Calculating the spawn index for Pacific Herring (*Clupea pallasii*) in British Columbia, Canada

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CALCULATING THE SPAWN INDEX FOR PACIFIC HERRING (*CLUPEA PALLASII*) IN BRITISH COLUMBIA, CANADA

by

Matthew H. Grinnell, Jake F. Schweigert, Matt Thompson, and Jaclyn S. Cleary

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Cat. No. Fs97-6/nnnn E ISBN xxx-x-xxx-xxxxx-x
Cat. No. Fs97-6/nnnn E-PDF ISSN 1488-5379

Correct citation for this publication:

Grinnell, M.H., Schweigert, J.F., Thompson, M., and Cleary, J.S. yyyy. Calculating the spawn index for Pacific Herring (*Clupea pallasii*) in British Columbia, Canada. Can. Tech. Rep. Fish. Aquat. Sci. nnnn: vii + 48 p.

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ABSTRACT

Grinnell, M.H., Schweigert, J.F., Thompson, M., and Cleary, J.S. yyyy. Calculating the spawn index for Pacific Herring (*Clupea pallasii*) in British Columbia, Canada. Can. Tech. Rep. Fish. Aquat. Sci. nnnn: vii + 48 p.

The spawn index time series is one component of Pacific Herring (*Clupea pallasii*) stock assessments in British Columbia, Canada. This document describes how we calculate the spawn index from spawn survey observations (e.g., spawn extent, number of egg layers, substrate type). There are three types of spawn survey observations: (1) observations of spawn taken from the surface usually at low tide; (2) underwater observations of spawn on giant kelp, Macrocystis (*Macrocystis* spp.); and (3) underwater observations of spawn on other types of algae and the substrate, which we refer to as 'understory.' We calculate the spawn index in several steps. First, we develop a conversion factor to convert Pacific Herring eggs to biomass, which is critical to calculating the spawn index. Then we calculate the spawn index for each of the three aforementioned spawn survey types: surface, Macrocystis, and understory. Finally, we combine the three spawn indices, and aggregate by stock assessment region and year to produce a relative index of combined sex spawning biomass. In addition, we identify uncertainties in spawn index calculations, and we describe how users can download the script to calculate the spawn index using an example database. Although we transform the spawn survey data from egg density to biomass in tonnes, the annual time series of egg density and biomass are relative indices of spawning biomass.

RÉSUMÉ

Grinnell, M.H., Schweigert, J.F., Thompson, M., and Cleary, J.S. yyyy. Calculating the spawn index for Pacific Herring (*Clupea pallasii*) in British Columbia, Canada. Can. Tech. Rep. Fish. Aquat. Sci. nnnn: vii + 48 p.

La série chronologique de l'indice de frai est une composante des évaluations des stocks de hareng du Pacifique (*Clupea pallasii*) en Colombie-Britannique, Canada. Ce document décrit comment nous calculons l'indice de frai à partir des observations du relevé du frai (par ex., l'étendue du frai, le nombre de couches d'œufs, le type de substrat). Il existe trois types d'observations du relevé des frayères: (1) les observations des frayères prélevées à la surface habituellement à marée basse; (2) les observations sous-marines des frayères sur varech géant, Macrocystis (*Macrocystis* spp.); et (3) les observations sous-marines des frayères sur les autres algues et le substrat, que nous appelons « sous-étage ». Nous calculons l'indice de frai en plusieurs étapes. Premièrement, nous quantifions la production d'œufs de hareng du Pacifique, ce qui est essentiel au calcul de l'indice du frai. Ensuite, nous calculons l'indice de frai pour chacun des trois types de relevés de frai susmentionnés: surface, Macrocystis, et sous-étage. Enfin, nous combinons les trois indices de frai, et les regroupons par région d'évaluation des stocks et par année pour produire un indice relatif de la biomasse combinée des géniteurs de sexe. De plus, nous identifions les incertitudes dans le calcul de l'indice de frai, et nous décrivons comment les utilisateurs peuvent télécharger le script pour calculer l'indice de frai à l'aide d'une base de données exemple. Bien que nous transformions les données du relevé de la densité des œufs en biomasse en tonnes, les séries chronologiques annuelles de la densité et de la biomasse des œufs sont des indices relatifs de la biomasse des géniteurs.

1 INTRODUCTION

Statistical age-structured stock assessment models rely on an annual indicator of relative population abundance to reconstruct a time series of estimated abundance. For Pacific Herring (*Clupea pallasii*), an index of relative population abundance is provided by monitoring the extent and intensity of the egg, or spawn, deposition throughout coastal British Columbia (BC), Canada (DFO 2019*b*). This document describes our calculations to convert spawn survey observations (e.g., spawn extent, number of egg layers, substrate type) to the spawn index for Pacific Herring in BC.

Hart and Tester (1934) first demonstrated that an estimate of Pacific Herring abundance could be determined from a count of egg deposition in a small set of sampling quadrats. The calculations in this document have been described elsewhere, in either published or informal, internal documents. The objective of this document is to collate the various calculations in their order of application. Spawn index calculations have been updated over the years as more data and analyses justify improvements; we restrict this document to describing the current method.

Annual monitoring surveys of egg deposition collect data used to calculate the spawn index. There are three types of spawn survey observations:

1. Observations of spawn taken from the surface usually at low tide,
2. Underwater observations of spawn on giant kelp, Macrocystis (*Macrocystis* spp.), and
3. Underwater observations of spawn on other types of algae and the substrate, which we refer to as 'understory.'

Surface spawn surveys are believed to be the least accurate of the three survey types, but they have the greatest temporal and spatial extent (Schweigert 1993). For example, surface spawn surveys were the only survey type prior to 1988, and they are still used extensively for minor spawns, remote spawns (i.e., outside stock assessment region boundaries; see below), as well as unusually early or late spawns. Macrocystis and understory spawn surveys are conducted using SCUBA gear, and have been used for all major spawns since 1988. Thus, we describe the spawn index as having two periods based on the predominant survey type: the surface survey period from 1951 to 1987, and the dive survey period from 1988 to present.

The inclusion of dive surveys in 1988 makes it challenging to compare the spawn index between these two periods. For example, surface surveys are less accurate than dive surveys (appendix A.1), and spawn surveyors used subjective intensity categories instead of direct egg layer estimates until 1981 (subsubsection 4.1.1). In addition, Pacific Herring spawn survey effort has been inconsistent over the time series due to available resources and departmental priorities. For example, in the past, surveyors often dedicated several months each year to spawn surveys; they used small vessels to search for spawn, and surface surveys to estimate spawn biomass. Currently, surveyors use flights to search for spawn, and underwater SCUBA surveys to estimate spawn biomass. Thus, widespread effort (both spatially and temporally) in the past has been replaced with intense effort in the present. Pacific Herring spawn surveys began in 1928, but are considered incomplete for indexing purposes prior to 1937 because many potential areas were not surveyed (Hay and Kronlund 1987).

Pacific Herring spawn survey observations have a nested hierarchical structure: samples, Macrocystis plants, and quadrats are nested within transects, transects are nested within spawns, and spawns are nested within Locations. To develop spawn indices, Locations are nested within Sections, Sections are nested within Statistical Areas, and Statistical Areas are nested within stock assessment regions (SARs) in BC (Figure 1 & Figure 2; Haist and Rosenfeld 1988). The major SARs are Haida Gwaii (formerly known as Queen Charlotte Islands), Prince Rupert District, Central Coast, Strait of Georgia, and West Coast of Vancouver Island. The minor SARs are Area 27 and Area 2 West; we do not develop spawn indices for minor SARs. We use the terms 'major' and 'minor' to describe relative differences in the geographic and biomass scales.

We calculate the spawn index in several steps. First, we develop a conversion factor to convert Pacific Herring eggs to biomass (section 2), which is critical to calculating the spawn index. Then we calculate the spawn index for each of the three aforementioned spawn survey types: surface (section 5), Macrocystis (section 6), and understory (section 7). Within each section, we separate each level of spatial aggregation (e.g., calculations at the quadrat, or transect level) into subsections. Finally, we combine the three spawn indices to get the total spawn index (section 8), and aggregate the total by SAR and year (Figure 3).

We developed this document while converting spawn index calculations implemented in a **Microsoft Access** database to an **R** (RCT 2017) script. We updated the calculations from a database to an **R** script for several reasons. First, the database has been used for various purposes over the last two decades and has incidental calculations that make it overly complex. Second, the database is difficult to troubleshoot because it is hard to differentiate between input (i.e., data) and derived values. Third, the **R** script is open and transparent; researchers can view and download the script and an example spawn survey database. For example, source the **R** script to implement the calculations described in this document using a small set of actual observations. Fourth, we consider it good practice to separate data from analyses. Fifth, an **R** script will facilitate future research to quantify spawn index uncertainty. Finally, an **R** script will allow us to generate dynamic documents in the spirit of reproducible research using **knitr** (Xie 2015; Marwick et al. 2018). Essentially, we have attempted to follow 'good enough' practices in scientific computing (Wilson et al. 2016).

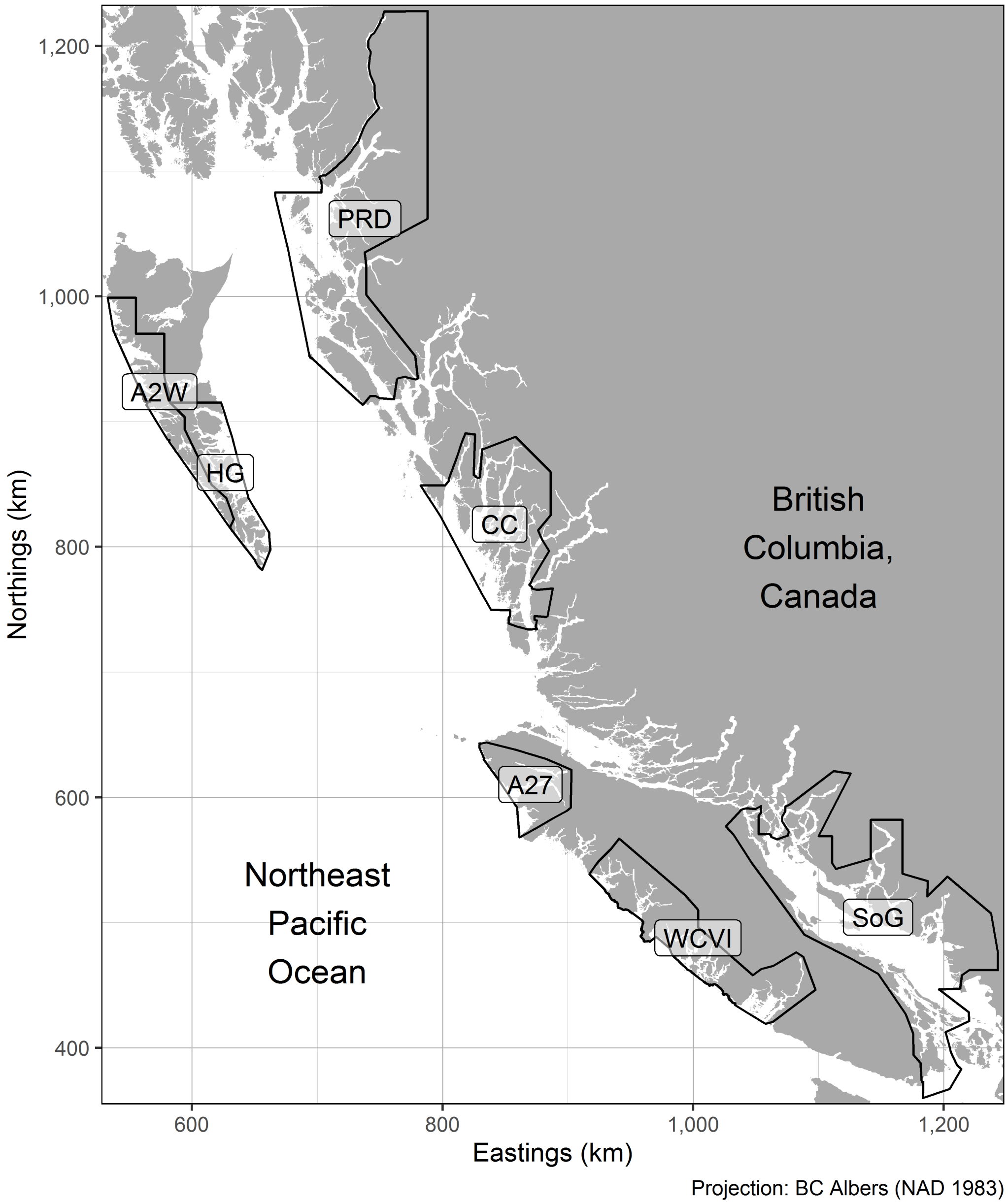
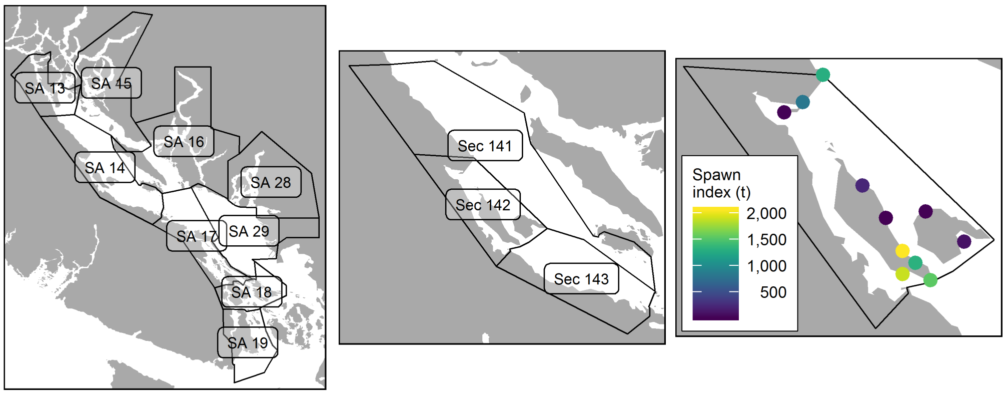


Figure 1. Spatial boundaries for Pacific Herring stock assessment regions (SARs) in British Columbia. There are five major SARs: Haida Gwaii (HG), Prince Rupert District (PRD), Central Coast (CC), Strait of Georgia (SoG), and West Coast of Vancouver Island (WCVI). There are two minor SARs: Area 27 (A27) and Area 2 West (A2W). Units: kilometres (km).



(a) Statistical Areas (SA) in the SoG SAR.

(b) Sections (Sec) in Statistical Area 14, SoG SAR.

(c) Spawn index in tonnes (t) by Location in Section 142, SoG SAR.

Figure 2. Statistical Areas (a), Sections (b), and spawn index by Location (c) for Pacific Herring spawns in 2018 in the Strait of Georgia (SoG) stock assessment region (SAR; Figure 1).

2 CONVERTING EGGS TO BIOMASS

Female Pacific Herring produce an average of approximately 200 000 eggs per kilogram (kg) of total body weight (Hay 1985; Hay and Brett 1988). We assume that females account for 50% of spawners, and we convert eggs to tonnes (t) of spawners using

$$ECF = fecundity \cdot pFemale \cdot \frac{10^3 \text{ kg}}{\text{t}} \tag{1}$$

where *fecundity* is the number of eggs per kilogram of total female body weight, *pFemale* is the proportion of spawners that are female, and *ECF* is the egg conversion factor (Table 1). Thus we convert eggs to biomass in tonnes of both sexes combined by dividing the number of eggs by *ECF*. Although Pacific Herring egg production is affected by environmental variability and other factors (Tanasichuk and Ware 1987; Hay and Brett 1988), we assume that bias to the spawn index from Equation 1 is insignificant in most areas

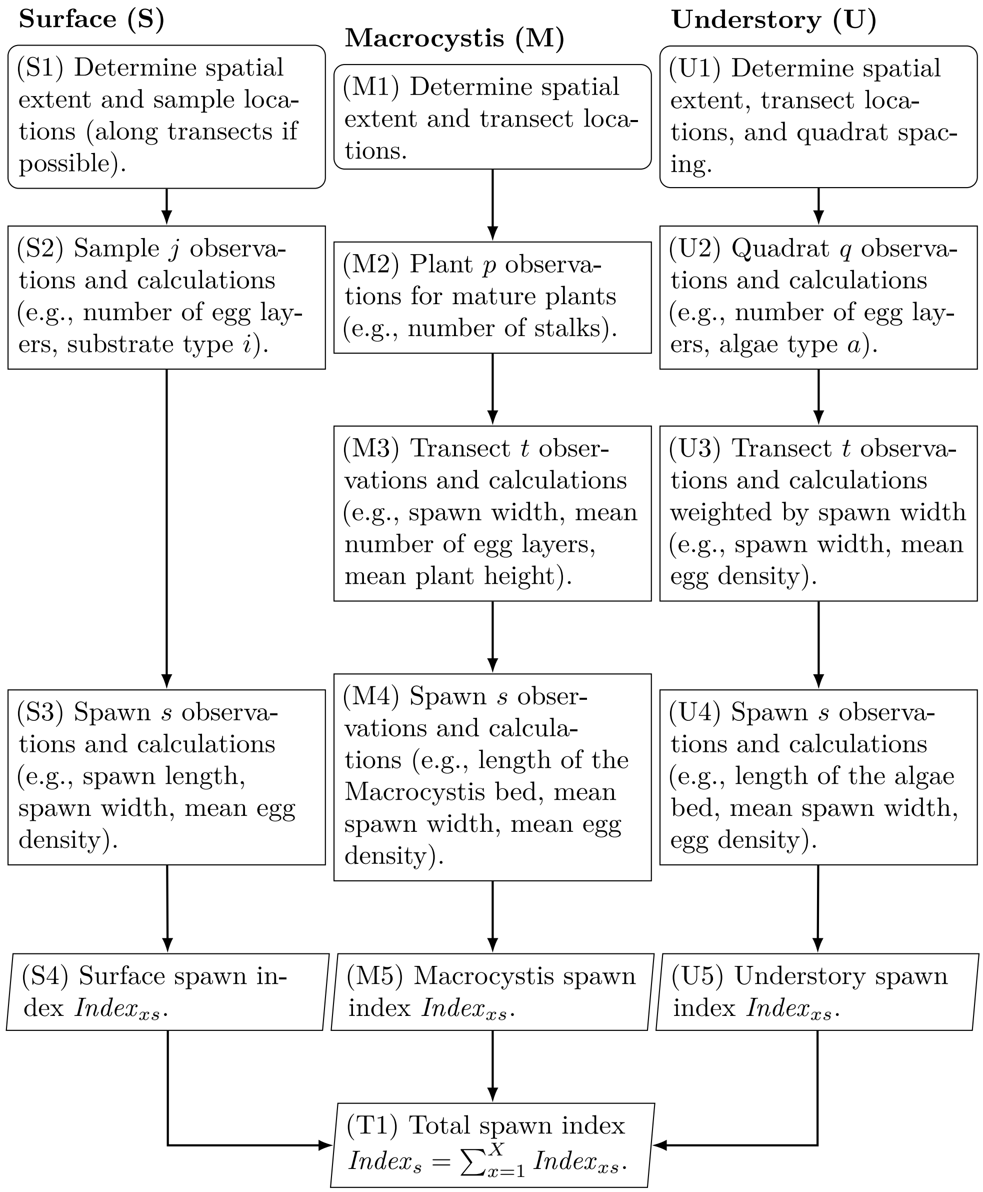


Figure 3. Sequence of steps (e.g., S1, S2) for Pacific Herring spawn index calculations for the three spawn survey types $x = \text{surface}, \text{Macrocystis}, \text{understory}$. Legend: rounded rectangles indicate start, rectangles indicate observations and calculations, parallelograms indicate output, and arrows show order of operation.

and years (Schweigert 1993; Ware 1985).

3 STATISTICAL FRAMEWORK

Historical and recent surface spawn surveys use an ad hoc sampling regimen, where surveys are often opportunistic given the state of the tide, as well as available sampling tools such as boats, rakes, and viewers. The data are analysed assuming simple random sampling, which likely generates a biased estimate of mean egg density.

In contrast, underwater dive surveys using SCUBA gear instituted in 1988 follow a two-stage systematic sampling design where transects are the first sampling stage, and individual quadrats within transects are the second sampling stage (Jessen 1978). Two steps are required to calculate mean understory egg density in each surveyed spawn s. First, mean egg density in eggs per square metre (m) for transect t in spawn s is

$$\overline{\rho_{ts}} = \frac{1}{Q}\sum_{q=1}^{Q}\rho_{qts} \tag{2}$$

where ρ_{qts} is egg density per square metre (m) for quadrat q, and Q is the number of quadrats in transect t. Before we calculate the mean egg density

Table 1. Notation for converting the number of Pacific Herring eggs to biomass. Legend: kilograms (kg), tonnes (t).

Name	Description	Value or unit
fecundity	Female fecundity	$200\,000$ eggs $\cdot$ kg^{-1}
pFemale	Proportion female	0.5
ECF	Egg conversion factor	eggs $\cdot$ 10^8 $\cdot$ t^{-1}

for spawn s, we determine the mean number of potential quadrats in spawn s

$$\overline{Q'_s} = \frac{1}{T}\sum_{t=1}^{T} Q' \tag{3}$$

where Q' is the number of potential quadrats in transect t (i.e., a function of spawn width). Then, we calculate mean egg density in eggs per square metre for spawn s

$$\overline{\rho_s} = \frac{1}{T\overline{Q'_s}}\sum_{t=1}^{T} Q'\overline{\rho_{ts}} \tag{4}$$

where T is the number of transects in spawn s, $\overline{Q'_s}$ is the mean number of potential quadrats in spawn s from Equation 3, Q' is the number of potential quadrats in transect t, and $\overline{\rho_{ts}}$ is the mean transect egg density from Equation 2. The egg density estimator from Equation 4 is unbiased, and the variance is

$$\sigma_s^2 = \frac{T'-T}{TT'}\sum_{t=1}^{T}\frac{\left(Q'\overline{\rho_{ts}} - \overline{Q'_s}\overline{\rho_s}\right)^2}{\overline{Q'_s}^2\left(T-1\right)} + \frac{f_t}{t^2}\sum_{t=1}^{T}\left(\frac{Q'}{\overline{Q'_s}}\right)^2\frac{\left(1-f_q\right)\sigma_{ts}^2}{Q} \tag{5}$$

where T' is the number of potential transects in spawn s (i.e., a function of spawn length), f_t is the transect sampling fraction for spawn s equal to $\frac{T}{T'}$, f_q is the quadrat sampling fraction for transect t equal to $\frac{Q}{Q'}$, and σ_{ts}^2 is the within transect egg density variance

$$\sigma_{ts}^2 = \frac{1}{Q-1}\sum_{q=1}^{Q}\left(\rho_{qts} - \overline{\rho_{ts}}\right)^2. \tag{6}$$

The calculation of the mean egg density for each spawn requires estimates of total spawn length, mean spawn width, length of each transect sampled, and estimated egg density in each sampling quadrat. The protocol to optimize sampling was determined through a series of studies conducted in the Strait of Georgia in 1981 and 1983 (Schweigert et al. 1985, 1990), and on the West

Coast of Vancouver Island in 1982 (Ref?). In the 1981 study, the location of transects and sampling quadrats along transects was determined using random allocation (Schweigert et al. 1985). However, this proved to be logistically difficult because neither the spawn length or width is known a priori, and divers had difficulty making the necessary calculations underwater. Nevertheless, data from these studies were used to determine a sampling protocol to estimate mean egg density with a standard error of no more than 25%. The results indicated that the sampling required to achieve this level of precision included surveying 3 transects per kilometre of spawn length, and sampling at least 5 quadrats per transect (i.e., spawn width). The sampling design was tested during a 1983 survey in the Strait of Georgia that applied a systematic rather than a random sampling protocol to simplify the logistics; variance estimates were similar to those from the 1981 study. This sampling protocol was further re-evaluated after additional surveys occurred in all areas of the coast during 1984 and 1985; the protocol was found to be robust and has been in routine use since 1988 (Schweigert et al. 1990). Although samples are collected systematically within each spawn, we assume that transects and quadrats are located randomly with respect to the underlying spawn distribution, and so these estimators are applicable. An analogous approach had previously been adopted in the sampling of various commercial fisheries where vessels arrive in port at random but are sampled in a systematic fashion to obtain a random sample (Quinn et al. 1983; Sen 1984).

Giant kelp, *Macrocystis* sp., requires a different sampling protocol than the aforementioned understory spawn survey protocol. Giant kelp routinely reach heights of 15 m, but once weighed down with herring eggs the plants can sink to lay flat on the bottom. After sampling dozens of giant kelp plants covered with herring eggs, it was determined that plant height, number of fronds per plant, and number egg layers per plant were key counts required to estimate the number of eggs per plant (Haegele and Schweigert 1990). The survey design employed to capture these data for each spawning bed rely on determining the average plant height, number of fronds in each plant holdfast, and number of giant kelp plants occurring within a 1 m swath on each side of the transect line. These data are used to determine the total egg deposition on *Macrocystis* sp. for each spawning bed (subsection 6.3).

4 SAMPLING PROTOCOL

The following is a brief summary of the spawn survey sampling protocol in the Pacific Herring spawn survey manual. Pacific Herring in BC primarily spawn in sheltered bays and inlets, depositing their eggs on rocks and algae between depths of 1.5 m above and 18 m below the 0-tide level (Humphreys and Hourston 1978; Haegele and Schweigert 1985). We identify distinct spawns (both spatially and temporally) by a unique combination of year, Location, and 'spawn number.' Spawns are numbered $s = 1, 2, 3, \ldots, S_{xlry}$ where S_{xlry} is the number of spawns in spawn survey type x (i.e., $x =$ surface, Macrocystis, understory), Location l, SAR r, and year y. A distinct spawn is a continuous stretch of shoreline with no detectable break in egg deposition; this is the finest scale at which we calculate the spawn index. A break in egg deposition is determined by the absence of Pacific Herring spawn on two consecutive transects, or by a temporal gap in spawning. Most spawns are also characterized by longitude and latitude, as well as start and end dates of spawning. Surveyors usually collect longitude and latitude at the start and end of each transect; for surface spawn surveys that don't use transects (subsection 4.1), surveyors collect longitude and latitude at the start and ends of the spawn (i.e., overall length and width).

Pacific Herring spawns typically extend along the shore; from above, spawns are identified by a milky or turquoise discolouration of the ocean caused by the release of milt, and often appear as bands running parallel to the shore (Figure 4). Thus, spawn 'length' refers to distance parallel to the shore, and 'width' refers to distance perpendicular to the shore. Similarly, Macrocystis bed length *LengthMacro* and algae bed length *LengthAlgae* refer to distances that Macrocystis and algae beds extend parallel to the shore, respectively.

Most areas of the BC coast have 'permanent transect' locations recorded on charts which enable surveyors to place transects in the same place each year. When permanent transects are unavailable for a given spawn, surveyors set new transects perpendicular to the shore, beginning 200 m in from one end of the spawn, and spaced 350 m apart along the length of the spawn. The end of the spawn is determined by the absence of eggs. We digitize new transects to make them available as permanent transects in subsequent surveys. Transects generally go from the deep edge of the spawn towards shore until divers reach the near-shore edge of the spawn; the near-shore edge can be out of the water depending on the tide height.

Pacific Herring spawn surveyors first determine the spatial extent of the spawn in terms of length of shoreline to survey (Figure 3, steps S1, M1, & U1); this is done by raking (subsection 4.1) or brief dives to determine the

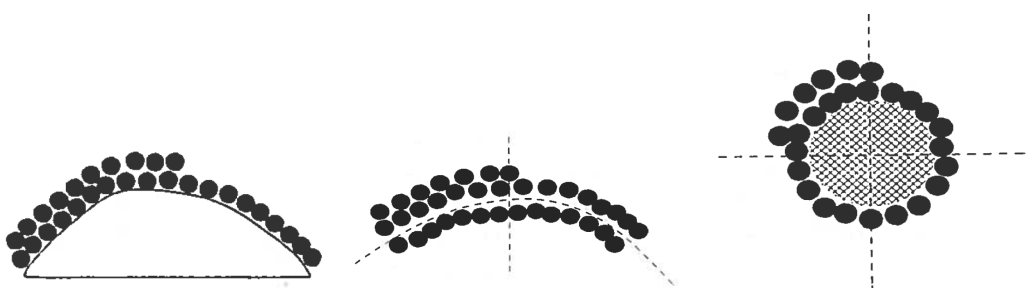
Figure 4. Aerial view of Pacific Herring spawn taken during a spawn reconnaissance flight in the Strait of Georgia. The spawn is identified by the band of discoloured water parallel to the shore.

presence or absence of spawn. Surveyors place the first transect in from one end (i.e., at the first permanent transect, or 200 m if there are no permanent transects) to avoid surveying areas with patchy and sparse egg layers. Within the spawn area, surveyors use transects to determine spawn width, quadrat placement, and which Macrocystis plants to survey. In some cases, we adjust spawn width to improve the accuracy of spawn index estimates (appendix A).

After determining the spatial extent, surveyors determine the number of egg layers on substrate and algae according to sampling protocols described in subsection 4.1, subsection 4.2, & subsection 4.3. For eggs on substrate, one egg layer is a layer of eggs one egg thick over the entire spawned surface (Figure 5a). For eggs on algae, surveyors count egg layers one of two ways depending on whether the algae is flat or round in cross-section. Egg layers on flat algae are counted on both sides of the algae (Figure 5b); egg layers on round algae are counted across the diameter of the algae (Figure 5c).

4.1 SURFACE SPAWN PROTOCOL

Surface spawn surveyors use the aforementioned transect interval when possible, but the sampling interval relies on surveyor judgement and available



(a) Substrate with 1.5 egg layers. (b) Flat algae with 2.5 egg layers. (c) Round algae with 2.5 egg layers.

Figure 5. Cross-sections showing the number of Pacific Herring egg layers on substrate (a), flat algae (b), and round algae (c). Diagrams copied with permission from the Pacific Herring spawn survey manual.

resources. If the spawn area is sufficiently large, surface surveyors usually sample along permanent transects. Small spawns can still be mapped as they were historically, with surveyors deciding how to sample the spawn. To sample, surveyors deploy specialized rakes throughout the spawn to determine algae type, number of egg layers, and percent cover. Surveyors may deploy a viewing box in shallow water, and at low tide a portion of the spawn may be visible for direct observation. We refer to these surface spawn observations as 'samples.'

Recall that there are two cases of surface spawn surveys: all surveys prior to 1988, and surveys since 1988 when dive surveys are not possible. Data from surface surveys are combined with data from dive surveys (i.e., Macrocystis, understory) to produce the total spawn index (section 8).

4.1.1 SPAWN INTENSITY CATEGORIES

From 1928 to 1978, surface spawn surveyors categorized spawn by subjective 'intensity' categories instead of directly estimating the number of egg layers (Table 2). From 1928 to 1968 there were five intensity categories described as very light, light, medium, heavy, and very heavy (numbered 1 to 5, respectively). Starting in 1969 there were nine intensity categories; the change from five to nine intensity categories was probably to accommodate the practice of reporting intermediate categories such as 3.5 (Hay and Kronlund 1987). Starting in 1979, spawn surveyors estimated the number of egg layers directly, and they continued to record intensity categories until 1981 to provide overlap between the two methods. In addition to the number of egg layers, intensity was sometimes recorded after being officially discontinued in 1981. We have converted spawn intensity observations in the Pacific Herring spawn survey database from five to nine categories for spawns that used the five category scale between 1951 and 1968. Thus, spawn data used for stock assessments is represented either by a nine category intensity scale, or a direct estimate of the number of egg layers.

Table 2. Spawn intensity descriptions, categories, and number of egg layers for Pacific Herring surface spawn surveys for the periods 1928 to 1968, and 1969 to 1978 (Hay and Kronlund 1987; Schweigert and Stocker 1988).

	Intensity category		
Description	1928 to 1968	1969 to 1978	Number of egg layers
Very light	1	1	0.5529
		2	0.9444
Light	2	3	1.3360
		4	2.1496
Medium	3	5	2.9633
		6	4.1318
Heavy	4	7	5.3002
		8	6.5647
Very heavy	5	9	7.8291

4.2 MACROCYSTIS SPAWN PROTOCOL

Macrocystis spawn surveyors take a census of Macrocystis plants within 1 m of the transect line, on both the left- and right-hand sides. We refer to the swath of substrate along Macrocystis transects as the transect swath, $Swath_{ts} = 2\,\mathrm{m}$ in transect t and spawn s. Divers categorize Macrocystis plants as either 'mature' or 'immature' based on stipe height; mature plants have stipes $\geq 1\,\mathrm{m}$ high, and are the only plants used for Macrocystis spawn index calculations. Immature plants are excluded because Pacific Herring spawn on Macrocystis fronds, not stipes; immature plants have limited fronds and slimy stipes that prevent egg adhesion. In addition, Pacific Herring typically deposit spawn higher up Macrocystis plants. For each mature plant, divers record the number of stalks. For each transect, divers record the average number of egg layers, and average plant height. Haegele and Schweigert (1990) provide a description of the sampling technique, and the basis for estimating the total number of eggs per plant.

4.3 UNDERSTORY SPAWN PROTOCOL

Understory spawn surveyors place quadrats along transects, with a target frequency of ≥ 5 quadrats per transect, given a minimum spacing of 2 m and a maximum spacing of 40 m. Similar to how the first transect is moved in from one end of the spawn, the first quadrat is moved in from the edge of the spawn to the first 5 m mark on the transect line to avoid surveying areas with patchy and sparse egg layers. Note that transect line position along permanent transects varies year to year: spawn location causes transects to shift seaward or shoreward, and spawn width causes transects to be shorter or longer. Understory spawn surveys use 0.5 m^2 quadrats; other sizes (e.g., 0.25 and 1.0 m^2) have been used for research, but are not used to calculate the spawn index (Schweigert 1993). Within each quadrat, divers record the dominant (i.e., most heavily spawned) substrate type, percentage of the quadrat covered by spawn, and number of egg layers. In addition, divers identify the three most abundant algae types that have spawn. For each of these algae types, divers record the percentage of the quadrat covered by the algae and number of egg layers.

5 SURFACE SPAWN CALCULATIONS

This section describes steps S2 to S4 in Figure 3. We simplify index notation in this section by suppressing subscripts for spawn survey type x, Location l, SAR r, and year y (Table 3). Surface spawn surveyors sample along transects or using their judgement. Surveyors collect data for at the substrate type i, sample j, and spawn s levels; we calculate metrics at the substrate type i, sample j, and spawn s levels (Table 4). Recall that surface spawn 'samples' include observations collected using specialized rakes and viewing boxes (subsection 4.1). Occasionally, we use field data sheets to fill-in missing egg layer information for surface survey data (appendix B).

Table 3. Index notation for Pacific Herring spawn index calculations. Legend: stock assessment region (SAR), spawn on kelp (SOK).

Name	Description	Value
i	Substrate type	1, 2, 3, ..., I
I	Number of substrate types in sample j, type x, spawn s, Location l, SAR r, and year y	
j	Sample	1, 2, 3, ..., J
J	Number of samples in type x, spawn s, Location l, SAR r, and year y	
p	Plant	1, 2, 3, ..., P
P	Number of plants in transect t, type x, spawn s, Location l, SAR r, and year y	
t	Transect	1, 2, 3, ..., T
T	Number of transects in type x, spawn s, Location l, SAR r, and year y	
a	Algae type	1, 2, 3, ..., A
A	Number of algae types in quadrat q, transect t, type x, spawn s, Location l, SAR r, and year y	
q	Quadrat	1, 2, 3, ..., Q
Q	Number of quadrats in transect t, type x, spawn s, Location l, SAR r, and year y	
x	Spawn survey type	1, 2, 3, ..., X
X	Number of spawn survey types in spawn s, location l, SAR r, and year y	
s	Spawn	1, 2, 3, ..., S
S	Number of spawns in Location l, SAR r, and year y	
l	Location	1, 2, 3, ..., L
L	Number of locations in SAR r and year y	
f	SOK fishery	1, 2, 3, ..., F
F	Number of SOK fisheries in SAR r and year y	
r	SAR	1, 2, 3, ..., R
R	Number of SARs	
y	Year	y_1, y_2, y_3, ..., Y
Y	Last year of time series	

Table 4. Notation for Pacific Herring surface spawn index calculations. We simplify index notation by suppressing subscripts for spawn survey type x, Location l, stock assessment region (SAR) r, and year y. Legend: metres (m), tonnes (t).

Name	Description	Value or unit
$Layers_{ijs}$	Number of egg layers on substrate i in sample j and spawn s	Number of egg layers
$Proportion_{ijs}$	Proportion of substrate i covered in eggs in sample j and spawn s	$(0, 1]$
$EggLyrs_{ijs}$	Number of egg layers on substrate i in sample j and spawn s	Number of egg layers
$EggLyrs_{js}$	Number of egg layers in sample j and spawn s	Number of egg layers
α	Regression intercept	14.698 eggs $\cdot\ 10^3 \cdot \mathrm{m}^{-2}$
β	Regression slope	212.218 eggs $\cdot\ 10^3 \cdot \mathrm{m}^{-2}$
$EggDens_{js}$	Egg density in sample j and spawn s	eggs $\cdot\ 10^3 \cdot \mathrm{m}^{-2}$
$\overline{EggDens_s}$	Mean egg density in spawn s	eggs $\cdot\ 10^3 \cdot \mathrm{m}^{-2}$
$Length_s$	Length of spawn s	m
$Width_s$	Width of spawn s	m
$Index_s$	Surface spawn index for spawn s	t

5.1 SAMPLE OBSERVATIONS AND CALCULATIONS

Each sample j (Figure 3, step S2) can have one or more substrate types i. The number of egg layers in substrate i, sample j, and spawn s is

$$EggLyrs_{ijs} = Layers_{ijs} \cdot Proportion_{ijs} \tag{7}$$

where $Layers_{ijs}$ is the number of egg layers on substrate i, and $Proportion_{ijs}$ is the proportion of substrate i covered with spawn. The total number of egg layers in sample j and spawn s is

$$EggLyrs_{js} = \sum_{i=1}^{I} EggLyrs_{ijs} \tag{8}$$

where $EggLyrs_{ijs}$ is the number of egg layers in substrate i from Equation 7. For the time period when surveyors recorded spawn 'intensity' categories instead of direct egg layer estimates, we convert intensity to number of egg layers $EggLyrs_{js}$ (Table 2). Schweigert et al. (1997) developed a predictive model of surface egg density as a function of number of egg layers using a linear regression model[1]

$$EggDens_{js} = \alpha + \beta \cdot EggLyrs_{js} \tag{9}$$

where α is the regression intercept, β is the regression slope, $EggLyrs_{js}$ is the total number of egg layers in sample j from Equation 8, and $EggDens_{js}$ is in $\text{eggs} \cdot 10^3 \cdot \text{m}^{-2}$ (Figure 6). Note that we only calculate $EggDens_{js}$ if $EggLyrs_{js} > 0$.

5.2 SPAWN OBSERVATIONS AND CALCULATIONS

The mean egg density in spawn s (Figure 3, step S3) is

$$\overline{EggDens_s} = \frac{\sum_{j=1}^{J} EggDens_{js}}{J} \tag{10}$$

where $EggDens_{js}$ is egg density in sample j from Equation 9, J is the number of samples, and $\overline{EggDens_s}$ is in $\text{eggs} \cdot 10^3 \cdot \text{m}^{-2}$. Two other metrics are required at the spawn level: spawn length $Length_s$, and spawn width $Width_s$, both in metres. We set $Width_s$ to the first non-missing value of median pool width, median Section width, median SAR width, or observed width (in that order;

[1] There is an error in Schweigert et al. (1997); surface egg density is in thousands per square metre ($\text{eggs} \cdot 10^3 \cdot \text{m}^{-2}$). Likewise, we report eggs in thousands (i.e., $\text{eggs} \cdot 10^3$) in this document and in the **R** script.

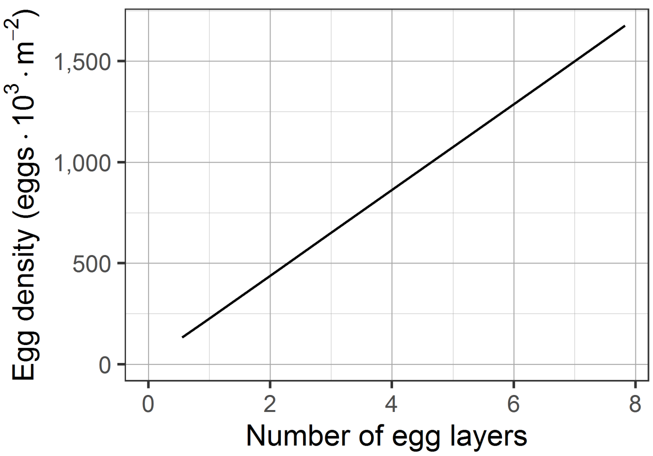


Figure 6. Egg density in thousands of eggs per square metre (m) as a function of number of egg layers for Pacific Herring surface spawn surveys (Equation 9; Schweigert et al. 1997). Note that number of egg layers can exceed those shown in this figure; values shown are for demonstration only.

subsection A.1). The surface spawn index in spawn s is

$$Index_s = \frac{\overline{EggDens_s} \cdot Length_s \cdot Width_s \cdot 10^3}{ECF} \tag{11}$$

where $\overline{EggDens_s}$ is mean egg density in spawn s from Equation 10, $Length_s$ is length of spawn s, $Width_s$ is width of spawn s, ECF is the egg conversion factor from Equation 1, and $Index_s$ is in tonnes (Figure 3, step S4).

6 MACROCYSTIS SPAWN CALCULATIONS

This section describes steps M2 to M5 in Figure 3. As with the previous section, we simplify index notation in this section by suppressing subscripts for spawn survey type x, Location l, SAR r, and year y. Macrocystis spawn surveyors use SCUBA gear to collect underwater data for individual plants p, transects t, and spawns s; we calculate metrics at the transect t, and spawn s levels (Table 5).

Table 5. Notation for Pacific Herring Macrocystis spawn index calculations. We simplify index notation by suppressing subscripts for spawn survey type x, Location l, stock assessment region (SAR) r, and year y. Legend: metres (m), tonnes (t).

Name	Description	Value or unit
$Stalks_{pts}$	Number of stalks on plant p in transect t in spawn s	Number of stalks
$Width_{ts}$	Width of transect t in spawn s	m
$Swath_{ts}$	Swath of transect t in spawn s	m
$Area_{ts}$	Area of transect t in spawn s	m^2
$\overline{Height_{ts}}$	Mean plant height on transect t in spawn s	m
$\overline{EggLyrs_{ts}}$	Mean number of egg layers on transect t in spawn s	Number of egg layers
$Stalks_{ts}$	Number of stalks on transect t in spawn s	Number of stalks
$LengthMacro_s$	Length of the Macrocystis bed in spawn s	m
$Length_s$	Length of spawn s	m
$\overline{Width_s}$	Mean width of spawn s	m
$Area_s$	Area of spawn s	m^2
$Plants_s$	Number of plants in spawn s	Number of plants
$Stalks_s$	Number of stalks in spawn s	Number of stalks
$\overline{Height_s}$	Mean plant height in spawn s	m
$\overline{EggLyrs_s}$	Mean number of egg layers in spawn s	Number of egg layers
$\overline{StalksPerPlant_s}$	Mean number of stalks per plant in spawn s	Number of stalks per plant
β	Regression slope	$0.073\ \mathrm{eggs} \cdot 10^3 \cdot \mathrm{plant}^{-1}$
γ	Regression exponent on $\overline{EggLyrs_s}$	0.673
δ	Regression exponent on $\overline{Height_s}$	0.932
ϵ	Regression exponent on $\overline{StalksPerPlant_s}$	0.703
$\overline{EggsPerPlant_s}$	Mean number of eggs per plant in spawn s	$\mathrm{eggs} \cdot 10^3 \cdot \mathrm{plant}^{-1}$
$\overline{EggDens_s}$	Mean egg density in spawn s	$\mathrm{eggs} \cdot 10^3 \cdot \mathrm{m}^{-2}$
$Index_s$	Macrocystis spawn index for spawn s	t

6.1 PLANT OBSERVATIONS

For each mature plant p in transect t and spawn s (Figure 3, step M2), surveyors count the number of stalks $Stalks_{pts}$.

6.2 TRANSECT OBSERVATIONS AND CALCULATIONS

At the transect t level (Figure 3, step M3), spawn width is $Width_{ts}$, and transect swath is $Swath_{ts} = 2\,\text{m}$, both in metres. We calculate the area in transect t and spawn s as

$$Area_{ts} = Width_{ts} \cdot Swath_{ts} \tag{12}$$

in square metres. In addition, divers estimate summary statistics for mature Macrocystis plants along transect t in spawn s: mean height $\overline{Height_{ts}}$ in metres, and mean number of egg layers $\overline{EggLyrs_{ts}}$. The total number of plants in transect t and spawn s is P. We calculate the total number of stalks in transect t and spawn s

$$Stalks_{ts} = \sum_{p=1}^{P} Stalks_{pts} \tag{13}$$

where $Stalks_{pts}$ is the number of stalks on plant p.

6.3 SPAWN OBSERVATIONS AND CALCULATIONS

At the spawn s level (Figure 3, step M4), we determine the length of the Macrocystis bed $LengthMacro_s$ in metres. If $LengthMacro_s$ is inadvertently not recorded, we set $LengthMacro_s$ to the spawn length $Length_s$. The mean width of spawn s is

$$\overline{Width_s} = \frac{\sum_{t=1}^{T} Width_{ts}}{T} \tag{14}$$

where $Width_{ts}$ is the transect width, T is the number of transects in spawn s, and $\overline{Width_s}$ is in metres. The total area of transects in spawn s is

$$Area_s = \sum_{t=1}^{T} Area_{ts} \tag{15}$$

where $Area_{ts}$ is the transect area from Equation 12, and $Area_s$ is in square metres. The total number of stalks in spawn s is

$$Stalks_s = \sum_{t=1}^{T} Stalks_{ts} \tag{16}$$

where $Stalks_{ts}$ is the number of stalks in transect t from Equation 13. The total number of plants in spawn s is

$$Plants_s = \sum_{t=1}^{T} P \tag{17}$$

where P is the number of plants in transect t. The mean plant height in spawn s is

$$\overline{Height_s} = \frac{\sum_{t=1}^{T} \overline{Height_{ts}}}{T} \tag{18}$$

where $\overline{Height_{ts}}$ is the mean plant height in transect t, T is the number of transects in spawn s, and $\overline{Height_s}$ is in metres. The mean number of egg layers in spawn s is

$$\overline{EggLyrs_s} = \frac{\sum_{t=1}^{T} \overline{EggLyrs_{ts}}}{T} \tag{19}$$

where $\overline{EggLyrs_{ts}}$ is the mean number of egg layers in transect t, and T is the number of transects in spawn s. The mean number of stalks per plant in spawn s is

$$\overline{StalksPerPlant_s} = \frac{Stalks_s}{Plants_s} \tag{20}$$

where $Stalks_s$ is the number of stalks in spawn s from Equation 16, and $Plants_s$ is the number of plants in spawn s from Equation 17.

Haegele and Schweigert (1990) developed a predictive model of number of eggs per plant as a function of number of egg layers on plants, plant height, and number of stalks per plant using a nonlinear multiple regression model

$$\overline{EggsPerPlant_s} = \beta \cdot \overline{EggLyrs_s}^{\gamma} \cdot \overline{Height_s}^{\delta} \cdot \overline{StalksPerPlant_s}^{\epsilon} \cdot 10^3 \qquad (21)$$

where β is the regression slope, $\overline{EggLyrs_s}$ is the mean number of egg layers in spawn s from Equation 19, γ is the regression exponent on $\overline{EggLyrs_s}$, $\overline{Height_s}$ is the mean plant height in spawn s from Equation 18, δ is the regression exponent on $\overline{Height_s}$, $\overline{StalksPerPlant_s}$ is the mean number of stalks per plant in spawn s from Equation 20, ϵ is the regression exponent on $\overline{StalksPerPlant_s}$, and $\overline{EggsPerPlant_s}$ is in eggs $\cdot\, 10^3 \cdot \text{plant}^{-1}$ (Figure 7). Mean macrocystis egg density in spawn s is

$$\overline{EggDens_s} = \frac{\overline{EggsPerPlant_s} \cdot Plants_s}{Area_s} \qquad (22)$$

where $\overline{EggsPerPlant_s}$ is the mean number of eggs per plant in spawn s from Equation 21, $Plants_s$ is the number of plants in spawn s from Equation 17, $Area_s$ is the total area of transects in spawn s from Equation 15, and $\overline{EggDens_s}$ is in eggs $\cdot\, 10^3 \cdot \text{m}^{-2}$. The Macrocystis spawn index in spawn s is

$$Index_s = \frac{\overline{EggDens_s} \cdot LengthMacro_s \cdot \overline{Width_s} \cdot 10^3}{ECF} \qquad (23)$$

where $\overline{EggDens_s}$ is the mean egg density in spawn s from Equation 22, $LengthMacro_s$ is the length of the Macrocystis bed in spawn s, $\overline{Width_s}$ is the mean width of spawn s from Equation 14, ECF is the egg conversion factor from Equation 1, and $Index_s$ is in tonnes (Figure 3, step M5).

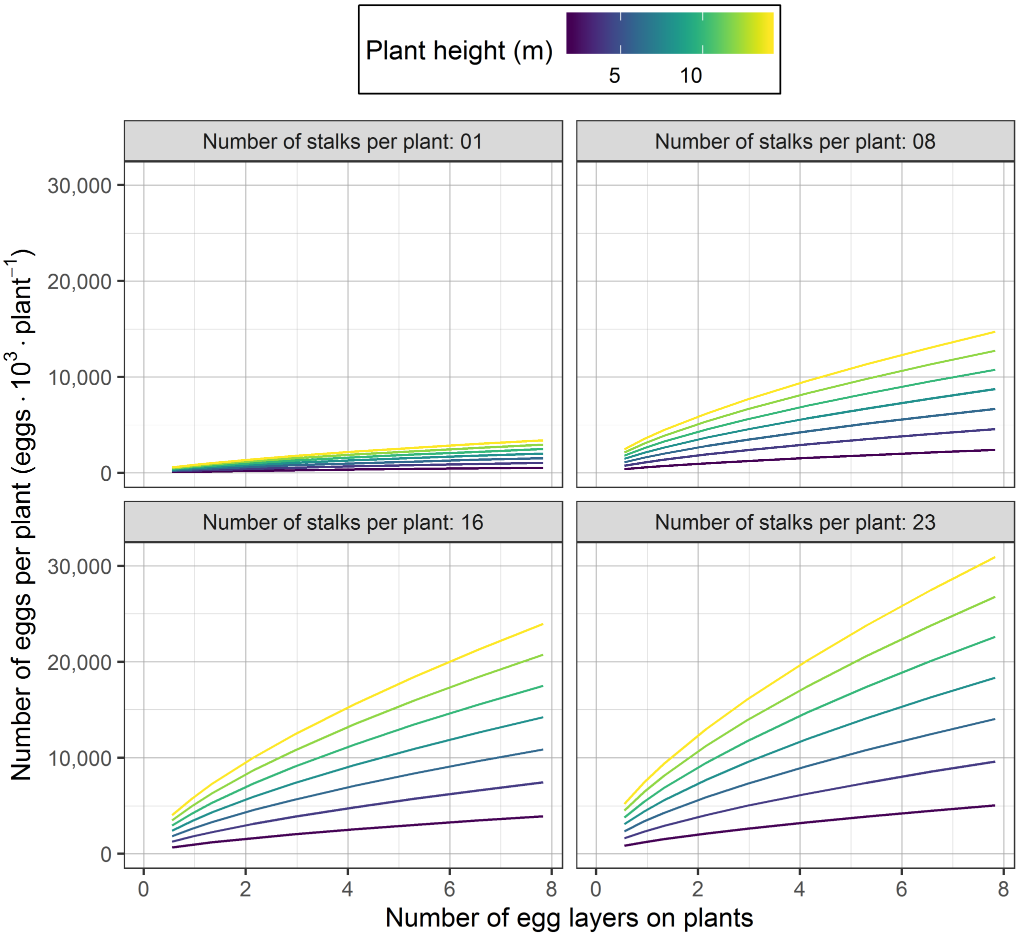


Figure 7. Number of eggs in thousands per Macrocystis plant as a function of number of egg layers on plants, plant height in metres (m), and number of stalks per plant for Pacific Herring Macrocystis spawn surveys (Equation 21; Haegele and Schweigert 1990). Note that number of egg layers, plant height, and number of stalks per plant can exceed those shown in this figure; values shown are for demonstration only.

7 UNDERSTORY SPAWN CALCULATIONS

This section describes steps U2 to U5 in Figure 3. As with the previous two sections, we simplify index notation in this section by suppressing subscripts for spawn survey type x, Location l, SAR r, and year y. Understory spawn surveyors use SCUBA gear to collect underwater data for algae types a, quadrats q, transects t, and spawns s; we calculate metrics at the algae type a, quadrat q, transect t, and spawn s levels (Table 6).

7.1 QUADRAT OBSERVATIONS AND CALCULATIONS

We calculate two separate estimates of egg density at the quadrat level (Figure 3, step U2): spawn on substrate, and spawn on algae a. Haegele et al. (1979) developed a predictive model of substrate egg density in quadrat q, transect t, and spawn s from egg layers using a linear regression model

$$EggDensSub_{qts} = \alpha \cdot SubLyrs_{qts} \cdot SubProp_{qts} \tag{24}$$

where α is the regression slope, $SubLyrs_{qts}$ is the number of egg layers on substrate in quadrat q, $SubProp_{qts}$ is the proportion of substrate in quadrat q covered by spawn, and $EggDensSub_{qts}$ is substrate egg density in eggs$\cdot 10^3 \cdot$m^{-2} (Figure 8).

Although quadrats have only one substrate, they can have one or more algae types a. Schweigert (2005) developed a predictive model of algae egg density from egg layers, proportion of the quadrat covered by algae, and an algae coefficient using a generalized linear model. Algae coefficients account for the effect of algae morphology on Pacific Herring egg density (Table 7). Egg density on algae a in quadrat q, transect t, and spawn s (Schweigert 2005) is

$$EggDensAlg_{aqts} = \beta \cdot AlgLyrs_{aqts}^{\gamma} \cdot AlgProp_{aqts}^{\delta} \cdot Coef_a \tag{25}$$

where β is the regression slope, $AlgLyrs_{aqts}$ is the number of egg layers on algae a, γ is the regression exponent on $AlgLyrs_{aqts}$, $AlgProp_{aqts}$ is the proportion of quadrat q covered by algae a, δ is the regression exponent on $AlgProp_{aqts}$, $Coef_a$ is the coefficient for algae a, and $EggDensAlg_{aqts}$ is in eggs $\cdot 10^3 \cdot$ m^{-2} (Figure 9). The total algae egg density for quadrat q in transect t and spawn

Table 6. Notation for Pacific Herring understory spawn index calculations. We simplify index notation by suppressing subscripts for spawn survey type x, Location l, stock assessment region (SAR) r, and year y. Legend: metres (m), tonnes (t).

Name	Description	Value or unit
$SubLyrs_{qts}$	Number of substrate egg layers in quadrat q, transect t, and spawn s	Number of egg layers
$SubProp_{qts}$	Proportion of substrate covered in eggs in quadrat q, transect t, and spawn s	$(0, 1]$
α	Regression slope	340 eggs $\cdot 10^3 \cdot \mathrm{m}^{-2}$
$EggDensSub_{qts}$	Substrate egg density in quadrat q, transect t, and spawn s	eggs $\cdot 10^3 \cdot \mathrm{m}^{-2}$
β	Regression slope	600.567 eggs$\cdot 10^3 \cdot \mathrm{m}^{-2}$
$AlgLyrs_{aqts}$	Number of egg layers on algae a in quadrat q, transect t, and spawn s	Number of egg layers
γ	Regression exponent on $AlgLyrs_{aqts}$	0.6355
$AlgProp_{aqts}$	Proportion of algae a covered in eggs in quadrat q, transect t, and spawn s	$(0, 1]$
δ	Regression exponent on $AlgProp_{aqts}$	1.4130
$Coef_a$	Coefficient for algae a	see Table 7
$EggDensAlg_{aqts}$	Algae a egg density in quadrat q, transect t, and spawn s	eggs $\cdot 10^3 \cdot \mathrm{m}^{-2}$
$EggDensAlg_{qts}$	Algae egg density in quadrat q, transect t, and spawn s	eggs $\cdot 10^3 \cdot \mathrm{m}^{-2}$
$EggDens_{qts}$	Egg density in quadrat q, transect t, and spawn s	eggs $\cdot 10^3 \cdot \mathrm{m}^{-2}$
$Width_{ts}$	Width of transect t in spawn s	m
$\overline{EggDensL}_{ts}$	Mean linear weighted egg density in transect t and spawn s	eggs $\cdot 10^3 \cdot \mathrm{m}^{-1}$
$Width_s$	Sum of spawn s widths	m
$\overline{Width}_s$	Mean width of spawn s	m
$LengthAlgae_s$	Length of the algae bed in spawn s	m
$Length_s$	Length of spawn s	m
$EggDensL_s$	Sum of linear weighted egg density in spawn s	eggs $\cdot 10^3 \cdot \mathrm{m}^{-1}$
$EggDens_s$	Egg density in spawn s	eggs $\cdot 10^3 \cdot \mathrm{m}^{-2}$
$Index_s$	Understory spawn index in spawn s	t

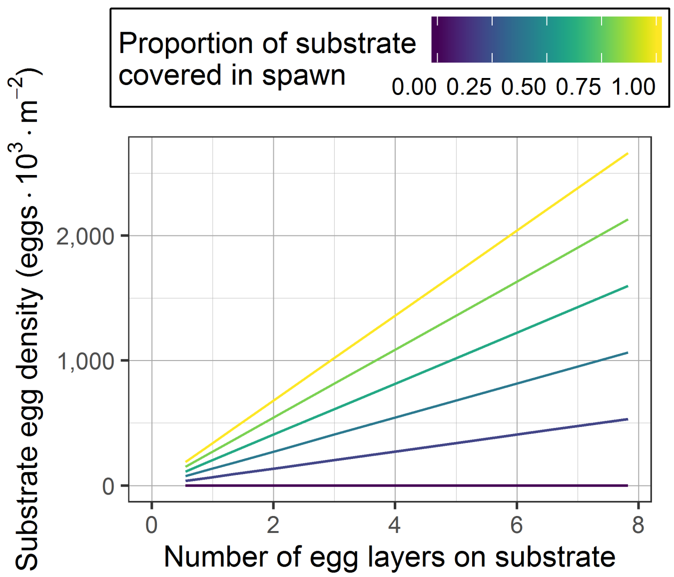


Figure 8. Substrate egg density in thousands of eggs per square metre (m) as a function of number of egg layers on substrate and proportion of substrate covered in spawn for Pacific Herring underwater spawn surveys (Equation 24; Haegele et al. 1979). Note that number of egg layers can exceed those shown in this figure; values shown are for demonstration only.

s is

$$EggDensAlg_{qts} = \sum_{a=1}^{A} EggDensAlg_{aqts} \tag{26}$$

where $EggDensAlg_{aqts}$ is egg density in algae a from Equation 25, and $EggDensAlg_{qts}$ is in eggs $\cdot\, 10^3 \cdot \mathrm{m}^{-2}$.

The total understory egg density for quadrat q in transect t and spawn s is

$$EggDens_{qts} = EggDensSub_{qts} + EggDensAlg_{qts} \tag{27}$$

where $EggDensSub_{qts}$ is substrate egg density from Equation 24, $EggDensAlg_{qts}$ is algae egg density from Equation 26, and $EggDens_{qts}$ is in eggs $\cdot\, 10^3 \cdot \mathrm{m}^{-2}$. Thus, we assume that eggs on substrate and algae are independent, and can be safely added without bias.

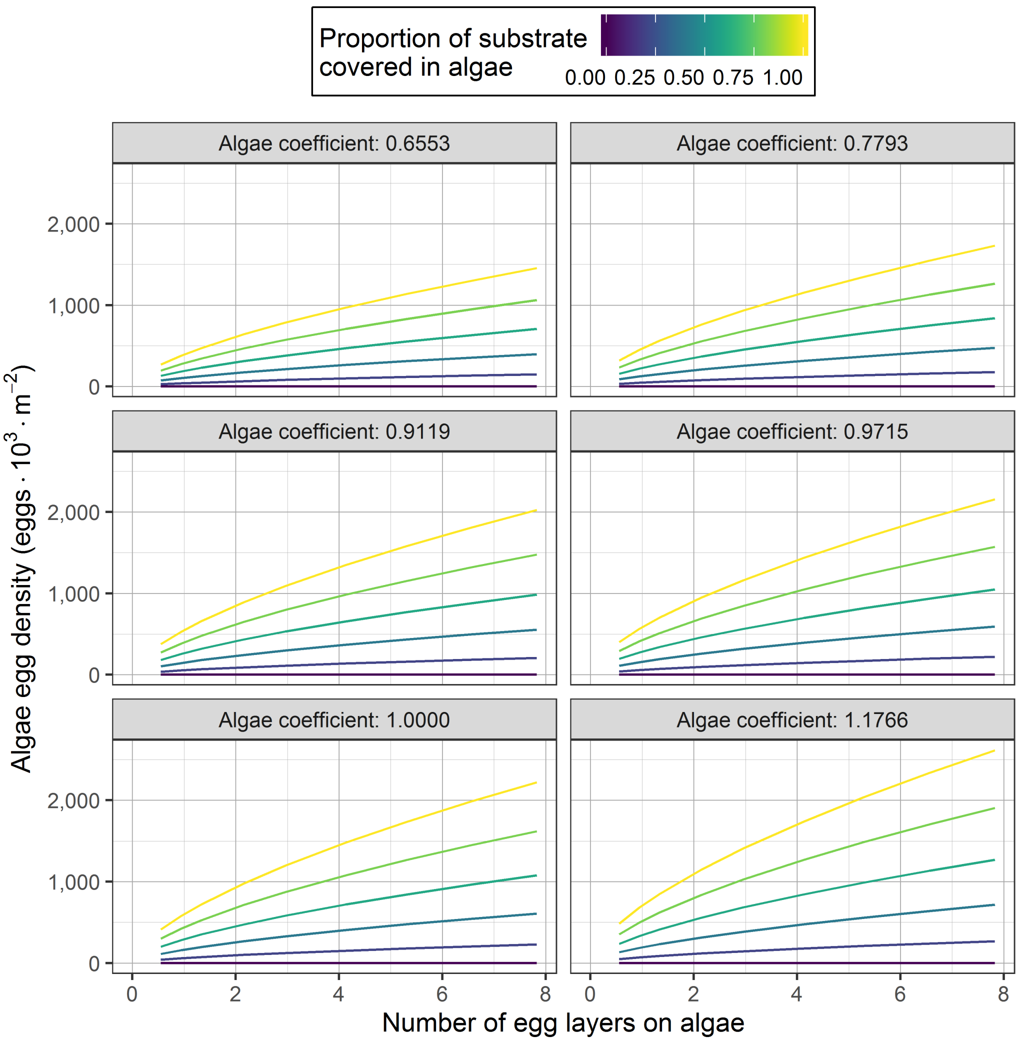


Figure 9. Algae egg density in thousands of eggs per square metre (m) as a function of number of egg layers on algae, proportion of substrate covered in algae, and algae coefficient (Table 7) for Pacific Herring underwater spawn surveys (Equation 25; Schweigert 2005). Note that number of egg layers can exceed those shown in this figure; values shown are for demonstration only.

Table 7. Algae types a and coefficients $Coef$ for Pacific Herring understory spawn surveys (Schweigert 2005).

Algae type a	Coefficient $Coef$
Grasses	0.9715
Grunge	1.0000
Kelp (flat)	0.9119
Kelp (standing)	1.1766
Leafy algae	0.6553
Rockweed	0.7793
Sargassum	1.1766
Stringy algae	1.0000

7.2 TRANSECT OBSERVATIONS AND CALCULATIONS

At the transect level (Figure 3, step U3), the mean linear understory egg density in transect t and spawn s is

$$\overline{EggDensL_{ts}} = \frac{\sum_{q=1}^{Q} EggDens_{qts}}{Q} \cdot Width_{ts} \tag{28}$$

where $EggDens_{qts}$ is total understory egg density in quadrat q from Equation 27, Q is the number of quadrats in transect t, $Width_{ts}$ is the spawn width in metres, and $\overline{EggDensL_{ts}}$ is in eggs $\cdot\, 10^3 \cdot \mathrm{m}^{-1}$. We calculate a weighted mean egg density because spawn width can vary along the spawn length; a weighted mean ensures that transects contribute proportionally to their area. Note that we update spawn width to correct for errors regarding the assumed accuracy of transect lines used to measure spawn width for understory surveys between 2003 and 2014 (subsection A.2).

7.3 SPAWN OBSERVATIONS AND CALCULATIONS

At the spawn level (Figure 3, step U4), the sum of spawn widths in spawn s is

$$Width_s = \sum_{t=1}^{T} Width_{ts} \tag{29}$$

where $Width_{ts}$ is the spawn width in transect t, and $Width_s$ is in metres. The mean width of spawn s is

$$\overline{Width_s} = \frac{Width_s}{T} \tag{30}$$

where $Width_s$ is the sum of widths from Equation 29, T is the number of transects in spawn s, and $\overline{Width_s}$ is in metres. The length of the algae bed in spawn s is $LengthAlgae_s$, also in metres. As with Macrocystis spawn calculations, if $LengthAlgae_s$ is inadvertently not recorded, we set $LengthAlgae_s$ to the spawn length $Length_s$. Thus, we assume that eggs on the substrate and eggs on algae are represented by the same length measurement. The sum of transect egg densities in spawn s is

$$EggDensL_s = \sum_{t=1}^{T} \overline{EggDensL_{ts}} \tag{31}$$

where $\overline{EggDensL_{ts}}$ is the mean linear weighted understory egg density in transect t from Equation 28, and $EggDensL_s$ is in eggs $\cdot\, 10^3 \cdot \mathrm{m}^{-1}$. Next we convert from linear density to area density. Understory egg density in spawn s is

$$EggDens_s = \frac{EggDensL_s}{Width_s} \tag{32}$$

where $EggDensL_s$ is the sum of transect egg densities in spawn s from Equation 31, $Width_s$ is the sum of spawn widths in spawn s from Equation 29, and

$EggDens_s$ is in eggs $\cdot\, 10^3 \cdot \mathrm{m}^{-2}$. The understory spawn index in spawn s is

$$Index_s = \frac{EggDens_s \cdot LengthAlgae_s \cdot \overline{Width_s} \cdot 10^3}{ECF} \tag{33}$$

where $EggDens_s$ is understory egg density from Equation 32, $LengthAlgae_s$ is the length of the algae bed, $\overline{Width_s}$ is the mean spawn width from Equation 30, ECF is the egg conversion factor from Equation 1, and $Index_s$ is in tonnes (Figure 3, step U5).

8 TOTAL SPAWN CALCULATIONS

This section describes step T1 in Figure 3. Unlike the previous three sections, we include subscripts for spawn survey type x, Location l, SAR r, and year y in the equations in this section (Table 8). The total spawn index in spawn s, Location l, region r, and year y is

$$Index_{slry} = \sum_{x=1}^{X} Index_{xslry} \tag{34}$$

where $Index_{xslry}$ is spawn index for surface, Macrocystis, and understory spawn surveys from Equation 11, Equation 23, and Equation 33, respectively, and $Index_{slry}$ is the total spawn index in tonnes (Figure 3, step T1). Finally, we aggregate the total spawn index by SAR r and year y

$$Index_{ry} = \sum_{l=1}^{L} \sum_{s=1}^{S} Index_{slry} \tag{35}$$

where $Index_{slry}$ is the total spawn index from Equation 34, and $Index_{ry}$ is a relative index of combined sex spawning biomass for SAR r and year y in tonnes. We use $Index_{ry}$ as an indicator of Pacific Herring relative population abundance in stock assessment models.

Table 8. Notation for Pacific Herring total spawn index calculations. Legend: stock assessment region (SAR), tonnes (t).

Name	Description	Value or unit
$Index_{xslry}$	Spawn index for spawn survey type x in spawn s, Location l, SAR r, and year y	t
$Index_{slry}$	Total spawn index in spawn s, Location l, SAR r, and year y	t
$Index_{ry}$	Total spawn index in SAR r and year y	t

9 SPAWN ON KELP CALCULATIONS

Spawn on kelp (SOK) fisheries collect Pacific Herring roe that adhere to algae such as Macrocystis after spawning. Other similar fisheries include spawn on bough, in which operators collect roe that adhere to tree boughs; we refer to these fisheries collectively as SOK in this document. There are two types of SOK fisheries in BC: 'open-pond' in which operators provide algae to spawning Pacific Herring, and 'closed-pond' in which operators impound spawning Pacific Herring in floating nets that contain algae (Shields et al. 1985). Although SOK fisheries do not directly remove spawning Pacific Herring, they do remove eggs that could otherwise have contributed to recruitment. Note that closed-pond operations also cause incidental mortality to spawning Pacific Herring (Shields et al. 1985), but we do not address this issue here. Thus, SOK fisheries present an issue in terms of their impact to the population, and accounting in stock assessment and monitoring (Schweigert et al. 2018). Although Pacific Herring stock assessments do not account for eggs removed by SOK fisheries at this time, there are a few options to account for the impact of SOK harvest. The most direct is to estimate the quantity of eggs removed from the population, and treat them as though they would have spawned and contributed to spawning biomass.

Shields et al. (1985) collected information on the relationship between the number of egg layers in SOK product, and proportion of product weight that consists of eggs and kelp. They determined that kelp represents an average of 12% of the total product weight. Since SOK product is universally brined at the time of harvest, it is necessary to also consider the uptake of salt by the eggs, which increases the overall product weight. However, there is uncertainty in the degree of brining that occurs prior to weighing the product. Nevertheless, Whyte and Englar (1977) determined that following a 24 hour brining period, wet product weight increases by about 13% due to salt uptake. By osmosis, brining would also draw some water from the eggs; unfortunately we are unable to account for osmosis at this time. The last factor to consider is the mean fertilized egg weight, which was determined by Hay and Miller (1982) to be $2.38 \cdot 10^{-6}$ kg.

We estimate spawning biomass removed from the population by SOK fishery f in SAR r and year y as

$$Biomass_{fry} = \frac{SOK_{fry} \cdot eggKelpProp \cdot eggBrineProp}{eggWt \cdot ECF} \tag{36}$$

where SOK_{fry} is the weight in kilograms of Pacific Herring SOK harvest in fishery f, SAR r, and year y, *eggKelpProp* is the proportion of SOK product that is eggs, not kelp, *eggBrineProp* is the proportion of SOK product that is eggs after brining, *eggWt* is the average weight in kilograms of a fertilized egg, and $Biomass_{fry}$ is spawning biomass in tonnes for SOK fishery f (Table 9).

Then we aggregate spawning biomass by stock assessment region (SAR) r and year y

$$Biomass_{ry} = \sum_{f=1}^{F} Biomass_{fry} \tag{37}$$

where $Biomass_{fry}$ is spawning biomass by fishery f from Equation 36, and $Biomass_{ry}$ is estimated spawning biomass removed by SOK fisheries in SAR r and year y in tonnes.

Table 9. Notation for Pacific Herring spawn on kelp (SOK) calculations. Legend: stock assessment region (SAR), kilograms (kg), tonnes (t).

Name	Description	Value or unit
SOK_{fry}	Weight of SOK harvest in fishery f, SAR r, and year y	kg
$eggKelpProp$	Proportion of SOK product that is eggs, not kelp	0.88
$eggBrineProp$	Proportion of SOK product that is eggs after brining	$\frac{1}{1.13}$
$eggWt$	Average weight of a fertilized egg	$2.38 \cdot 10^{-6}$ kg
$Biomass_{fry}$	Spawning biomass in SOK fishery f, SAR r, and year y	t
$Biomass_{ry}$	Spawning biomass for SOK fisheries in SAR r and year y	t

10 SOURCES OF UNCERTAINTY

Like all biological models, spawn index calculations are affected by various potential sources of uncertainty including natural variability, observation error (e.g., bias, precision), measurement error, and model structural complexity (Link et al. 2012). Three examples illustrate these sources of uncertainty (Table 10). First, natural variability could affect Pacific Herring fecundity, and the sex ratio of spawning Pacific Herring (Equation 1). Fecundity could be influenced by biological processes such as the observed non-stationarity of weight-at-age, or a truncated age distribution caused by selective fishing or high natural mortality. Second, measurement error could affect input data such as the number of egg layers, while model structural complexity could affect estimated prediction model parameters, or the form of their relationship, or both (e.g., Equation 9). In addition, spawn index prediction models are dated, and our understanding of these processes could have changed in the intervening years. Third, fixed parameters are used as data without error (e.g., Equation 9). Despite these assumptions and potential sources of uncertainty, the spawn index has typically been reported without quantifying uncertainty (but see Schweigert et al. 1993). Reporting the spawn index without uncertainty may perpetuate the misconception that the spawn index is observed data, whereas it is derived data with assumptions and uncertainties.

There are several potential benefits to addressing spawn index uncertainty. First, quantifying uncertainty could identify parameters to target with future research. Potential analyses to quantify spawn index uncertainty include:

1. Investigate factors that influence fecundity and sex ratios (Equation 1);
2. Quantify and report variability in estimated prediction model parameters and equations (e.g., Equation 9);
3. Bootstrap observed input data (see Schweigert 1993); and
4. Conduct sensitivity analyses.

Second, acknowledging uncertainty can reduce another source of uncertainty: inadequate communication among scientists, managers, and stakeholders, which can lead to misapplication of scientific advice (Link et al. 2012). Finally, acknowledging uncertainty will increase transparency, and enable users to assess potential impacts to Pacific Herring stock assessments in a management strategy evaluation (MSE) approach (e.g., DFO 2019*a*). Addressing data and model uncertainty is a required component of MSE approaches (Punt et al. 2016).

Quantifying uncertainty may also identify options to increase survey program efficiency, in terms of data precision and accuracy. Sampling surveys trade off precision of estimated quantities versus survey effort or cost. Ideally, reducing survey effort does not result in biased target variable estimates. Therefore, understanding this trade-off is important if, for example, budget reductions cause reduced survey effort. Potential strategies to improve spawn survey efficiency include:

Table 10. Uncertainty in Pacific Herring spawn index parameters. Legend: kilograms (kg), metres (m).

Equation	Name	Description	Value or unit	Uncertainty	Reference
1	$fecundity$	Female fecundity	$200\,000$ eggs $\cdot$ kg^{-1}	Unknown	Hay (1985); Hay and Brett (1988)
1	$pFemale$	Proportion female	0.5	Unknown	Hay (1985); Hay and Brett (1988)
9	α	Regression intercept	14.698 eggs $\cdot\ 10^3 \cdot$ m^{-2}	Unknown	Schweigert et al. (1997)
9	β	Regression slope	212.218 eggs $\cdot\ 10^3 \cdot$ m^{-2}	Unknown	Schweigert et al. (1997)
21	β	Regression slope	0.073 eggs $\cdot\ 10^3 \cdot$ plant^{-1}	Unknown	Haegele and Schweigert (1990)
21	γ	Regression exponent on $\overline{EggLyrs_s}$	0.673	Unknown	Haegele and Schweigert (1990)
21	δ	Regression exponent on $\overline{Height_s}$	0.932	Unknown	Haegele and Schweigert (1990)
21	ϵ	Regression exponent on $\overline{StalksPerPlant_s}$	0.703	Unknown	Haegele and Schweigert (1990)
24	α	Regression slope	340 eggs $\cdot\ 10^3 \cdot$ m^{-2}	Unknown	Haegele et al. (1979)
25	β	Regression slope	600.567 eggs $\cdot\ 10^3 \cdot$ m^{-2}	Unknown	Schweigert (2005)
25	γ	Regression exponent on $AlgLyrs_{aqts}$	0.6533	Unknown	Schweigert (2005)
25	δ	Regression exponent on $AlgProp_{aqts}$	1.4130	Unknown	Schweigert (2005)
36	$eggKelpProp$	Proportion of SOK product that is eggs, not kelp	0.88	Unknown	Shields et al. (1985)
36	$eggBrineProp$	Proportion of SOK product that is eggs after brining	$\frac{1}{1.13}$	Unknown	Whyte and Englar (1977)
36	$eggWt$	Average weight of a fertilized egg	$2.38 \cdot 10^{-6}$ kg	Unknown	Hay and Miller (1982)

1. Conduct underwater surveys for major spawns in core areas, and surface surveys for other spawns;
2. Quantify the precision and accuracy of spawn width estimates (e.g., appendix A);
3. Review transect and quadrat spacing (section 3; see Schweigert 1993);
4. Review egg prediction model accuracy (Equation 1);
5. Review temporal stability of egg layer estimates; and
6. Conduct periodic versus annual surveys.

Even with a stable budget, there is a trade-off between high survey effort in some areas, versus low survey effort or no information in other areas.

11 FUTURE RESEARCH

Many of the parameters and prediction models used to calculate the spawn index are dated; these analyses could be checked with new information, and updated if required. Parameters include *fecundity*, *pFemale*, *eggKelpProp*, *eggBrineProp*, and *eggWt*. Prediction models include Equation 9, Equation 21, Equation 24, and Equation 25. In addition, parameter and prediction model uncertainty should be propagated through the calculations to quantify spawn index uncertainty (section 10). One approach to account for prediction model uncertainty is to incorporate the underlying data that informs these equations into spawn index calculations. In addition, future work could review the assumed statistical framework.

12 DOWNLOAD

The **R** script to calculate the Pacific Herring spawn index, `SpawnIndex.R` is publicly accessible on the Pacific Herring spawn index repository. The repository includes instructions, and an example database of Pacific Herring spawn survey observations to use with the script. When sourced, the **R** script imports tables from the database, and follows the calculations described in this document. This document is meant to accompany the **R** script, which has complete details regarding how we calculate the spawn index. Sections in this document correspond to functions in the **R** script. For example, 'Surface spawn' (section 5) follows the **R** function `CalcSurfSpawn`. In addition, variable names in this document correspond to variable names in the script. Finally, we have commented the **R** script to promote accessibility and transparency.

13 ACKNOWLEDGEMENTS

The authors thank Ashleen Benson for translating the calculations in the **Microsoft Access** database to an **R** script, which they referenced when developing `SpawnIndex.R`. They also thank Paul Starr and Rob Kronlund for their thorough reviews that improved this document.

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APPENDIX A SPAWN WIDTH ADJUSTMENTS

Spawn width is a critical component in spawn index calculations. There are two cases where we adjust spawn width estimates to improve spawn index accuracy: surface surveys in all years from 1951 to present, and understory dive surveys between 2003 and 2014.

A.1 SURFACE SURVEY WIDTH

Surface surveys were the only survey type prior to 1988, while the majority of spawns since 1988 have been surveyed using SCUBA gear. Recall that we describe the spawn index as having two periods based on the predominant survey type: the surface survey period from 1951 to 1987, and the dive survey period from 1988 to present.

One issue with comparing these two partly overlapping protocols is that surface surveyors tend to underestimate spawn width (Hay and Kronlund 1987). To improve the consistency of spawn index estimates throughout the time period from 1951 to present, we adjust surface spawn width estimates using underwater estimates when available (Schweigert et al. 1993). Our preferred width is the median width from all dive surveys within a 'pool.' A pool is a group of Locations within a Section that are often adjacent, contain similar algae and substrate, and can be treated as a group with likely similar widths. We summarise spawn width by the median instead of the mean because widths are not normally distributed (Schweigert et al. 1993). If there are no dive data that meet those criteria, we use the median width from all dive surveys within the Section, or within the SAR if there are no dives within the Section. If there are still no dive data that meet those criteria, we use the observed width from the surface survey. We update the aforementioned median width values periodically, not annually.

A.2 UNDERSTORY SURVEY WIDTH

In 2013, Fisheries and Oceans Canada (DFO) staff realized that they were inadvertently underestimating spawn width for Pacific Herring understory dive surveys (Cleary et al. 2017). The issue was caused by the assumed accuracy of transect lines used by spawn surveyors to measure spawn width. Spawn surveyors determine spawn width by placing transects perpendicular to the shore. Surveyors use weighted lead lines to ensure that lines rest on the substrate; these lines are marked in 1 m increments, and are standardized to 20 m segments. Segments refer to individual sections of line, which may be linked together to make complete transects.

Sometime in the mid- to late-1990s, spawn surveyors observed that the

20 m segments shrank by approximately 1 m during the first season of use, and continued to shrink over time. DFO staff noticed that this issue was occurring coastwide, and began re-measuring lead lines each season. They also modified the lead line marking protocol to account for shrinkage by marking 1.15 m increments; thus, segments were extended to 23 m. DFO staff derived this 15% increase by measuring and re-marking lead lines each year. Lead lines are made of a mix of polypropylene and nylon; nylon tightens up under repeated use, which is thought to explain the shrinkage. DFO staff re-measured lead line increments in about 2005, and found that they still shrank from 1.15 m to 1.0 m, and continued to use the modified protocol.

In 2013, spawn surveyors observed that lead line increments were consistently 1.15 m, and no longer appeared to be shrinking. Following this observation, DFO staff re-measured additional lead lines and found that lead lines were made up of a combination of 1.0 m and 1.15 m increments. The combination of observed increment lengths is explained by the lifespan of lead lines: lead lines are replaced every 5 to 10 years, with some segments being replaced more frequently (i.e., inner segments are replaced more frequently than seaward segments, and segments in some SARs are replaced more frequently than in other SARs). DFO staff believe that a change in lead line manufacturing prevents new lead lines from shrinking.

The earliest written instructions that describe the modified protocol of marking 1.15 m increments is from 2003, and this protocol was used until 2013. Note that some SARs continued to use old lead lines in 2014. The practice of annually re-measuring lead line increments ceased around 2005; thus we are unable to determine when lead lines ceased shrinking. Given the observations summarized above, we adjust spawn width estimates based on written instructions for the marking protocol in 2003. Accordingly, our best estimate of years impacted by marking lead lines at 1.15 m increments (when shrinking no longer occurred) is from 2003 to 2014. However, not all SARs r and years y are impacted equally by this issue (Table A.1): some

SARs and years had all 1.0 m increment lengths (no correction factor needed; $WidthFac_{ry} = 1.0$), others had all 1.15 m increment lengths ($WidthFac_{ry} = 1.15$), and others had a combination of 1.0 m and 1.15 m increment lengths which we assume to be in equal proportion ($WidthFac_{ry} = 1.075$). We correct understory spawn widths by multiplying the observed width by the correction factor

$$Width_{txslry} = WidthObs_{txslry} * WidthFac_{ry} \tag{A.1}$$

where $WidthObs_{txslry}$ is the observed spawn width for transect t in spawn survey type x, spawn s, Location l, SAR r, and year y, $WidthFac_{ry}$ is the spawn width correction factor for SAR r and year y (Table A.1), and $Width_{txslry}$ is the corrected understory spawn width in metres. Instead of updating the database permanently, we adjust spawn widths in the **R** script to be transparent, and to prevent a mismatch between the original data sheets and the database.

APPENDIX B SURFACE SPAWN MANUAL UPDATES

One record in the surface spawn database since 1951 requires an update to fill-in missing egg layer information. As with understory spawn width updates, we make this update in the **R** script. This affects the following record:

1. Update ‘intensity’ from 0 to 1 for the 1 record in the year 1962, Statistical Area 14, Section 142, Location code 820, and with intensity = 0. We update intensity from 0 to 1 because spawn was surveyed but not reported.

Spawn survey records prior to 1951 have additional missing or inaccurate egg layer information, and are unreliable for indexing purposes. Therefore, we do not include spawn data prior to 1951 in stock assessments.

While reviewing the spawn index calculations and translating them from the **Microsoft Access** database to **R**, we found several cases where good

Table A.1. Spawn width correction factors $WidthFac_{ry}$ for Pacific Herring understory spawn surveys by stock assessment region (SAR, r) and year y. Legend: Haida Gwaii (HG), Prince Rupert District (PRD), Central Coast (CC), Strait of Georgia (SoG), West Coast of Vancouver Island (WCVI), Area 27 (A27), and Area 2 West (A2W).

	SAR						
Year	HG	PRD	CC	SoG	WCVI	A27	A2W
2003	1.000	1.075	1.075	1.075	1.075	1.075	1.000
2004	1.000	1.075	1.075	1.075	1.075	1.075	1.000
2005	1.000	1.075	1.075	1.075	1.075	1.075	1.000
2006	1.000	1.075	1.075	1.075	1.075	1.075	1.000
2007	1.000	1.075	1.075	1.075	1.075	1.075	1.000
2008	1.000	1.075	1.075	1.075	1.075	1.075	1.000
2009	1.150	1.075	1.075	1.075	1.075	1.075	1.150
2010	1.150	1.075	1.075	1.075	1.075	1.075	1.150
2011	1.150	1.075	1.075	1.075	1.075	1.075	1.150
2012	1.150	1.075	1.075	1.075	1.075	1.075	1.150
2013	1.150	1.150	1.075	1.075	1.075	1.000	1.150
2014	1.150	1.150	1.000	1.000	1.000	1.000	1.150

quality spawn index data were being over-written with no documented reason. These updates have been omitted, and affected the following records:

1. Update $EggLyrs_{js}$ to 2.1496 for the 15 records in the year 1979, Statistical Area 2, and with intensity 4;

2. Update $EggLyrs_{js}$ to 0.5529 for the 4 records in the year 1981, Statistical Area 24, and with $EggLyrs_{js} = 0.0$;

3. Update $EggLyrs_{js}$ to 1.3360 for the 7 records in the year 1982, Statistical Area 23, and with intensity 3;

4. Update $EggLyrs_{js}$ to 2.3300 for 41 records in the year 1984, Statistical Area 24, and with intensity 0; and

5. Update $EggLyrs_{js}$ to 2.9800 for 14 records in the year 1982, Statistical Area 27, and with $EggLyrs_{js} = 0.0$.

In the first three cases, $EggLyrs_{js}$ was updated using intensity categories (Table 2); in the last two cases, $EggLyrs_{js}$ was updated using historical averages. These changes had negligible effects on spawn index values.