- A Model For Sea Lice (Lepeophtheirus salmonis)
- 2 Dynamics In A Seasonally Changing Environment
- Matthew A. Rittenhouse^a, Crawford W. Revie^b, and Amy Hurford^{a, c}
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- ^a Department of Biology, Memorial University of Newfoundland, St. John's, NL, Canada
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- ⁵ Centre for Veterinary Epidemiological Research, Department of Health Management,
- 8 Atlantic Veterinary College, University of Prince Edward Island, 550 University Avenue,
- 9 Charlottetown, Prince Edward Island, Canada C1A 4P3
- ^c Department of Mathematics and Statistics, Memorial University of Newfoundland, St.
- John's, NL, Canada A1C 5S7
- Matthew A. Rittenhouse¹ mar665@mun.ca
- 13 Crawford W. Revie crevie@upei.ca
- Amy Hurford ahurford@mun.ca
- ¹ Corresponding Author

1 Introduction

The control of parasitic organisms is a major concern in marine aquaculture. In particular, sea lice (Lepeophtheirus salmonis and Caligus spp.) cause substantial economic losses on salmon farms (Costello 2009). Due to their economic importance, control of sea lice 19 on salmon farms has been named one of the top priorities in aquaculture research by both scientists and aquaculture practitioners (Jones et al. 2014). Adequate control of sea lice is predicated on the ability to predict future lice levels from current population and environmental trends, as well as predicting the effectiveness of different treatment regimes. These two needs can be accomplished through mathematical modelling and it is imperative that tractable and biologically sound models are developed to aid practitioners in decisions regarding sea lice dynamics. Seasonal environmental variability plays a major role in the dynamics of many disease systems (Altizer et al. 2006). Temperature and salinity affect several characteristics of sea lice life history (summarized in Table 1), thus models of sea lice dynamics must be able to incorporate the effects of seasonally varying temperature and salinity on the sea louse lifecycle.

Table 1: The effect of water temperature and salinity on characteristics of sea lice life history.

Characteristic	Increasing Temp.	Increasing Salinity	Ref.	
Nauplius maturation rate	Increase	No effect	Stien et al. 2005	
Chalimus maturation rate	Increase	No effect	Stien et al. 2005	
Egg viability	No effect	Increase	Johnson & Albright 1991	
Nauplius mortality rate	No effect	Decrease	Johnson & Albright 1991	
Copepodid mortality rate	No effect	Decrease	Johnson & Albright 1991; Bricknell et al. 2006	
Chalimus mortality rate	No effect	Decrease	Connors et al 2008	
Adult mortality rate	No effect	Decrease	Connors et al 2008	

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A variety of deterministic (Revie et al. 2005, Stien et al. 2005, Robbins et al. 2010, 33 Gettinby et al. 2011, Aldrin et al. 2013, Groner et al. 2014, Kristoffersen et al. 2014) and stochastic (Aldrin et al. 2013, Groner et al. 2014) models have been derived to predict sea lice dynamics. Revie et al. (2005) derived a life stage model with fixed delays and constant mortality rates that formed the basis for the simulation tool, SLiDESim. Robbins et al. (2010) utilized SLiDESim to search for optimal treatment strategies in Scottish farms. Gettinby et al. (2011) tested the SLiDESim model on sea lice collection data from the Hardangerfjord in south-west Norway. These authors concluded that for the model to be utilized in evaluating treatment strategies, a better understanding of the underlying biological and environmental factors, including temperature-dependent maturation and salinity-dependent survival, was necessary. Kristoffersen et al. (2014) used Bělehrádek functions derived in Stien et al. (2005) to estimate degree-days needed for sea lice to mature from one stage to the next and derived a life stage model with temperature-based delays and constant mortality rates. Aldrin et al (2013) used a stochastic spatio-temporal model to show how seawater temperatures, fish stock population, and distance between farms contributed to predicted sea lice counts. Groner et al. (2014) created a stochastic matrix population model to examine the effects of seasonally varying temperature on treatment schemes and louse mate limitation. A deterministic model capable of accounting for both the effects of seasonally varying 50 temperature on sea lice maturation, and the effects of seasonally varying salinity on sea lice mortality has not yet been developed. Stien et al. 2005 suggest that the delay differential equation models described in Nisbet & Gurney (1983) as a method to address this need. Delay differential equations models of this type have been successfully used in epidemiological models of koi herpes virus (Omori & Adams 2010) and malaria (Beck-Johnson et al. 2014).

We present a delay differential equation model of the sea lice lifecycle with temperature-

dependent stage durations, salinity-dependent mortality, and time-dependent temperature/salinity.

Where possible, model parameters are fitted to values from the literature for the species,

Lepeophtheirus salmonis. The time dependent basic reproductive ratio, $R_0(t)$, is derived nu-

60 merically to quantify seasonal differences in sea lice replenishment at sites in British Columbia

and southern Newfoundland, Canada. Additionally, a sensitivity analysis is conducted to

62 identify the parameters that most substantially affect the model predictions.

$_{ ext{\tiny 53}}$ 2 The Model

64 We developed a model for sea lice dynamics on salmon farms that includes temperature-

dependent maturation delays and salinity-dependent mortality. Lepeophtheirus salmonis

exhibit 8 distinct life stages, consisting of nauplius I/II, copepodids, chalimus I/II, pre-adult

₆₇ I/II, and adults (Hamre et al 2013). For the purpose of modelling, we assume that sea lice

may be in 1 of 4 possible functional states: planktonic non-infectious nauplii (P), infectious

69 copepodids (I), non-reproductive chalimus and pre-adults (C), or adult females, (A). Each

individual matures through the states in order from nauplius (P), to copepodid, (I), to

chalimus (C), and finally to adult female (A) (Fig. 1).

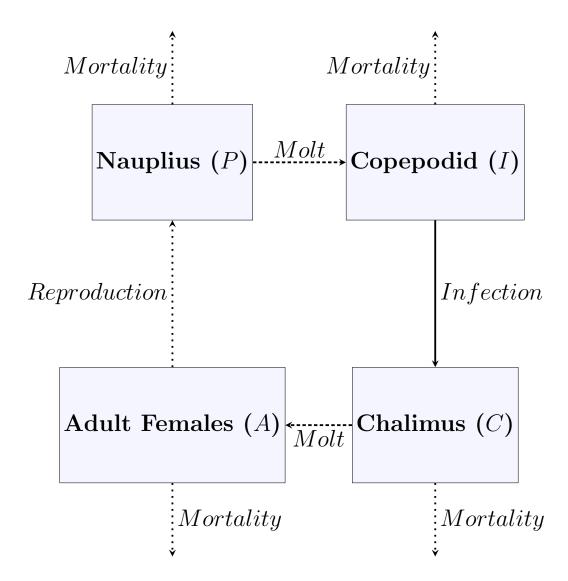


Figure 1: Modelled life cycle of the sea louse. Dashed arrows indicate aspects of the life history affected by temperature. Dotted arrows indicated aspects of the life history affected by salinity.

The length of time that a nauplius or chalimus requires to mature to their respective next life stages depends on water temperature (Table 1). Let $\gamma_x(T(t))$ be a function that describes the rate of change in the level of development for a given stage $x \in \{P, C\}$ as it depends on temperature (T), which changes over time (t). For notational simplicity, we write simply $\gamma_x(t)$, because given functions that describe how temperature changes with respect to time (T(t)), and how the development rate changes with respect to temperature $(\gamma_x(T))$, we can then determine how the development rate changes with respect to time $(\gamma_x(t))$ without needing to explicitly reference the dependence on temperature.

The waiting times associated with maturation are such that a cohort exiting a state x at time t, will all have entered that stage at $t - \tau_x(t)$. The waiting time, $\tau_x(t)$, depends on the development rate, $\gamma_x(t)$, and is defined as the length of time that it takes sea lice to reach

a threshold development level, \bar{q}_x , given that they entered the stage x with a development

level $q_x = 0$. As such, $\tau_x(t)$ is implicitly defined as,

$$\bar{q}_x = \int_{t-\tau_x(t)}^t \gamma_x(t') \, \mathrm{d}t' \tag{1}$$

85 (Nisbet & Gurney, 1983).

Natural mortality occurs in all stages at a per capita rate $\mu_y(S(t))$, where $y \in \{P, I, C, A\}$.

Natural mortality is a function of salinity S(t), which is a function of time (t). For notational simplicity, we write simply $\mu_y(t)$, because given functions that describe how salinity changes with respect to time (S(t)), and how the mortality rate changes with respect to salinity $(\mu_y(S))$, we can then determine how the mortality rate changes with respect to time $(\mu_y(t))$ without needing to explicitly reference the dependence on salinity. Not all members of a cohort who enter a stage x at time $t - \tau_x(t)$ survive to mature at time t. The proportion of the cohort that survive the maturation period is,

$$\phi_x = e^{-\int_{t-\tau_x(t)}^t \mu_x(t')dt'}.$$

The proportion of eggs that produce viable nauplii is a function of salinity, v(t). All other

events in the sea lice life history do not depend on temporally varying quantities and are
assumed to occur at constant per capita rates. The complete model is a system of delay
differential equations,

$$\frac{dP}{dt} = \eta \epsilon v(t) A(t) - \eta \epsilon v(t - \tau_P) A(t - \tau_P) \frac{\gamma_P(t)}{\gamma_P(t - \tau_P)} \phi_P - \mu_P(t) P(t), \tag{2}$$

$$\frac{dI}{dt} = \eta \epsilon v(t - \tau_P) A(t - \tau_P) \frac{\gamma_P(t)}{\gamma_P(t - \tau_P)} \phi_P - \iota f I - \mu_I(t) I(t), \tag{3}$$

$$\frac{dC}{dt} = \iota f I(t) - \iota f I(t - \tau_C) \frac{\gamma_C(t)}{\gamma_C(t - \tau_C)} \phi_C - \mu_C(t) C(t), \tag{4}$$

$$\frac{dA}{dt} = \frac{1}{2} \iota f I(t - \tau_C) \frac{\gamma_C(t)}{\gamma_C(t - \tau_C)} \phi_C - \mu_A(t) A(t), \tag{5}$$

$$\frac{d\tau_P}{dt} = 1 - \frac{\gamma_P(t)}{\gamma_P(t - \tau_P)},\tag{6}$$

$$\frac{d\tau_C}{dt} = 1 - \frac{\gamma_C(t)}{\gamma_C(t - \tau_C)},\tag{7}$$

where η is the number of eggs per egg string, ϵ is the rate of egg string production, ι is the rate of infection per fish, f is the number of fish on the farm, and all model parameters are summarized in Table 2. Equations (6) and (7) arise from differentiating equation (1) with respect to time (Nisbet and Gurney, 1983). The $\gamma_x(t)/\gamma_x(t-\tau_x)$ terms arise because we wanted to ensure a correspondence between our model (which lumps all individuals with a development level $q_x < \bar{q}_x$ together into one state) and a model that treats the development level as a continuous quantity (Nisbet & Gurney, 1983; see Appendix A for further details).

Table 2: Model Parameters

Parameter	Description	Units	Value	Ref.		
η	eggs per clutch	eggs	592	Heuch et al. 2000		
ϵ	egg string production rate	$\frac{1}{day}$	0.0476	Heuch et al. 2000		
l	infection rate	$\frac{1}{day}$	0.01	NA		
f	number of fish	fish	1000	NA		
a	average annual temperature	$^{\circ}\mathrm{C}$	9	NA		
b	temperature amplitude	$^{\circ}\mathrm{C}$	4	NA		
c	average annual salinity	psu	25	NA		
d	salinity amplitude	psu	2	NA		
β_P	nauplius development at $10^{\circ}C$	$\frac{1}{\sqrt{day}}$	0.525	Stien et al. 2005		
β_C	chalimus development at $10^{\circ}C$	$\frac{1}{\sqrt{day}}$	0.177	Stien et al. 2005		
T_{REF}	reference temperature for β_x	$^{\circ}\mathrm{C}$	10	Stien et al. 2005		
α_P	nauplius development = 0 $(T_{REF} - \alpha_P)$	$^{\circ}\mathrm{C}$	13.01	Stien et al. 2005		
α_C	chalimus development = 0 $(T_{REF} - \alpha_P)$	$^{\circ}\mathrm{C}$	11.94	Stien et al. 2005		
σ_{0E}	egg salinity-viability intercept	NA	-0.458	Johnson & Albright 1991		
σ_{1E}	egg salinity-viability slope	$\frac{1}{psu}$	0.037	Johnson & Albright 1991		
σ_{0P}	nauplius salinity-mortality intercept	$\frac{1}{day}$	0.4492	Brooks & Stucchi 2006		
σ_{1P}	nauplius salinity-mortality slope	$\frac{1}{day*psu}$	-0.01484	Brooks & Stucchi 2006		
σ_{0I}	copepodid salinity-mortality intercept	$\frac{1}{day}$	0.4492	Brooks & Stucchi 2006		
σ_{1I}	copepodid salinity-mortality slope	$\frac{1}{day*psu}$	-0.01484	Brooks & Stucchi 2006		
σ_{0C}	chalimus salinity-mortality intercept	$\ln hours$	4.12	Connors et al. 2008		
σ_{1C}	chalimus salinity-mortality slope	$\ln \frac{hours}{psu}$	0.124	Connors et al. 2008		
σ_{0A}	adult salinity-mortality intercept	$\ln hours$	4.12	Connors et al. 2008		
σ_{1A}	adult salinity-mortality slope	$\ln \frac{hours}{psu}$	0.124	Connors et al. 2008		

It should be noted that our model lacks a mechanism for density dependence in the sea lice population. Sea lice could exhibit density dependence due to an Allee effect caused by difficulties in mate finding at low densities (Krkošek et al. 2012, Groner et al. 2014). They may also exhibit density dependence due to host mortality or decreased reproduction at high

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intensities. In an aquaculture setting, managers will typically intervene with chemotherapeutics before any natural density-dependent regulation of the sea louse population can 111 occur. 112

3 **Model Parameterization**

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For our model, the maturation rate is the inverse of the Bělehrádek functions describing minimum development time as a function of temperature (Stien et al. 2005), 115

$$\gamma_x(T) = \left(\frac{\beta_x(T - T_{REF} + \alpha_x)}{\alpha_x}\right)^2.$$

The shape of the function is described by the duration of the life stage (β_x^{-2}) at the reference temperature (T_{REF}) and by the location of an asymptote at $T_{REF} - \alpha_x$. The terms β_P and 117 β_C are identical to the β_2 terms for time from hatching to copepodid and time from infection 118 to adult females in Stien et al. (2005). The α_P and α_C terms are the product of β_1 and β_2 119 from Stien et al. (2005). The reference temperature is 10°C. 120 Water temperature (T(t)) and salinity (S(t)) on salmon farms varies over time. We use 121

sinusoidal functions to describe the general annual patterns,

$$T(t) = a - b \cos\left(\frac{2\pi t}{365}\right), \tag{8}$$

$$S(t) = c - d\cos\left(\frac{2\pi t}{365}\right),\tag{9}$$

where a is the average annual temperature, c is the average annual salinity, and b and d are

the respective amplitudes of the cosine functions. Sinusoidal functions of the form,

$$T(t) = a + b_1 \sin\left(\frac{2\pi t}{365}\right) + b_2 \cos\left(\frac{2\pi t}{365}\right), \tag{10}$$

$$T(t) = a + b_1 \sin\left(\frac{2\pi t}{365}\right) + b_2 \cos\left(\frac{2\pi t}{365}\right),$$

$$S(t) = c + d_1 \sin\left(\frac{2\pi t}{365}\right) + d_2 \cos\left(\frac{2\pi t}{365}\right),$$
(10)

were fit to monthly temperature and salinity data from a salmon farm in the Broughton 125 Archipelago of British Columbia (Marty et al., 2010; Fig. 4D and E) and to quarter-hourly temperature and salinity data from a salmon farm on the southern coast of Newfoundland 127 (data provided by the Aquaculture Real-Time Integrated Environmental System; Fig. 4D 128 and E). 129 Mortality is related to salinity via linear and log-linear relationships from the literature. 130 The salinity-mortality relationship $(\mu_A(S))$ is log-linear for adult sea lice (Connors et al. 2008). We assume that the mortality rate for adults and chalimi $(\mu_C(S))$ is similar. The 132 salinity-mortality relationship for nauplii $(\mu_P(S))$ is a linear model from Brooks & Stucchi 133 (2006), which is fit to data from Johnson & Albright (1991). We assume that the mortality rate for nauplii and copepodids $(\mu_I(S))$ is similar. 135 The number of eggs per egg string (η) was parameterized using data from Heuch et al. 136 (2000). The lower bound on the egg string production rate (ϵ) is taken from Mustafa et 137 al. (2000) and is used as the default egg string production rate. The upper bound for 138 the egg string production rate, used in the uniform distribution for the sensitivity analysis, 139 was taken from Heuch et al. (2000). Egg viability (v(S)) consists of a linear model fit to 140 salinity-survivorship data from Johnson & Albright (1991). The infection success rate (ι) is dependent on a number of variables and is not well 142 understood at the farm level. As such, a wide range of values, from 0.001 to 0.9, were used

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in the sensitivity analysis and a value of 0.01 was chosen as the default value.

5 4 Model Dynamics

We numerically solved the system of equations (2)-(7) using the PBSddesolve package in R. Due to the lack of density dependence our model produces either unbounded growth or extinction (Figs. 2 and 3). The abrupt changes in the trajectory slope shown in Fig. 2 indicate the beginning of successive generations. When the initial infection occurs on the coldest day of the year, the second cohort will take less days to mature to adulthood than the first, due to warming temperatures. Sites with higher average annual temperatures will take less time for cohorts to reach adulthood than sites with lower average annual temperatures (Fig. 2).

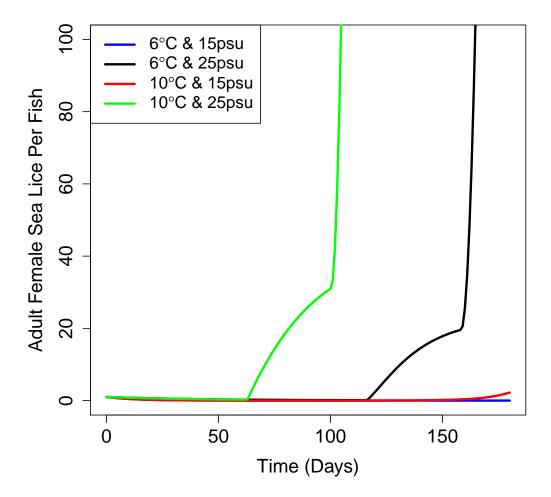


Figure 2: Four model simulations under high/high, high/low, low/high, low/low average annual temperature and salinity values. Amplitudes were 4 °C and 2 ppt for all simulations. All simulations began on the coldest, least saline day of the year. Note that increasing temperature increases parasite numbers and decreases generation time. Note also that generation time decreases as the simulation progresses, due to warming temperatures. The sea lice population dies off under the low/low temperature/salinity condition.

We ran the model with average annual temperatures of 6, 7, 8, or 9°C and average annual salinities of 14, 16, 18, or 20 psu. After simulating four years, environmental conditions that

resulted in > 1 adult female per fish were considered to be favourable to sea lice, while environmental conditions with < 1 adult female per fish were considered unfavourable. Sea lice in low temperature/low salinity environments will die out. As either temperature or salinity increases, conditions for the sea lice population improve. Sea lice populations are viable at ≥ 18 psu at all temperatures investigated and sea lice populations can persist at lower salinities in warmer climates (Fig. 3).

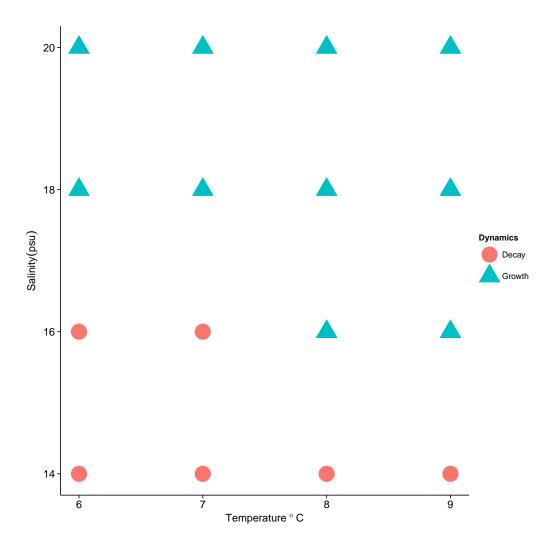


Figure 3: Long term persistence/extinction dynamics at sixteen temperature and salinity combinations. Amplitudes were 4 °C and 2 psu for all simulations. Environmental conditions that result in > 1 adult female sea louse per fish after four years are shown as triangles. Environmental conditions that result in < 1 are shown as circles.

$$_{^{162}}$$
 5 $R_0(t)$

The basic reproductive ratio, R_0 , is commonly used as a measure of reproductive success in populations. The basic reproductive ratio can be defined as the "expected number of

secondary individuals produced by an individual in its lifetime" (Heffernen et al. 2005) and acts as a threshold condition that indicates either population persistence or extinction 166 (Caswell 2009). When $R_0 \geq 1$ the population will grow with each subsequent generation and persist, whereas when $R_0 \leq 1$ the population will shrink with each subsequent generation 168 until extinction. In seasonal systems, R_0 will depend on the time that the infection is 169 introduced to the system. We define $R_0(t)$ such that it is the number of second generation adult females produced by a single adult female, who enters the system at time, t. As such, 171 $R_0(t)$ depends on the probability that the nauplius survives each successive life stage and the timing and duration of egg string hatching during the adult female stage (see Appendix 173 B). We determined $R_0(t)$ numerically by augmenting the system of equations (2)-(7) with a 174 delay differential equation describing the number of adult females in the second generation, where this second generation does not reproduce (see Appendix B for details). 176 In the Broughton Archipelago, temperatures are favourable year round (5th percentile = 177 6.61, median = 8.90, 95th percentile = 11.75), while salinity levels are very favourable to sea 178 lice survival in the winter and very unfavourable during the summer (5th percentile = 16.11, 179 median = 27.30, 95th percentile = 32.15). The environmental conditions most favourable to sea lice growth are asynchronous because high sea surface temperatures coincide with low 181 salinities and visa versa. We find $R_0(t)$ to be highest in December, when salinity is high, but 182 temperatures are low (Fig. 4A). As such, sea lice that enter the farm in December will go 183 on to produce the most offspring despite having longer generation times than sea lice that 184 hatch in the summer months (Fig. 4B). The value of $R_0(t)$ is not < 1 at any point during the year, so an adult female that enters at any given time can reasonably be expected to 186 replace itself over the course of its lifetime. $R_0(t)$ reaches a peak of 475.13 in December and 187 a low of 23.68 in June and has a mean value of 220.98.

In southern Newfoundland, temperatures are colder (5th percentile = 2.48, median = 189 6.00, 95th percentile = 13.20) than the Broughton Archipelago site and sea lice maturation 190 can take a long time during the winter months. Salinity is mostly constant year round 191 (5th percentile = 19.92, median = 22.77, 95th percentile = 26.06), with a low in the spring 192 months. In southern Newfoundland, the environmental conditions that are most favourable 193 to sea lice growth are synchronous: both high temperatures and high salinities occur towards the end of the summer. We find that $R_0(t)$ is highest in August (Fig. 4A), after maturation 195 times plummet during the summer months (Fig. 4B) and when time to maturity is shortest 196 (Fig. 4B). The value of $R_0(t)$ is not < 1 at any point during the year, so an adult female 197 that enters at any given time can reasonably be expected to replace itself over the course of 198 its lifetime. $R_0(t)$ reaches a peak of 125.41 in August, a low of 3.27 in December and has a mean of 46.72.

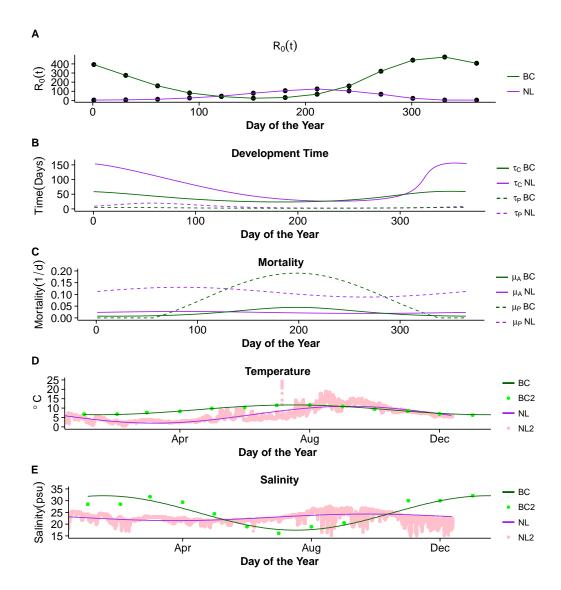


Figure 4: $R_0(t)$ is presented for sites in Newfoundland and British Columbia (A). $R_0(t)$ is affected by maturation time (B) and the mortality rates of parasitic stages $\mu_A(t)$ and free-living stages $\mu_P(t)$ (C). Temperature (D) and salinity (E) for the two sites affect development time (B) and mortality (C), respectively.

6 Sensitivity Analysis

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We conducted a sensitivity analysis on seven model parameters to analyze the effects of
their variation on the abundance of female adult sea lice (A). Parameter distributions were
estimated from the literature (Table 3). A Latin Hypercube Sampling (LHS) scheme was
used to sample the parameter space and partial rank correlation coefficients (PRCC) were
used as a test statistic for the sensitivity analysis (Blower & Dowlatabadi 1994).

Table 3: Parameter values for sensitivity analysis

Parameter	Units	Min	Max	Median	STD	PDF	Ref.
η	eggs	NA	NA	492	200	Normal	Heuch et al. 2000
ϵ	$\frac{1}{day}$	0.0476	0.0576	NA	NA	Uniform	Heuch et al. 2000, Mustafa et al. 2000
ι	$\frac{1}{day}$	0.001	0.9	0.01	NA	Triangle	Groner et al. 2014
a	$^{\circ}\mathrm{C}$	7	10	NA	NA	Uniform	Beamish & Jones 2011
c	psu	20	30	NA	NA	Uniform	Beamish & Jones 2011
β_P	$\frac{1}{\sqrt{day}}$	NA	NA	0.525	0.017	Normal	Stien et al. 2005
β_C	$\frac{1}{\sqrt{day}}$	NA	NA	0.177	0.006	Normal	Stien et al. 2005

The simulation begins with only adult females and the only parameter that affects adult mortality is mean salinity (c, Fig. 5A). After the cohorts start maturing the size of the adult female population is also affected by parameters relating to maturation, infection, and reproduction (Fig. 5A and B). The three most sensitive parameters at 180 days were mean temperature (a), mean salinity (c), and the number of eggs per egg clutch (η ; Fig. 5A and B). Female sea lice abundance is more sensitive to the temperature-development relationship of the chalimus and pre-adult stages (β_C) than it is to the temperature-development relationship of the nauplius stage (β_P ; Fig. 5A). Despite a large level of uncertainty about the value of

the infection rate, the model is least sensitive to infection rate, out of the seven parameters examined (Fig. 5B).

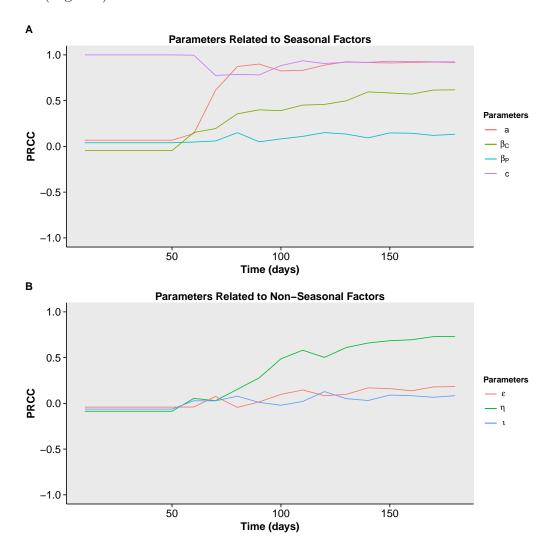


Figure 5: Sensitivity to 7 parameters. The PRCC values at 180 days (from highest PRCC to lowest) are $c=0.92,~a=0.92,~\eta=0.73,~\beta_C=0.62,~\epsilon=0.18,~\beta_P=0.13,~{\rm and}~\iota=0.08.$

7 Discussion

Sea lice control is one of the top priorities in aquaculture research. Temperature and salinity affect maturation rates, mortality, and egg viability; so control of sea lice relies on

understanding their population dynamics in relation to their environment. We derived a deterministic model of the sea louse lifecycle, with temperature-dependent maturation and 222 salinity-dependent mortality. We conducted numerical analyses to: characterize sea lice pop-223 ulation dynamics for different environmental conditions, determine the time-dependent basic 224 reproductive number, R0(t), for BC and southern Newfoundland, and perform a sensitivity 225 analysis. There is a substantial difference between the timing of the peak in $R_0(t)$ for British 227 Columbia and Newfoundland. We found that the peak value of $R_0(t)$ for both the British 228 Columbia and the Newfoundland sites occurred during peak salinity levels, although in 220 Newfoundland the salinity levels were fairly constant and the highest salinity levels also 230 coincided with the highest sea surface temperatures. As such, optimal treatment schemes will also differ between these two sites. In both the Broughton Archipelago in British Columbia 232 and southern Newfoundland, fecund sea lice are able to replace themselves at all times of 233 the year, however, well-timed treatments may result in slower population growth, ultimately 234 leading to fewer required treatments over a production cycle. 235 We also found that the mean $R_0(t)$ was much higher at the British Columbia site than at 236 the Newfoundland site. In considering just the environmental data, it is not clear that this 237 would necessarily be the case. On the one hand, British Columbia has higher mean temper-238 atures, higher mean salinity, and higher maximum salinity: all conditions that are conducive 230 to sea lice growth, while on the other hand, British Columbia also has lower minimum salin-240 ity and favourable environmental conditions for sea lice growth are not synchronous as they are in Newfoundland. Contrary to our results, it is generally acknowledged that control of 242

sea lice is typically more straightforward in British Columbia than in Newfoundland. Our

model suggests that something other than temperature and salinity may be responsible for

the more successful management of sea lice in British Columbia. This may be the result of farm-level and regional-level management decisions, population size of wild hosts, potential 246 genotype differences in sea lice populations (Yazawa et al. 2008; Skern-Mauritzen et al. 2014), or hydrodynamical differences. 248 The comparison of the British Columbia and the Newfoundland sites suggest no general 240 patterns. To understand the dynamics of sea louse fecundity at other sites, environmental data would need to be provided for analysis using our model. This is especially pertinent 251 as salinity patterns may vary substantially over small spatial scales due to their proximity 252 to rivers, and even two sites within the same broad geographic region potentially could have 253 very different salinity patterns. 254 It is important to note that the $R_0(t)$ we calculate is not a threshold condition for sea lice 255 epidemics, since subsequent generations will hatch throughout the year and experience their 256 own $R_0(t)$ values. $R_0(t)$ also does not show the size of, or even the instantaneous growth 257 rate of, the population. Rather, our $R_0(t)$ provides a means of comparing reproductive 258 output at different times of the year. The methods outlined in Zhao (2015) provide a 250 framework to analytically determine how environmental conditions affect the threshold for sea lice outbreaks. One of the advantages to using a deterministic delay differential equation 261 approach is that the theoretical approaches outlined in Zhao (2015) can be utilized. 262 Our sensitivity analysis found that adult female sea lice abundance is most sensitive to 263 average annual temperature and salinity. This is likely because a large number of parameters 264 depend on temperature $(\tau_P(t))$ and $\tau_C(t)$, salinity $(\mu_P(t), \mu_I(t), \mu_C(t), \mu_C(t))$ or both $(\phi_P(t), \phi_C(t))$. Our findings that lice abundance is more sensitive to the development rate of 266 the combined chalimus/pre-adult stage than to the development rate of the nauplius stage is

in line with sensitivity analyses conducted by Revie et al. (2005) and Groner et al. (2014),

who found that sea lice numbers were most sensitive to the survival through the pre-adult stage, of which development time plays a major role.

In addition to the need for more empirical studies into the temperature-maturation rela-271 tionship of chalimus/pre-adult stages that has already been highlighted, a number of avenues 272 exist to improve model accuracy and usability. Because eggs per clutch (η) and egg clutch 273 production rate (ϵ) are two parts of the same product, the difference in PRCC values between them is solely due to the distribution we sampled from for each parameter. Egg clutch size 275 is highly variable and the number of eggs per egg string has been suggested to be dependent 276 on the temperature history of the female louse during development (Ritchie et al. 1993; 277 Heuch et al. 2000). Sea lice that develop in colder temperatures are suggested to produce 278 more eggs per egg string as a compensatory strategy for slower development, although the exact mechanism linking temperature history and egg string length is unknown (Ritchie et 280 al. 1993). As such, a better understanding of the relationship between temperature and 281 egg production is needed before it can be incorporated into mechanistic models of sea lice 282 development. Difficulties in mate finding at low densities has also been indicated as a major 283 facet of reproductive success (Stormoen et al. 2013; Groner et al. 2014). Future models may wish to explore the impacts of these biological complexities. 285

Salmon farms regularly treat for sea lice, which impacts population numbers at a level greater than environmental factors. The timing of treatment, in regards to the typical salmon farming cycle has been shown to have a large impact on sea lice numbers (Revie et al. 2005). Because salmon can be introduced to saltwater pens at most times of the year, our model is well suited to examining the effects of treatment timing on sea lice numbers. We conclude by recommending that future modelling studies incorporate detailed seasonal characterists of their chosen study site into models of sea lice population dynamics.

References

- ²⁹⁴ [1] Aldrin, M., B. Storvik, A.B. Kristoffersen, P.A. Jansen, M. Krkošek. 2013. Space-time
- modelling of the spread of salmon lice between and within norwegian marine salmon
- farms. PLoS ONE 8:e64039. DOI: 10.1371/journal.pone.0064039.
- ²⁹⁷ [2] Altizer, S., A. Dobson, P. Hosseini, P. Hudson, M. Pascual, P. Rohani. 2006. Seasonality
- and the dynamics of infectious diseases: Seasonality and infectious diseases. Ecology
- Letters 9:467-484. DOI: 10.1111/j.1461-0248.2005.00879.x.
- ³⁰⁰ [3] Beamish, R., and S. Jones. 2011. Salmon lice: an integrated approach to understanding
- parasite abundance and distribution. John Wiley & Sons. Ames, Iowa.
- [4] Beck-Johnson, L.M., W.A. Nelson, K.P. Paajimans, A.F. Read, M.B. Thomas, and O.N.
- Bjørnstad. 2014. The effect of temperature on anopheles mosquito population dynamics
- and the potential for malaria transmission. PLoS ONE 8:e79276. DOI: 10.1371/jour-
- nal.pone.0079276.
- [5] Blower, S.M., and H. Dowlatabadi. 1994. Sensitivity and Uncertainty Analysis of Com-
- plex Models of Disease Transmission: An HIV Model, as an Example. International Sta-
- tistical Review / Revue Internationale de Statistique 62:229-243. DOI: 10.2307/1403510.
- ₃₀₉ [6] Boxaspen, K. 2006. A review of the biology and genetics of sea lice. ICES Journal of
- Marine Science 63:1304-1316. DOI: 10.1016/j.icesjms.2006.04.017.
- ³¹¹ [7] Bricknell, I.R., Dalesman, S.J., O'Shea, B., Pert, C.C., A.J. Mordue Luntz. 2006. Effect
- of environmental salinity on sea lice *Lepeophtheirus salmonis* settlement success. Diseases
- of Aquatic Organisms 71:201-212. DOI: 10.3354/dao071201.

- Brooks, K.M., and D.J. Stucchi. 2006. The effects of water temperature, salinity, and currents, on the survival and distribution of the infective copepodid stage of the salmon louse (*Lepeophtheirus salmonis*) originating on Atlantic salmon farms in the Broughton Archipelago of British Columbia, Canada (Brooks, 2005)—A Response to the Rebuttal of Krkošek et al. (2005a). Reviews in Fisheries Science 14:13-23. DOI: 10.1080/10641260500448893.
- [9] Caswell, H. 2009. Stage, age and individual stochasticity in demography. Oikos 118:1763 1782. DOI: 10.1111/j.1600-0706.2009.17620.x.
- [10] Connors, B.M., E. Juarez-Colunga, and L.M. Dill. 2008. Effects of varying salinities on *Lepeophtheirus salmonis* survival on juvenile pink and chum salmon. Journal of Fish Biology 72:1825-1830. DOI: 10.1111/j.1095-8649.2008.01839.x.
- [11] Costello, M.J., 2009. The global economic cost of sea lice to the salmonid farming industry. Journal of Fish Diseases 32:115-118. DOI: 10.1111/j.1365-2761.2008.01011.x.
- [12] Gettinby, G., C. Robbins, F. Lees, P.A. Heuch, B. Finstad, R. Malkenes, C.W. Revie.

 2011. Use of a mathematical model to describe the epidemiology of *Lepeophtheirus salmo-*nis on farmed Atlantic salmon Salmo salar in the Hardangerfjord, Norway. Aquaculture

 320:164-170. DOI: 10.1016/j.aquaculture.2011.03.017.
- [13] Groner, M.L., G. Gettinby, M. Stormoen, C.W. Revie, R. Cox, E. Sotka. 2014. Modelling the impact of temperature-induced life history plasticity and mate limitation on the epidemic potential of a marine ectoparasite. PLoS ONE 9:e88465. DOI: 10.1371/journal.pone.0088465.

- 135 [14] Hamre, L.A., Eichner, C., Caipang, C., Dalvin, S.T., Bron, J.E., Nilsen, F., Boxshall,
- G., R. Skern-Mauritzen. 2013. The Salmon Louse Lepeophtheirus salmonis (Cope-
- poda: Caligidae) life cycle has only two Chalimus stages. PloS ONE 8:e73539. DOI:
- 338 10.1371/journal.pone.0073539.
- ³³⁹ [15] Heffernan, J.M., Smith. R.J., L.M. Wahl. 2005. Perspectives on the basic reproductive
- ratio. Journal of The Royal Society Interface 2:281-293. DOI: 10.1098/rsif.2005.0042
- ³⁴¹ [16] Heuch, P.A., Nordhagen, J.R., T.A. Schram. 2000. Egg production in the salmon louse
- [Lepeophtheirus salmonis (Kroyer)] in relation to origin and water temperature. Aqua-
- culture Research 31:805-814. DOI: 10.1046/j.1365-2109.2000.00512.x.
- Johnson S.C., and L.J. Albright. 1991. Development, growth, and survival of Lepeoph-
- theirus salmonis (Copepoda: Caligidae) under laboratory conditions. Journal of the Ma-
- rine Biological Association of the United Kingdom 71:425-436.
- ³⁴⁷ [18] Jones, A.C., A. Mead, M.J. Kaiser, M.C.V. Austen, A.W. Adrian, N.A. Auchterlonie,
- K.D. Black, L.R. Blow, C. Bury, J.H. Brown, G.M. Burnell, E. Connolly, A. Dingwall, S.
- Derrick, N.C. Eno, D.J.H, Gautier, K.A. Green, M. Gubbins, P.R. Hart, J.M. Holmyard,
- A.J. Immink, D.L. Jarrad, E. Katoh, J.C.R. Langley, D.O. Lee, L. Le Vay, C.P. Leftwich,
- M. Mitchell, A. Moore, A.G. Murray, E.M.R., McLaren, H. Norbury, D. Parker, S.O.
- Parry, D. Purchase, A. Rahman, F. Sanver, M. Siggs, S.D. Simpson, R.J. Slaski, K. Smith,
- M.L.Q. Syvret, C. Tibbott, P.C. Thomas, J. Turnbull, R. Whitely, M. Whittles, M.J.
- Wilcockson, J. Wilson, L.V. Dicks, W.J. Sutherland. 2014. Prioritization of knowledge
- needs for sustainable aquaculture: a national and global perspective. Fish and Fisheries.
- DOI: 10.1111/faf.12086.

- ³⁵⁷ [19] Kristoffersen, A.B., Jimenez, D., Hildegunn, V., Grøntvedt, R., Stien, A., P.A. Jansen.
- 2014. Large scale modelling of salmon lice (Lepeophtheirus salmonis) infection pressure
- based on lice monitoring data from Norwegian salmonid farms. Epidemics 9:31-39. DOI:
- 10.1016/j.epidem.2014.09.007.
- [20] Krkošek, M., Connors, B.M., Lewis, M.A., R. Poulin. 2012. Allee Effects May Slow the
- Spread of Parasites in a Coastal Marine Ecosystem. The American Naturalist 179:401-
- 412. DOI: 10.3354/aei00014.
- ³⁶⁴ [21] Marty, G.D., Saksida, S.M. T.J. Quinn II. 2010. Relationship of farm salmon, sea
- lice, and wild salmon populations. Proceedings of the National Academy of Sciences
- 107:22599-22604. DOI: 10.1073/pnas.1009573108.
- ³⁶⁷ [22] Mustafa A., Conboy, G.A., J.F. Burka. 2000. Lifespan and reproductive capacity of sea
- lice, (Lepeophtheirus salmonis), under laboratory conditions. Aquaculture Association of
- Canada, Special Publication 4:113-114.
- ³⁷⁰ [23] Nisbet, R.M., and W.S.C. Gurney. 1983. The systematic formulation of population mod-
- els for insects with dynamically varying instal duration. Theoretical Population Biology
- ³⁷² 23:114-135.
- ₃₇₃ [24] Omori, R., and B. Adams. 2010. Disrupting seasonality to control diseases out-
- breaks: The case of koi herpes virus. Journal of Theoretical Biology 271:159-165. DOI:
- ³⁷⁵ 10.1016/j.jtbi.2010.12.004.
- ³⁷⁶ [25] Revie, C.W., C. Robbins, G. Gettinby, L. Kelly, and J.W. Treasurer. 2005. A mathe-
- matical model of the growth of sea lice, *Lepeophtheirus salmonis*, populations on farmed

- Atlantic salmon, Salmo salar L., in Scotland and its use in the assessment of treatment
- strategies. Journal of Fish Diseases 28:603-613. DOI: 10.1111/j.1365-2761.2005.00665.x.
- ³⁸⁰ [26] Ritchie, G., Mordue, A.J., Pike, G.H. Rae. 1993. The reproductive output of Lepeoph-
- theirus salmonis adult females in relation to seasonal variability of temperature and
- photoperiod. In G. Boxshall & D. Defaye (Eds.), Pathogens of wild and farmed fish: Sea
- lice 153-165. New York: Ellis Horw.
- Robbins, C., Gettinby, G., Lees, F., Baillie, M., Wallace, C., C.W. Revie.
- 2010. Assessing topical treatment interventions on Scottish salmon farms using a
- sea lice (*Lepeophtheirus salmonis*) population model. Aquaculture 306:191-197. DOI:
- ³⁸⁷ 10.1016/j.aguaculture.2010.05.006.
- ³⁸⁸ [28] Skern-Mauritzen, R., Torrissen, O., K.A. Glover. 2014. Pacific and Atlantic Lepeoph-
- theirus salmonis are allopatric subspecies: Lepeophtheirus salmonis salmonis and L.
- salmonis oncorhynchi subspecies novo. BMC Genetics 15:32. DOI: 10.1186/1471-2156-
- ³⁹¹ 15-32.
- ³⁹² [29] Stien, A., P.A. Bjørn, P.A. Heuch, and D.A. Elston. 2005. Population dynamics of
- salmon lice Lepeophtheirus salmonis on Atlantic salmon and sea trout. Marine Ecology
- Progress Series 290:263-275. DOI: 10.3354/meps290263.
- ³⁹⁵ [30] Stormoen, M., Skjerve, E., A. Aunsmo. 2013. Modelling salmon lice, Lepeophtheirus
- salmonis, reproduction on farmed Atlantic salmon Salmo salar L. Journal of Fish Diseases
- 36:25-33. DOI: 10.1111/j.1365-2761.2012.01415.x.
- ³⁹⁸ [31] Yazawa, R., Yasuike, M., Leong, J., von Schalburg, K.R., Cooper, G.A., Beetz-Sargent,
- M., Robb, A., Davidson, W.S., Jones, S.R.M., B.F. Koop. 2008. EST and Mitochon-

- drial DNA Sequences Support a Distinct Pacific Form of Salmon Louse, Lepeophtheirus
- salmonis. Marine Biotechnology 10:741-749. DOI: 10.1007/s10126-008-9112-y.