 1$ for $L^* > 87$ cm (this is a very large size for coho salmon and rarely occurs). Note also that $L^* < 28.26/1.47$ cm is also very rare for a mature female.

We computed the length-specific fitness of females that inherit a specific k and mature at length L^* after spending z months at sea, $\omega_{\rm f}(L^*, z)$, as the product of survival to adulthood (eq. [9]), fecundity (eq. [10]), and the probability of nest survival (eq. [11]):

$$\omega_{\rm f}(L^*,z) = S_{\rm FW}(12) \cdot S_{\rm SW}(z)$$
$$\cdot g[L_{\rm SW}(z|k)] \cdot p[S_{\rm nest} \mid L_{\rm SW}(z|k)]. \quad (12)$$

All four terms in equation (12) are size specific and thus involve genetic and bottom-up effects on fitness. The two survival terms also describe top-down effects on fitness. We substituted $L_{\text{SW}}(z|k)$ for L^* in equations (10) and (11), where z=6, 18, or 30 months.

Males. Male coho use two reproductive tactics, sneaking and fighting (Gross 1985). Two important components of the successful use of either tactic are body size and frequency with which each tactic is used on the spawning

grounds (Gross 1985; Repka and Gross 1995). Previous models based on game theory have assumed that phenotypic variation in reproductive tactics is maintained by negative frequency dependence (e.g., Hutchings and Myers 1994; Repka and Gross 1995). Here, our focus is on how environmental conditions in both freshwater and saltwater influence development (growth) and reproductive tactic. We argue that decisions regarding which reproductive tactic will be used must be made before gaining any knowledge of the social structure (relative frequency of individuals employing each tactic) on the spawning ground, and so in this exercise we do not consider frequency dependence in the success of reproductive strategies. Our work complements the application of game theory; we predict the distribution of phenotypes in a population, while game theory can predict the distribution of genotypes.

We used gonadosomatic indexes (GSIs) to describe a component of male reproductive potential and estimate a proxy of sperm volume, although the influence of sperm volume on final fitness will be small compared to that of access to mating opportunities facilitated by body size. In general, male salmonids that mature early invest about 37%-63% more in relative gonad mass than do males that mature later (table 2). The cultured males observed by Bessey et al. (2004) are phenotypically similar to wild, early-maturing individuals, and therefore we used GSI values from Bessey et al. (2004) to develop our index of sperm volume, V(z):

$$V(z) = (L^*)^3 \cdot GSI(z), \tag{13}$$

where z is months at sea. For z = 6, we used the GSI for males that mature early; for z = 18 or 30, we used the GSI for males that mature late (table 2).

The reproductive potential of male coho is also influenced by the relationship between body size and the tactic-specific ability to achieve mating opportunities. Large body size is beneficial to the fighting tactic but not to the sneaking tactic (Gross 1985). Using least squares, we fitted logistic functions to the tactic-specific data from Gross (1985) that relates male length to proximity to nesting females. We assumed that reproductive success is proportional to proximity and estimated the probability of successfully mating given use of the sneaking tactic and length at maturity as

$$p(M|\text{sneak}, L^*) = 0.77[1 + \exp(0.45L^* - 17.44)]^{-1}.$$
 (14)

We estimated the probability of successfully mating given use of the fighting tactic and length at maturity as

$$p(M|\text{fight}, L^*) = \exp(0.21L^* - 10.48)$$

 $\cdot [1 + \exp(0.21L^* - 10.48)]^{-1}.$ (15)

Table 2: Gonadosomatic indexes (GSIs) for early- and late-maturing salmonids

		•	•			
Source	Species	Description early	Description late	GSI early	GSI late	GSI late/ GSI early (%)
Bessey et al.		•		,		
2004	Coho	3–4-year-old cultured males; phenotypically jacks	3-year wild	.070 ± .010 (SD)	.040 ± .010 (SD)	57.1
Vladić and Järvi		1 11 17				
2001	Atlantic	Mature parr	Anadromous males	$.110 \pm .039 (SD)$	$.041 \pm .011 \text{ (SD)}$	37.1
Gage et al.						
1995	Atlantic	Mature parr	Anadromous males	$.047 \pm .004 \text{ (SE)}$	$.023 \pm .0024 \text{ (SE)}$	50.1
Foote et al.						
1997	Sockeye	Jacks	Males over 525 mm	.043	.021	48.8
Koseki and						
Maekawa						
2002:						
Shikaribetsu Lake	Masu	Mature parr	Anadromous males	$.106 \pm .026 \text{ (SD)}$	$.067 \pm .016 \text{ (SD)}$	63.2
Shumarinai						
Reservoir	Masu	Mature parr	Anadromous males	$.131 \pm .031 \text{ (SD)}$	$.064 \pm .015 \text{ (SD)}$	48.9

Note: Variability is reflected as either standard deviation (SD) or standard error (SE), depending on what is provided in the source document. Species listed include coho salmon (Oncorhynchus kisutch), Atlantic salmon (Salmo salar), sockeye salmon (Oncorhynchus nerka), and masu salmon (Oncorhynchus masou).

In equation (14), z = 6, and in equation (15), z = 18 or

We computed the length-specific fitness of male coho that inherit k and mature after z months at sea as the product of survival (eq. [9]), sperm volume (eq. [13]), and ability to achieve mating opportunities (eqq. [14], [15]). The resulting fitness functions for the sneaking tactic and the fighting tactic are

$$\omega_{\rm m}(L^*, z | {\rm sneak}) = S_{\rm FW}(12) \cdot S_{\rm SW}(z)$$

$$\cdot V(z) \cdot p[M | {\rm sneak}, L_{\rm SW}(z | k)] \quad (16)$$

and

$$\omega_{\rm m}(L^*, z|{\rm fight}) = S_{\rm FW}(12) \cdot S_{\rm SW}(z)$$
$$\cdot V(z) \cdot p[M|{\rm fight}, L_{\rm SW}(z|k)], \quad (17)$$

respectively. Similar to the fitness functions for females, all eight terms on the right sides of equations (16) and (17) describe genetic and bottom-up effects on fitness, while the four survival terms also describe top-down effects. We considered scenarios for use of the fighting tactic (eq. [17]) for z = 6, or maturation after 6 months at sea, and use of the sneaking tactic (eq. [16]) for z = 18 or 30, or maturation after 18 or 30 months at sea. Under these conditions, the resulting fitness curves were at or near 0 and were always less than values achieved when z = 6 in equation (16) (use of the sneaking tactic by males that

mature early) and when z = 18 or 30 in equation (17) (use of the fighting tactic by males that mature late). Hence, from this point forward we consider only z = 6in equation (16) and z = 18 or 30 in equation (17).

Fitness and Age at Maturity

We evaluated the fitness functions with a variety of parameter combinations; we used seven values of ψ , three values of \overline{E}_{sw} , three distributions of k, and 10 survivorship schedules each in freshwater and saltwater. We simulated the survival and maturation of 200,000 fry in each simulation. Each fry was randomly assigned a k value from one of the three gamma distributions. Note again that we interpret individual variation in k as phenotypic variation and variation in \overline{k} as genotypic variation. Additionally, we did not explore variation in \overline{k} to make inferences about most-fit genotypes; rather, we aimed to infer the relative degree of genetic control over variation in age at maturity. Growth was simulated on a monthly time step using equations (5)-(8), and mortality was simulated by drawing random numbers, one for each individual, from a U(0, 1) distribution and comparing these random numbers to the monthly survival rates predicted by equation (9). Using equations (12), (16), and (17), we determined the optimal age at maturity for each individual by computing its fitness, conditioned on survival, after 6, 18, and 30 months at sea. The time at sea that maximized fitness defined the optimal age at maturity, where age at maturity = 6 months for egg E146

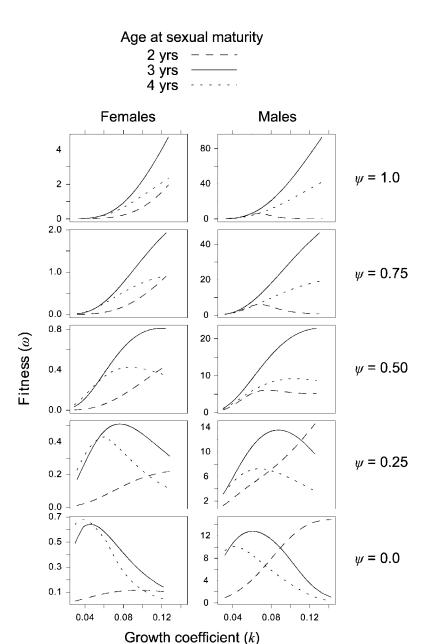


Figure 1: Fitness curves for males and females at five values of ψ . The value of \overline{E}_{sw} was 6.4, and the proportion of mortality attributable to size-independent factors was \approx 50% in both freshwater and saltwater. Note that these curves are depicted by connecting point estimates of age-specific fitness from many individuals with values of k ranging from 0.03 to 0.18. Also note that the Y-axes are independently scaled.

incubation + 12 months for growth in freshwater + number of months at sea that maximized fitness. Throughout the remainder of this article, we present detailed results for simulations in which overall survival at sea was approximately 0.08; we also consider overall survival rates of about 0.06 and 0.10 and note the general effects of such changes in "Results."

Results

Fitness curves at $\psi = 1.0$, 0.75, 0.5, 0.25, and 0 (those values considered by Snover et al. [2005]) using intermediate survival schedules where size-independent mortality accounted for about 50% of total mortality are illustrated in figure 1. Early maturation maximized fitness

for large male smolts only when $\psi < 0.5$. Similarly, for females, delaying maturation and spawning at 4 years of age maximized fitness for small smolts only when ψ < 0.5. For both sexes, $\psi \ge 0.5$ predicted that fitness would always be maximized by maturing at 3 years old. Since coho salmon do not strictly mature at 3 years of age (Sandercock 1991), we limit our presentation of all further results to those from simulations in which ψ < 0.5. Figure 1 also illustrates how switch points based on growth performance in freshwater (i.e., switch points based on k) were predicted to be influenced by environmental conditions in the ocean. The values of k at which fitness switched from being maximized at one age at maturity to the next age at maturity varied among panels.

For males, the bottom-up effects of resource density $(\overline{E}_{sw};$ increases across rows in fig. 2) and resource distribution (ψ ; decreases down columns in fig. 2) had greater impacts on the likelihood of early maturation than did the top-down effects of mortality in either freshwater or saltwater (fig. 2). The spatial distribution of the resources (ψ), also interpreted by Snover et al. (2005) as the effectiveness of aggressive behavior, had the greatest influence on the likelihood of early maturation by males (fig. 2; compare down columns). The abundance of resources (\overline{E}_{SW}) also had an effect on the likelihood of early maturation, although the effect was not as dramatic as that of resource distribution (fig. 2; compare across rows). In general, early maturation was predicted to be less likely when resources were more available at sea. Early maturation was also predicted to be less likely when overall survival in saltwater was increased. From the diagonal nature of the contours illustrated in figure 2, it appears that top-down factors in freshwater and saltwater were approximately equivalent in their effects on early maturation. Nevertheless, increasing the proportion of size-independent mortality in freshwater decreased the likelihood of early maturation, while in saltwater, it increased that likelihood. Overall, however, the effects of both top-down controls were much less than those of the two bottom-up controls.

The predictions for females were different (fig. 3). Both of the bottom-up factors at sea (resource abundance and distribution) had little impact on the proportion of a cohort delaying maturation to 4 years. The same could be said for top-down effects in freshwater. Top-down control in saltwater, however, strongly influenced the likelihood of delayed maturation. Decreases in size-independent mortality (or increases in size-dependent mortality) lead to increased age at maturity for females. Increases in overall survival in saltwater also increased the likelihood of delayed maturation.

Genotypic and freshwater bottom-up effects on the likelihood of early and late maturation were intermediate to the bottom-up and top-down effects identified in the previous paragraphs (figs. 4, 5). Increasing mean k (equivalent to increasing smolt length) increased the likelihood of early maturation (fig. 4). Likewise, the likelihood of delayed maturation was reduced with increased k (smolt length;

Discussion

For the management of Pacific salmon populations, it is important to understand how survival and ecosystem productivity can modify age at maturity because variability in this trait affects the age composition of adults on the spawning grounds and the amount of genetic mixing between cohorts. Genetic diversity is a key component of viability in salmon populations (McElhany et al. 2000). There are three views on how conditions in freshwater and saltwater interact to control salmon abundance (Bisbal and McConnahan 1998). The first is that conditions in freshwater alone control abundance; the second is that variability in ocean conditions accounts for most of the variation in abundance. The third, more recent view is that salmon abundance is influenced by the combination of conditions in both habitats (Bisbal and McConnahan 1998). Our results support the third view; the likelihoods of both early and late maturation are influenced by processes that occur in freshwater and at sea. Interestingly, however, we predict that processes at sea may have more influence on maturation than genetics and processes in freshwater, but the latter effects are consistent, whereas the relative influences of effects at sea change over time. Genetic and freshwater effects appear to form a template that constrains the outcomes of processes at sea. Here, we focus on relatively simple, idealized life histories of coho salmon, considering only 1 year of freshwater residency, followed by an obligate migration to sea. Applications of this model to different life histories with longer freshwater residencies or life cycles spent entirely in freshwater may yield different results.

Age at Maturity in Males

A striking prediction of our model is that for $\psi \geq 0.5$, the sneaking tactic, with maturation at 2 years of age, never results in higher overall individual fitness (fig. 1). It is only when $\psi \le 0.5$ that maturation at 2 years (sneaking tactic) results in higher overall fitness for large smolts, while medium-sized smolts are always predicted to mature at 3 years (fighting tactic). These results are consistent with empirical studies on the relationship between smolt length and age at maturity by coho salmon (e.g., Vøllestad et al. 2004). Snover et al. (2005) demonstrated that growth potential at sea increases with increasing ψ for large smolts. Hence, the predictions demonstrated in figure 1 support

Contoured probabilities that male coho salmon will mature early and reproduce as 2-yr olds

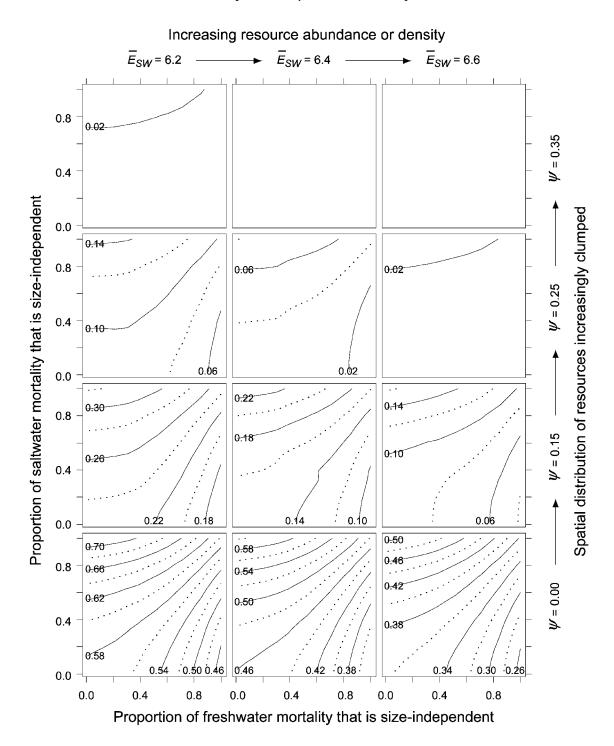


Figure 2: Individual contour plots show how the proportion of size-independent mortality in both freshwater and saltwater influences the percentage of males maturing early at 2 years of age. In the matrix of plots, $\overline{E}_{\rm SW}$ increases across rows, and ψ decreases down columns to demonstrate the potential impacts of bottom-up effects in saltwater on the likelihood of early maturation. For all cohorts, $\bar{k}=0.07$.

Contoured probabilities that female coho salmon will delay maturation and reproduce as 4-yr olds

Increasing resource abundance or density $\overline{E}_{SW} = 6.4$ $\overline{E}_{SW} = 6.6$ 0.02 8.0 0.4 Proportion of saltwater mortality that is size-independent Spatial distribution of resources increasingly clumped 0.0 8.0 0.420.50 0.0 8.0 0.420.50 0.66 0.0 8.0 0.4 0.0 0.0 0.0 0.4 8.0 0.4 8.0 0.0 0.4 8.0 Proportion of freshwater mortality that is size-independent

Figure 3: Individual contour plots show how the proportion of size-independent mortality in both freshwater and saltwater influences the percentage of females delaying maturation until 4 years of age. In the matrix of plots, $\overline{E}_{\rm SW}$ increases across rows, and ψ decreases down columns to demonstrate the potential impacts of bottom-up effects in saltwater on the likelihood of delayed maturation. For all cohorts, $\bar{k}=0.07$.

Contoured probabilities that male coho salmon will mature early and reproduce as 2-yr olds

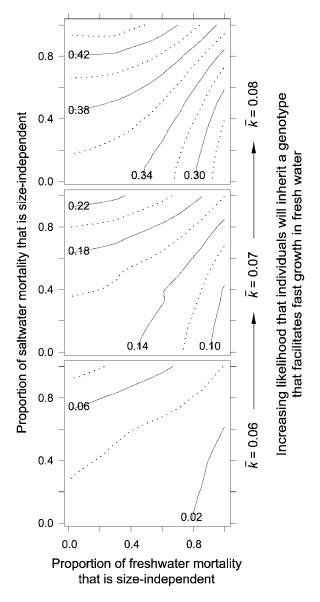


Figure 4: Individual contour plots show how the proportion of size-independent mortality in both freshwater and saltwater influences the percentage of males maturing early at 2 years of age. The value of \bar{k} increases from 0.06 to 0.08 up the column of plots, demonstrating the impact of genotypic and bottom-up environmental effects in freshwater. In all plots, $\bar{E}_{\rm SW}=6.4$ and $\psi=0.15$.

the theory that large smolts mature early as a result of decreased growth potential at sea (Jonsson and Jonsson 1993; Snover et al. 2005).

The results of our model also highlight that environmental conditions alone are enough to maintain the two reproductive strategies within a population. In all of the scenarios presented in figure 2, both reproductive tactics are maintained within cohorts, although at the extreme ranges ($\psi = 0$), sneaking frequencies were higher and possibly unrealistic. Such high frequencies would probably not occur, and here is where it would be important to consider negative-frequency-dependent issues.

Resource distribution at sea, a bottom-up process, is predicted to have the greatest influence on the likelihood of early maturation by males. The parameter ψ describes both the distribution of resources and the effectiveness of behaviors (e.g., aggression) individuals use to acquire those resources (Snover et al. 2005). Over the range of ψ values we considered, the likelihood of early maturation in males changed by an order of magnitude. Males were most likely to mature early when $\psi = 0$, when there is equal access to resources regardless of size or behavior and large smolts have reduced growth potential at sea (Snover et al. 2005). When resources are aggregated and more defensible $(0 < \psi \le 1)$, growth potential for large smolts increases (Snover et al. 2005), and here our model predicts decreased rates of early maturation. Access to food at sea, modulated through resource distribution (e.g., dispersed vs. clumped resources) and behavior, affects fitness by influencing the trade-off between the likelihood of future reproductive success and survival. If food is accessible, it seems profitable to risk mortality and stay in the ocean. The converse also applies; if food is not accessible, the risk of mortality is not worthwhile.

Vøllestad et al. (2004) found a negative relationship, as did we, between marine growth potential and the proportion of male coho that mature early, although no relationship was detected for chinook salmon (Oncorhynchus tshawytscha). Vøllestad et al. (2004) indicate that their results contrast with the general theoretical principle of increased growth rates leading to earlier age at maturity (Day and Rowe 2002). Our model shows how the results of Vøllestad et al. (2004) can actually be congruent with the theory discussed by Day and Rowe (2002). Increased growth rates in freshwater lead to younger age at maturity by decreasing growth potential at sea (Jonsson and Jonsson 1993; Snover et al. 2005). It should be noted that the populations studied by Vøllestad et al. (2004) are from a hatchery and are therefore somewhat artificial. However, other studies have demonstrated negative relationships between smolt length and net sea growth (Shapovalov and Taft 1954; Hager and Noble 1976; Holtby and Healey 1986; Mathews and Ishida 1989). Most of these relationships were not highlighted in the respective articles, but the data were summarized by Snover et al. (2005).

The second most important factor influencing the likelihood of early maturation by males is k, which represents both genotypic and bottom-up factors in freshwater. An

Contoured probabilities that female coho salmon will delay maturation and reproduce as 4-yr olds

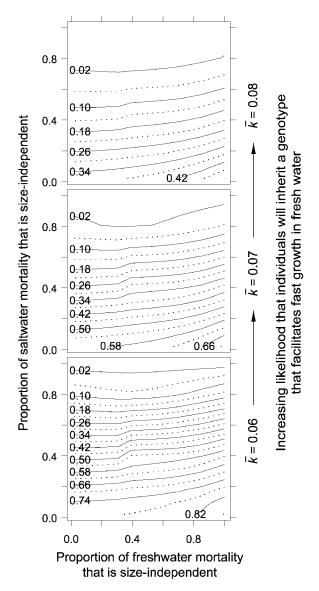


Figure 5: Individual contour plots show how the proportion of sizeindependent mortality in both freshwater and saltwater influences the percentage of females delaying maturation until 4 years of age. The value of \bar{k} increases from 0.06 to 0.08 up the column of plots, demonstrating the impact of genotypic and bottom-up environmental effects in freshwater. In all plots, $\overline{E}_{\rm SW}=6.4$ and $\psi=0.15$.

increase in k from 0.06 to 0.08 resulted in nearly an order of magnitude increase in rates of early maturation. This prediction is consistent with the literature on coho salmon, which indicates that for males, larger smolts are more likely to mature early (Garrison 1971; Hager and Noble 1976).

Resource density at sea, another bottom-up effect, seems likely to have a small, negative effect on the likelihood of early maturation. Over the range of E_{SW} values considered here, the likelihood that males would mature early approximately doubled; \overline{E}_{sw} was negatively related to the likelihood of early maturation, suggesting that improved conditions at sea are predicted to result in lower likelihoods of early maturation. When resources are less dense or abundant overall, growth rates and, hence, survival rates are lowered. Correlative studies suggest that "poor" environmental conditions in the ocean do result in decreased salmon survival (Cole 2000; Hobday and Boehlert 2001). Cole (2000) found that coho survival was decreased when upwelling was reduced and sea-surface temperatures were increased. Cole (2000) attributed decreased survival to reduced food availability, increased metabolic costs, and increased predation. Hobday and Boehlert (2001) found that a deep mixed layer results in decreased survival. They suggested that deep mixed layers result in lower plankton densities. Thus, it appears that the effect of resource density (abundance) acts in a manner similar to that of resource distribution. More abundant resources make it worthwhile to risk future mortality in the ocean because the potential reproductive payoff is great. Less abundant resources do not make the risk worthwhile.

Top-down factors are predicted to have a much smaller influence on the likelihood of early maturation than bottom-up factors. Nevertheless, when $\overline{E}_{\rm SW}$ and ψ were constant, the likelihood of early maturation was as much as doubled (or halved) by changing the mix of sizeindependent and size-dependent mortalities in both habitats. The combination of mortality terms most representative of natural systems is difficult to know. It is likely that the sources of size-independent and size-dependent mortality vary regionally and over time within a region. The opposite effects of mortality sources in freshwater and saltwater (e.g., increasing size-independent mortality in freshwater and decreasing it in saltwater to decrease the likelihood of early maturation) seem, again, related to growth potential at sea.

Age at Maturity in Females

Contrary to our findings for early maturation by males, top-down processes at sea are important for females delaying maturation to 4 years. Over the range of saltwater mortality schedules we considered, the likelihood of delayed maturation by females changed by more than an order of magnitude. At low levels of size-independent mortality, there is little difference between total survivorship after two growing seasons in comparison to three growing seasons, but the gains in body size are substantial. Female fecundity increases with length (van den Berghe and Gross 1989), and for males, increased length corresponds to an increased probability of securing mating opportunities using the fighting tactic (Gross 1985). The mortality risk associated with delaying maturity and becoming a 4-year-old spawner may be worthwhile if the same process, growth, simultaneously acts to increase reproductive potential and survival (with the latter effect being substantive only if size-independent mortality is low).

Although female coho do return as 4-year-old spawners, it is interesting to speculate why this age at maturity is not more common. Mortality rates, specifically the proportions of size-independent mortality, may be different between the sexes. Studying coho, Spidle et al. (1998) found that while the sex ratio of smolts outmigrating from Big Beef Creek, Washington, did not differ significantly from 1:1, the sex ratio between same-aged adults on the spawning ground was strongly biased toward males. Holtby and Healey (1990) suggest males may be more risk averse at sea, resulting in higher survival rates but potentially lower growth rates. In contrast, risk-prone behaviors by females may increase the proportion of mortality attributable to size-independent sources acting on this sex and thereby decrease the likelihood of delayed maturation.

Top-down controls on salmon production have been observed in the field. Willette et al. (2001) observed top-down control of juvenile pink salmon (*Oncorhynchus gorbuscha*) that resulted from prey switching by predators. When zooplankton resources were abundant, the primary predators of pink salmon fry, herring (*Clupea pallasi*) and walleye pollock (*Theragra chalcogramma*), fed heavily on the zooplankton, and predation on pink salmon was minimized. When zooplankton abundance fell below a threshold, herring and pollock switched and preyed more heavily on pink salmon fry.

Smolt Length and Fitness

Our predictions relating growth performance in freshwater to fitness and age at maturity are robust to environmental variability. Under all conditions with $\psi \leq 0.35$, large smolts (individuals with the largest values of k) maximize fitness by maturing early, and small smolts (individuals with the smallest values of k) maximize fitness by delaying maturation. For the middle range of smolt lengths, maturing at 3 years (which is the most common age at maturity for coho salmon) maximizes fitness. Snover et al. (2005) assessed the potential values of ψ for coho salmon. They concluded that $0 \leq \psi \leq 0.5$ during the first growing season at sea, but in subsequent growing seasons, ψ may increase (such an increase might, for example, result from an increasing ability of coho to defend larger feeding territories as the fish grow). Here we show that predicted fitness is

consistent with what is known about the relationship between freshwater growth and age at maturity in coho salmon only when $\psi < 0.5$. Hence, it seems likely that values of ψ , with the associated interpretations of relatively less clumped and less defensible resources, are descriptive of coho, at least during the first growing season at sea, and influence the likelihood of early maturation. Our qualitative predictions about the relationship between age at maturity and growth performance in freshwater are also robust to changes in mortality schedules and $\overline{E}_{\rm sw}$. In general, our predictions are consistent with evidence from the literature that large male smolts are more likely to mature early (Garrison 1971; Hager and Noble 1976) and that early maturation is predicted to be a rare event for females.

The details of our model suggest a proximate mechanism for why large smolts are observed to mature early. The reduced growth potential at sea for large smolts results in early maturation maximizing fitness (Snover et al. 2005). Large smolts cannot attain a large enough size to be competitive in the fighting tactic (males) or to maximize nest survival and fecundity (females). Gross (1996) assumes that large smolts are high status and, as such, their choice of the sneaking tactic indicates that this tactic has the highest overall fitness. Our results suggest that length at smolt transformation may not be the appropriate measure of "status" but that remaining growth potential is more appropriate (Jonsson and Jonsson 1993). Growth potential is affected by environmental conditions in the ocean as well as growth performance in freshwater.

Conclusions

In this study we modeled how phenotypes (i.e., age at maturity and reproductive tactic) are maintained in a population via variation in top-down and bottom-up factors. Our model demonstrates that these factors, in conjunction with limits on growth potential, are sufficient to maintain multiple phenotypes for both males and females within a population. We found switch points based on the growth coefficient k (or length at the smolt transformation), at which the fitness of two phenotypes are equivalent, with the fitness of one phenotype predominating over the other as individual ks deviate from the switch points. Studies of the maintenance of alternative reproductive tactics based on game theory suggest that negative frequency dependence is necessary to maintain alternative strategies in a population (e.g., Hutchings and Myers 1994; Repka and Gross 1995; Roff 1996). Here, we turned the question around and considered how the genotype-by-environment interaction can determine phenotype. Game-theoretic modifications (including negative frequency dependence) to the effective transmission of genotypes between generations may create an interesting feedback loop. For a more complete understanding of the processes involved in maintaining alternative reproductive strategy, our modeling approach ultimately needs to be considered in conjunction with the game-theoretic approach.

Nevertheless, our modeling approach can provide insights for understanding how the diversity of salmon life histories results from the interplay of ecosystem and genetic controls. The salmonids in general exhibit great diversity in the degree of anadromy, the amount of time spent in freshwater, and age at maturity (Stolz and Schnell 1991; Behnke 2002). Some salmonids (e.g., lake [Salvelinus namaycush], bull [Salvelinus confluentus], and golden [Oncorhynchus aguabonita trout) are essentially freshwater species, while others (e.g., pink [O. gorbuscha] and chum [Oncorhynchus keta] salmon) are obligate anadromous species. A large proportion of the species (e.g., chinook [O. tshawytscha], coho, sockeye [Oncorhynchus nerka], and Atlantic salmon; steelhead [Oncorhynchus mykiss], brown [Salvelinus trutta], and cutthroat [Oncorhynchus clarki] trout; Dolly Varden [Salvelinus malma]; and arctic [Salvelinus alpinus and eastern brook [Salvelinus fontinalis] char) are adaptive or optional anadromous species. These latter species can either be anadromous or complete their life histories in freshwater without migrating to sea. The obligate anadromous species, pink and chum salmon, migrate to sea at age 0; the facultative anadromous species migrate to sea over a range of ages and have the greatest variability in age at maturity. By appropriately structuring an environmentally explicit growth model and models of expected reproductive success and by choosing appropriate parameters, rather than fitting them to a focal species, we can expand our modeling approach to generate each of the salmonid life histories, describing variations in age at outmigration and return (age at maturity) and how genotype-by-environment interactions, including topdown and bottom-up controls, influence this variability. Ultimately, it should be possible to describe how degrees of anadromy map onto a variable background of ecosystem control. For example, Hutchison and Iwata (1997) found a positive relationship between degrees of aggressive behavior and the duration of stream residency. From the insights gained here, we predict that, given access to the sea, facultative anadromy evolves when aggressive behaviors are linked to growth performance and under strong genetic control. Obligate anadromy should evolve when growth performance is not linked to aggressive behavior and such behaviors are not controlled genetically. These predictions derive from the findings that the interplay of behavior and resource distribution has strong effects on expected growth potential at sea (Snover et al. 2005) and that genetics controls age at maturity throughout the life history. Confirming such predictions is, however, the subject of another article.

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APPENDIX

Further Details on Our Growth Model, Survival Rates, and Female Fitness Model

Growth Model

Activity Metabolism versus Standard Metabolism. The growth model of Snover et al. (2005) implicitly discriminates between energy costs associated with maintenance and energy costs associated with activity; however, this discrimination needs clarification. Ursin (1979) provides a nice mathematical exposition of energy accumulation and utilization and how growth rates of fishes derive from the difference of these two processes, which is the basic concept underlying von Bertalanffy's growth model. The gross instantaneous rate of energy accumulation is equal to βfhw^m (see eq. [5] in Ursin 1979), where β is the fraction of ingested food that is absorbed, f is the "feeding level" (using Ursin's terminology), h is the "coefficient of food consumption," w is body weight, and m is an allometric scaling parameter. Importantly, there are two components of energy utilization: "losses due to feeding and assimilation activities (feeding catabolism)" and losses that are "independent of feeding (fasting catabolism)." The instantaneous rate of feeding catabolism is equal to $\alpha\beta fhw^m$ (see eq. [6] in Ursin 1979), where α determines how much of the absorbed food is used to pay for the energetic demands of feeding and assimilation. The instantaneous rate of fasting catabolism is equal to cwⁿ (see eq. [7] in Ursin 1979, but use the parameter name c instead of Ursin's original parameter k), where c is a coefficient and n is another allometric scaling parameter. Note that the instantaneous rate of fasting catabolism is the standard metabolic rate (Ursin describes how to make this rate temperature dependent if desired). The growth rate (dw/dt) is equal to the difference between the rate of energy accumulation and the summed rates of feeding and fasting catabolism: $dw/dt = \beta fhw^m - (\alpha \beta fhw^m + cw^n) =$ $(1 - \alpha)\beta fhw^m - cw^n$. Thus, the growth rate is the difference between net energy accumulated by feeding (where net accumulation is the difference between gross accumulation

and feeding catabolism) and the standard metabolic rate. If we set $H = (1 - \alpha)\beta fh$, we see that Ursin's model is the same as equation (1) in Snover et al. (2005): $dw/dt = Hw^m - cw^n$. Both Ursin (1979) and Snover et al. (2005) use a weight-length relationship of the form $w = ql^3$ and set m = 2/3 and n = 1 to provide a model for describing growth in length (1): dl/dt = E - kl. Here, *q* is a scalar, $E = H/3q^{1/3}$, and k = c/3. Snover et al. (2005) set q = 1, and we used this formulation in equation (1) here. Note that the first term on the right side of the equation for growth in length is still the net rate of energy accumulation, and the second term is still the standard metabolic rate. Thus, the separation of standard metabolic costs and other metabolic costs is implicit in the model we use for growth. Finally, we note that Ursin (1979), who, again, provided an exposition in which activity metabolism was separated from standard metabolism, also arrived at the equation $L_{\infty} = E/k$. Thus, while it is reasonable to say that our relationship between k and L_{∞} is an "artifact" of our model, it is also an artifact of the separation between standard and activity metabolism.

To maintain body mass, fish with high standard metabolic rates (related to k; Snover et al. 2005) require increased food intake compared to fish with lower standard metabolic rates (Jonsson and Jonsson 1993; Forseth et al. 1994). When food is limiting, fish with high metabolism are the first to detect the shortage, and Forseth et al. (1994) suggest that such fish will be the first to react to such limitation by, for example, migrating between habitats and maturing. The growth model described by Snover et al. (2005) incorporates these ideas of variability in metabolism and food availability into individual growth rates, and predictions from that model generally support the mechanism suggested by Forseth et al. (1994). Under conditions of limited food resources at sea, a fish with high k should be more likely to mature sooner than one with low kbecause the former individual will have less potential to grow at sea. There is considerable empirical evidence supporting this idea, and salmonids displaying increased growth rates in freshwater (the largest smolts) are more likely to mature early (e.g., Skilbrei 1989; Nicieza and Braña 1993).

Assumptions regarding ψ , E, and k. Snover et al. (2005) limited ψ in equation (4) to the interval [0, 1], and here we provide further interpretation of the growth model when ψ is at the extremes of this interval. From equation (4), when $\psi = 0$, $E = \gamma$ and is constant across all individuals. Under this condition, from equation (3), animals with a higher k will have a smaller asymptotic length and lower growth potential than animals with lower k (Jonsson and Jonsson 1993; Forseth et al. 1994). Coho with higher metabolism (higher k) are more aggressive than those with

lower metabolism (Metcalfe et al. 1995) but do not necessarily have a foraging advantage over them. Resources must be distributed in a way that facilitates resource acquisition via aggressive behavior (e.g., resources should be aggregated rather than dispersed). When $\psi=1$, $\gamma=L_{\infty}$ and is constant across all individuals. Biologically, this implies that animals with higher k must have a higher k and therefore must be able to acquire more resources than animals with lower k. This would imply that resources are distributed (e.g., resources are clumped; Snover et al. 2005) such that aggressive behaviors are effective for securing food in competitive interactions with conspecifics. Intermediate values of ψ provide intermediate interactions between resource distribution and behavior.

Aggressive coho seem to have an advantage in freshwater streams where food tends to be heterogeneously distributed, and the establishment of territories can provide an aggressive individual with access to more resources than a nonaggressive individual (Puckett and Dill 1985; Nielsen 1992; Martel 1996). This may not be the case for other salmonids (e.g., Atlantic salmon; see Martin-Smith and Armstrong 2002; Harwood et al. 2003), but predictions from the growth model presented by Snover et al. (2005) are robust, provided there is a positive relationship between k and smolt length, which is the case for $\psi > 0.6$.

We made the assumption that k remains constant in an individual. Snover et al. (2005) acknowledged that since k is related to metabolism and coho salmon are ectothermic, it will vary with temperature; however, these metabolic fluctuations will be relatively constant across all individuals from a single cohort. Hence, if k is allowed to decrease upon outmigration to sea, it will decrease uniformly across all individuals, and the qualitative predictions from our models will be the same as with the assumption of constant k.

We also made the assumption that E remains constant from year to year in saltwater. Obviously, this is not realistic because ocean conditions change dramatically both seasonally and interannually. Nevertheless, annual variations in $E_{\rm SW}$ around a mean do not change the qualitative results of our model.

Coho Life Cycle. Our models are based on the predominant life cycles for coho salmon (Sandercock 1991). After emergence from the gravel, fry grow in freshwater for 12 months. The smolt transformation is assumed to occur in month 12, and growth in saltwater occurs while coho are 13–42 months old. Fish that mature early spend 6 months at sea; they experience no sea winters and one growing season. Females are almost never observed to mature early. Most coho, both males and females, spend 18 months at sea and mature at 3 years of age; these fish experience one winter at sea and two growing seasons. Fish that delay

maturation experience two sea winters and three growing seasons, spending a total of 30 months at sea and maturing at 4 years of age. Typically, male coho do not delay maturation by spending 30 months at sea.

Survival Rates

There are numerous estimates of overall survival for coho salmon. Shapovalov and Taft (1954) estimated 1.2%-1.6% survival from egg to smolt for coho from Waddell Creek, California. They also estimated that survival from egg to emerging fry is 65%-85%, which suggests about 1.8%-1.9% survival from emergence to smolt. Coronado and Hilborn (1998) found that marine survival in coho salmon from a variety of stocks ranged from 0.92% to 6.21% for 3-year-old returning adults. McGurk (1996) compiled estimates of survival from smolt to adult in coho salmon. The mean of these estimates was 13.5%. Table 1 details the values for m_0 and m_1 used in the fitness functions.

Components of Female Fitness

In addition to egg production, breeding competition is an important source of selection for size in semelparous female salmonids (Fleming and Gross 1994). Larger female salmonids dig deeper nests (van den Berghe and Gross 1984; Holtby and Healey 1986) and may have longer postreproductive survivorship, allowing for longer duration of nest defense (van den Berghe and Gross 1986; but see McPhee and Quinn 1998). While these factors do not guarantee the nest will not be superimposed by a subsequent female (Steen and Quinn 1999), together they decrease the likelihood of superimposition. Deeper nests may also result in decreased probability of nest loss due to scour (Montgomery et al. 1996; Steen and Quinn 1999). It is difficult to incorporate a relationship between nest depth and female size into a fitness equation, so we used the relationship between percentage of nests still intact at the end of the breeding season and female length presented by van den Berghe and Gross (1989).

In some systems there may be a penalty on large female body size. For example, Quinn et al. (2001a) suggest that in sockeye salmon (Oncorhynchus nerka), access to spawning grounds and size-biased predation by bears may select against larger females. Nevertheless, the studies we found for coho indicate positive relationships between adult female body size and egg survival (van den Berghe and Gross 1989; Fleming and Gross 1994). The timing of arrival to the spawning ground is also important because late arrivals are less likely to have nests superimposed (McPhee and Quinn 1998); however, we did not consider this factor here.

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