# Estimation of an $\mathrm{F}_{\text {MSY }}$ Proxy Reference Point for Spiny Dogfish 

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## INTRODUCTION

The purpose of this report is to refine the estimate of the $\mathrm{F}_{\text {MSY }}$ proxy for spiny dogfish. In 2010 the MAFMC Science and Statistical Committee (SSC) approved a new estimator of the Spawning Stock Biomass (SSB) and a new swept area SSB target of $159,288 \mathrm{mt}$. The target is defined as the SSB corresponding to the estimated maximum recruitment in a Ricker stock recruitment model (Rago and Sosebee 2010). The swept area target corresponds to $30.343 \mathrm{~kg} / \mathrm{tow}$ in the NEFSC spring bottom trawl survey and a trawl survey footprint of $0.0119 \mathrm{~nm}^{2}$.

Previous attempts to develop a fully integrated stock assessment model for spiny dogfish have not been successful (e.g. NEFSC 1994, 1998, 2003, 2006, and TRAC 2010a, b) and it has not been possible to develop an absolute measure of population abundance. However several lines of evidence (see Rago and Sosebee 2009) suggest that the NEFSC trawl captures about $50 \%$ of the spiny dogfish encountered between the trawl doors. The ratio of the distance between the doors and wings of the net is about 2 which, owing to known herding behavior in dogfish, implies an wing-towing efficiency near $100 \%$. The current management of spiny dogfish is based on an assumption that the spawning stock biomass can be adequately indexed by the NEFSC spring bottom trawl survey. Juvenile spiny dogfish, particularly those less than 40 cm are thought to be more pelagic and hence less vulnerable to trawl gear. Ontogenetic changes in behavior result in increasing availability to demersal habitats with size and age. The unknown selectivity pattern for juvenile spiny dogfish complicates the determination of fishing mortality reference points. Since spiny dogfish recruits may be underestimated compared to adult spawners, conventional analyses of a stock recruitment function cannot be conducted. For example many of the observed recruitments fall below the replacement line when $\mathrm{F}=0$. This further emphasizes the problem of establishing the scale of the dogfish recruits (i.e., individuals less than 36 cm TL ).

Target and threshold fishing mortality rates have been based on a life history model presented in Rago et al. (1998). The length-based life history model derives an estimate of the first year survival rate So calibrated to a stanza (1987-1994) of population growth corresponding to a finite rate of increase of 1.044( See Rago et al. 1998, Fig.7). It was noted that the model estimates are highly sensitive to the average size at entry to the fishery (see Rago et al. Fig. 8). Using a knife edge Draft Working Paper for Predissemination Peer Review Only
selection pattern, the replacement $F$ ranged from 0.08 at 65 cm to 0.25 at 84 cm . The life history model has subsequently been improved to incorporate a logistic selectivity pattern (Rago and Sosebee 2009). Analyses of that model for the selectivity pattern estimated in 2006-2008 (Rago and Sosebee 2010) resulted in an F threshold of 0.325 when the median size at entry in the fishery was 90 cm . The target fishing mortality rate, defined as an F which would allow 1.5 female pups per recruit, was 0.207 . Both of these values are dependent on a derived first year survival rate of 0.68 . This corresponds to the first year survival rate in Rago et al. 1998 p. 174 of 0.34 (this value includes both male and female pups at a 50:50 sex ratio)

Following adoption of the revised biomass reference points, the MAFMC SSC examined the population trajectories corresponding to the life history model estimates and noted that the population trajectory associated with $\mathrm{F}=0.325$ ( and $\mathrm{L}_{50}=90 \mathrm{~cm}$ ) and noted that the decline in population size was not consistent with a stable equilibrium. The SSC rejected the estimate of the Fmsy proxy of 0.325 and instead chose an $\mathrm{F}=0.207$ (with expected lifetime pup production of 1.5 female pups per recruit) as an interim value for setting an Acceptable Biological Catch. They further requested an update of the estimate of Fmsy proxy to achieve a value consistent with the projection model. This report summarizes a re-evaluation of the fishing mortality reference points for spiny dogfish with a focus on using a length-based projection model rather than the equilibrium life-history model.

## METHODS

## Life History Model

The life history model estimates pups per recruit as a function of length specific growth, maturation and fecundity. Details on the methodology may be found in Rago et al. (1998, p 167168). Only the summary equation is presented below. A modification made in Rago and Sosebee (2009) was the incorporation of length specific selectivity for fishing mortality.

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$$
\begin{align*}
& P P R=S_{o}\left(R_{1}+\sum_{j=2}^{J} \prod_{i=1}^{j-1} \frac{S_{i} R_{j}}{\lambda^{T_{j}}}\right) \\
& \text { where } \\
& T_{j}=\sum_{i=1}^{j} t_{i} \tag{1}
\end{align*}
$$

Instead of a knife-edge function, we used a smooth function for selectivity as follows:

$$
\begin{equation*}
S_{j}=e^{-\left(\operatorname{Sel}_{L_{j}} F+M\right) \Delta t_{j}} \tag{2}
\end{equation*}
$$

where the selectivity at length $L_{j}$ is described in Eq. 1. A von Bertalanffy growth model is used to estimate length at age and the duration of each age class at length $L_{j}$. A maximum size of 110 cm was used for females. The parameterization of the model is very sensitive to the size of the time steps associated with the duration of each 1 cm increment in length.

## Estimation of Population Size

The variability of initial population size is a primary determinant of the short term dynamics and precision of catch estimates for a given harvest strategy. The magnitude of the population is not important for defining the target fishing mortality rates since this depends only on the population's rate of change in a linear model.

As the details of the population uncertainty are less important for this exercise, the details of this approach are summarized in Appendix 1.

## Projection Model

Since inception of the FMP a length-based stochastic projection model has been used to evaluate effects of alternative fishing mortality scenarios and guide management advice. Short term
management advice on catches relies predominantly on the current non-equilibrium age structure. Transient oscillatory effects of the current age structure are expected to dominate the pattern of landings for the next decade. The magnitude of fishing mortality can offset those effects by damping the oscillations. The model has implications for long-term effects of harvest policies because it can provide insights into the expected population growth rates.

The model incorporates sex specific rates of growth and fishing mortality. Discard mortality is assumed to act equally all size ranges of both sexes. Reproduction in the model is assumed to be proportional to stock abundance. The basic model can be written in terms of two matrix equations as

$$
\begin{align*}
& N_{f, t+1}=S_{f, Z, t} P_{f} N_{f, t}+N_{f, t}^{T} \operatorname{Pup} S_{o} \varphi R_{f}^{o} \\
& N_{m, t+1}=S_{m, Z, t} P_{m} N_{m, t}+N_{f, t}^{T} \operatorname{Pup} S_{o}(1-\varphi) R_{m}^{o} \tag{3}
\end{align*}
$$

where
$\mathbf{N}_{\mathrm{f}, \mathrm{t}}=$ Vector of female population abundance at length. Dimension $=\left(l_{\max }-l_{\min }+1\right)$
$\mathbf{N}_{\mathbf{m}, \mathbf{t}}=$ Vector of male population abundance at length. Dimension $=\left(l_{\max }-l_{\min }+1\right)$
$\mathrm{S}_{\mathrm{f}, \mathrm{Z}, \mathrm{t}}=$ Diagonal matrix of composite survival from instantaneous fishing and natural mortality rates for females at time $t$. Dimensions $=\left(l_{\max }-l_{\min }+1, l_{\max }-l_{\min }+1\right)$
$\mathbf{S}_{\mathbf{m}, \mathbf{Z , t}}=$ Diagonal matrix of composite survival from instantaneous fishing and natural mortality rates for males at time $t$. Dimensions $=\left(l_{\max }-l_{\min }+1, l_{\max ^{-}} l_{\min ^{\prime \prime}}+1\right)$
$\mathbf{R}^{\mathbf{0}}=$ Vector of proportions at length of new recruits. Dimension $=\left(l_{\text {max }}-l_{\text {min }}+1\right)$
$\mathbf{P}_{\mathbf{f}}=$ Growth projection matrix for females. Dimensions $=\left(l_{\max }-1_{\min }+1,1_{\max }-1_{\min }+1\right)$


Pup $=$ Vector of length specific pup production rates for mature females. Dimension $=\left(l_{\text {max }}-\right.$ $1_{\text {min }}+1$ )

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$\mathrm{S}_{\mathrm{o}}=$ Scalar first year survival rate of newborn pups. Derived from analysis of life history model
$\mathrm{T}=$ Transpose operator
$\varphi=$ proportion of female pups at birth; 0.5 implies an equal sex ratio.
The projection equation for males is a function of the numbers of recruits produced by females. The component processes of the matrix model and quantities derived from the population states are described below.

Growth in length at age is modeled by the von Bertalanffy equation applied separately to each sex. The model parameters are taken from Nammack et al. (1985). The projection matrices, $\mathbf{P}_{\mathbf{f}}$ and $\mathbf{P}_{\mathbf{m}}$ for females and males, respectively are defined as square matrices consisting of 0,1 elements. The non-zero elements in cell $\mathrm{i}, \mathrm{j}$ indicate the projection of individuals from cell i to cell j . The growth of individual dogfish from length $i$ to length $j$ is modeled by first inverting the von Bertalanffy equation to obtain the age of individuals of length $i$ to obtain age $\mathrm{i}_{\mathrm{i}}$. The projected length at age $\mathrm{i}_{\mathrm{i}+1}$ by sex then obtained substituting age $\mathrm{i}_{\mathrm{i}}+1$ back into the von Bertalanffy equation to obtain length j as follows:

Step 1. Find age for $L_{i}$

$$
a_{f, i}=\frac{\log \left(1-\frac{L_{i}}{L_{\infty}}\right)}{K}+t_{o}
$$

Step 2.Compute $L$ in next time step

$$
\begin{equation*}
L_{j}=L_{\infty}\left(1-e^{-K\left(a_{i}+1-t_{o}\right)}\right) \tag{4}
\end{equation*}
$$

## Step 3. Compute element of projection matrix

$$
P\left(\operatorname{int}\left(L_{j}\right), \operatorname{int}\left(L_{i}\right)\right)=1
$$

Natural mortality was assumed equal to 0.092 for all length classes. Fishing mortality in year t , defined as $F_{t}$, is multiplied by sex-specific selectivity functions to estimate the sex- and lengthDraft Working Paper for Predissemination Peer Review Only
specific fishing mortality rates. The diagonal matrices that decrement the populations for fishing and natural mortality are defined as $\mathbf{S}_{\mathbf{f}, \mathbf{Z , t}}$ and $\mathbf{S}_{\mathbf{m}, \mathbf{Z , t}}$ with elements defined by

$$
\begin{align*}
& S_{f, Z, t}(\ell, \ell)=e^{-\left(\operatorname{sel}_{f}(\ell) F_{t}+M\right)} \\
& S_{m, Z, t}(\ell, \ell)=e^{-\left(\operatorname{sel}_{m}(\ell) F_{t}+M\right)} \tag{5}
\end{align*}
$$

The total number of pups produced is written at the product of the length-specific pup production rates and the number of females alive in year t .

$$
\begin{equation*}
\operatorname{Pup}_{\text {тот }, t}=S_{o} N_{f, t}^{T} \text { Pup } \tag{6}
\end{equation*}
$$

The numbers of recruits produced by length and size category is estimated by splitting the total pup number by sex and multiplying by the observed proportion of dogfish at length for lengths $<35 \mathrm{~cm}$ (assumed to be less than one year old at the time of the survey). The resulting numbers of pups by sex produced is written as:

$$
\begin{align*}
& \text { female pups }=R_{f, t}=\varphi \text { Pup }_{\text {тот }, t} R_{f}^{o}  \tag{7}\\
& \text { male pups }=R_{m, t}=(1-\varphi) \text { Pup }{ }_{\text {тот }, t} R_{m}^{o}
\end{align*}
$$

The $\mathbf{R}^{\mathbf{0}}$ and $\mathbf{R}_{\mathbf{m}}^{\mathbf{0}}$ vectors of length ( $1_{\max }-1_{\min }+1$ ) represent the proportions by length class and was estimated from the proportions in the 2006-2008 spring surveys. The male and female vectors have equivalent proportions but differ with respect to vector length, owing to the larger maximum size attained by females. Spawning stock biomass is expressed in terms of female biomass only and is defined as the sum of mature females. Females are assumed to be mature at 80 cm so that the spawning stock biomass can be written as

$$
\begin{equation*}
S S B_{t}=\sum_{j=80}^{\ell_{\max }} N_{f, t}(j) W_{f}(j) \tag{8}
\end{equation*}
$$

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The initial condition of the population was defined as the 3-yr average (2008-2010) of dogfish abundance in the NEFSC spring R/V trawl survey. The projection model incorporates the variation in abundance defined by survey abundance. Variation in mean abundance is used to scale the index numbers at length by generating values of mean abundance over 500 equally-spaced probability intervals.

When length specific fecundity is held constant and none of the other model parameters vary as a function of population size, Equation 3 is a linear system of equations and has properties similar to a stage-based Leslie matrix model. Briefly, the linear system of the form $\mathrm{AX}_{\mathrm{t}}=\mathrm{X}_{\mathrm{t}+1}$ with a positive definite matrix A will be expected to achieve a constant rate of increase $\lambda$ such that $\mathrm{AX}_{\mathrm{t}}=\lambda \mathrm{X}_{\mathrm{t}}$ as t approaches infinity. Generally we don't have to wait that long and the model equilibrates after a modest number of iterations. Alternatively one can estimate the dominant eigenvalue and eigenvector of A to derive the steady state growth rate and population size structure.

Model complexity can be increased by introducing a stock recruitment model

$$
\begin{equation*}
\text { Pup }_{\text {тот }, t}=G\left(S S B_{t} \mid \Theta\right) \tag{9}
\end{equation*}
$$

Where SSB is defined by Eq. 8 and $\Theta$ represents a vector of parameters. For example the stock recruitment model defined in Rago and Sosebee (2010) could be used. Details on this exploratory exercise are described in Appendix 2.

## RESULTS

The predicted female pups per female recruit from the equilibrium life history model are depicted in Fig. 1. The model incorporates the selectivity pattern for 2006-2008 fishery (Appendix 1). Fthreshold ( 0.325 ) corresponds to a lifetime production of 1 female pup per female recruit. Ftarget ( 0.207 ) corresponds to a lifetime production of 1.5 female pups per female recruit.

The inconsistency between the predicted population trajectories associated with $\mathrm{F}=0.325$ (and $\mathrm{L}_{50}$ $=90 \mathrm{~cm}$ ) is demonstrated in Fig. 2. When $\mathrm{F}=0.207$ the population increases at about $1 \%$ per year Draft Working Paper for Predissemination Peer Review Only
( $\lambda=1.00892$ ) but when $\mathrm{F}=0.325$ the population declines at about $1.9 \%$ per year $(\lambda=1.00892)$. The differences between the life-history model predictions and the projection model are small in absolute terms but result in large differences over the time span of 50 years or more.

The Newton-Raphson algorithm was used to estimate a fishing mortality rate that would achieve a $(\lambda=1.00)$ given the initial conditions of the population (Table 1 ). When $\mathrm{F}=0.2439$ the population stabilizes at a target biomass of about $125,000 \mathrm{mt}$. However, since the model is linear, the population can be stabilized at any population size by applying an alternative F until the desired population SSB is achieved. A two stage harvest policy is applied in Fig. 4 wherein a fishing mortality rate of 0.235 is applied from 2010 to 2129 and followed by the threshold $\mathrm{F}=0.2439$ from 2130 to 2259 . The harvest policy allows the population to grow between 2010 and 2129. The population fluctuates slightly when the fishing mortality increases and then returns to equilibrium at the target female SSB of $159,288 \mathrm{mt}$. A detailed summary of the statistical properties of the stable population is provided in Table 2.

Using a harvest control rule of $75 \%$ of the Fmsy proxy gives an F of 0.18293 . Expected annual population growth is $1.52 \%$ per year or a finite rate of increase of 1.01527 (Fig. 5). The maximum population growth rate, attainable occurs when $\mathrm{F}=0.0$, is estimated to be about $7 \%$ per year $(\lambda=1.07)$ (Fig. 6). The rate of population growth decreases linearly with F (Fig. 7).

As shown by numerous authors in the ecological and fisheries literature, Leslie matrix type models are very sensitive to the first year survival rate $\left(\mathrm{S}_{\mathrm{o}}\right)$ The effect of variation in So on the equilibrium $F$ is shown in the text tables below. If So increases from 0.68 to 0.892 the equilibrium Fmsy proxy would increase to 0.3269 . Similarly, a decline in So to 0.468 would reduce the Fmsy proxy to 0.15289 .

| So | F | r | lambda |
| ---: | ---: | ---: | ---: |
| 0.892 | 0.2439 | 0.018433 | 1.018603936 |
| 0.892 | 0.3649 | -0.00842455 | 0.991610837 |
| 0.892 | 0.326945287 | 0 | 1 |


| So | F | r | lambda |
| ---: | ---: | ---: | ---: |
| 0.4688 | 0.2439 | -0.024720102 | 0.975582937 |
| 0.468 | 0.12195 | 0.008405458 | 1.008440884 |
| 0.468 | 0.152894251 | 0 | 1 |

## DISCUSSION

The revised estimate of the Fmsy proxy for spiny dogfish is 0.2439 . Using a $75 \%$ of Fmsy proxy as a candidate target F results in population growth rate of $1.5 \%$ per year. The effects of these alternative fishing mortality reference points on predicted USA landings, total catch and SSB in 2011 are summarized in Table 3 and 4 for Fmsy proxy and 75\% Fmsy proxy, respectively. These summaries are done for illustrative purposes only and are meant to show the properties of the new estimators on predicted catch using the same initial conditions as employed during the 2010 specifications setting process. Choice of a target F reference point is typically based on a desired population growth rate or buffer associated with uncertainty in the Overfishing Limit (OFL).

There are multiple factors influencing the uncertainty in the fishing mortality rate estimator for spiny dogfish in the projection model. While it is possible to develop a parametric Monte Carlo sampling approach to estimate the uncertainty in Fmsy, it would be desirable to discuss the potential range of input parameters. Variations in growth rate, selectivity, maturity and first year survival would likely be key parameters to vary. The derived sampling distribution of Fmsy proxies could then be convolved with the uncertainty in population abundance to obtain a better approximation of the OFL uncertainty.

Ultimately it would be desirable to develop an integrated stock assessment model for spiny dogfish. The TRAC made considerable progress toward that objective. The projection model described herein could be used as a basis for developing a length based assessment model. Alternatively, something like Sullivan's CASA model could be applied. One of the key obstacles for such a model is the estimation of selectivity for the more pelagic pups. It is hypothesized that juvenile dogfish gradually become more demersal with age, but the age or size at full recruitment to the bottom habitat has not been estimate. The difficulties of rescaling the recruitment estimates to be

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consistent with the predictions of the Ricker stock recruitment model (Rago and Sosebee 2010) are examined in Appendix 2.

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Fig. 1. Current estimates of fishing mortality reference points for spiny dogfish based on equilibrium life history model. Model incorporates selectivity pattern for 2006-2008 fishery. Fthreshold ( 0.325 ) corresponds to a lifetime production of 1 female pup per female recruit. Ftarget ( 0.207 ) corresponds to a lifetime production of 1.5 female pups per female recruit.

## Stochastic Projections at $\mathrm{F}=0.207$ and $\mathrm{F}=0.325$



Fig 2. Stochastic projections of SSB at current fishing mortality threshold ( $\mathrm{F}=0.325$ ) and target ( $\mathrm{F}=0.207$ ) fishing mortality rates defined in Rago and Sosebee (2010). Threshold and target Fs are based on life history model. Horizontal dashed lines represent biomass target and threshold values of $159,288 \mathrm{mt}$ and 79644 mt , respectively. Projections depict $2.5 \%$, $50 \%$ and $97.5 \%$ iles for each scenario. The expected finite rate of population increase at $\mathrm{F}=0.325$ is 0.98128 or about $1.9 \%$ decline per year. The finite rate of population increase at $\mathrm{F}=0.207$ is 1.00892 or about a $0.9 \%$ increase per year.

## Longterm Projection at Fmsyproxy given 2010 Initial Condition



Fig. 3. Median projection of female spawning stock biomass (mt) and projected US landings (mt) using estimated fishing mortality rate of 0.2439 that achieves a finite rate of increase equal to 1.0. Horizontal dashed lines represent biomass target and threshold values of $159,288 \mathrm{mt}$ and 79644 mt , respectively. Transient population dynamics prior to 2050 are induced by the non-equilibrium initial condition of population size structure.

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## Transitional $\mathrm{F}=0.235$ to $\mathrm{F}=0.244$ Attain SSBmsy proxy



Fig. 4. Median projection of female spawning stock biomass ( mt ) and projected US landings ( mt ) using a two stage fishing mortality harvest policy designed to achieve a target biomass of $159,288 \mathrm{mt}$. The harvest policy allows the population to grow between 2010 and 2129. A fishing mortality rate of 0.235 from 2010 to 2129 and followed by the threshold $\mathrm{F}=0.2439$ from 2130 to 2259 . The population fluctuates slightly when the fishing mortality increases and then returns to equilibrium at the target female SSB. Horizontal dashed lines represent biomass target and threshold values of 159,288 mt and 79644 mt , respectively.
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## Longterm Projection at 75\% Fmsyproxy given 2010 Initial Condition



Fig. 5. Median longterm projection of female spawning stock biomass ( mt ) and projected US landings (mt) using a harvet policy of $75 \%$ of the Fmsy proxy ( $=0.2439$ ). Trajectories represent SSB and landings under a fishing mortality rate of 0.18293 . Expected annual population growth is $1.52 \%$ per year or a finite rate of increase of 1.01527 Horizontal dashed lines represent biomass target and threshold values of $159,288 \mathrm{mt}$ and 79644 mt , respectively.

## Longterm Projection at F=0.0 given 2010 Initial Condition



Fig. 6. Median longterm projection of female spawning stock biomass (mt) and projected US landings ( mt ) using a harvet policy of $\mathrm{F}=0.0$ Expected annual population growth is $7 \%$ per year or a finite rate of increase of 1.07. Horizontal dashed lines represent biomass target and threshold values of $159,288 \mathrm{mt}$ and 79644 mt , respectively.


Fig. 7. Predicted instantaneous population growth rates and finite rate of increase for spiny dogfish as a function of fishing mortality rate. The first year survival rate is fixed at 0.68 .

Table 1. Estimates of population growth rate for varying levels of F and So

| Description | So | $\boldsymbol{F}$ | $\boldsymbol{\operatorname { l n } ( \boldsymbol { \lambda } )}$ | $\boldsymbol{\lambda}$ |
| :--- | :--- | :--- | :--- | :--- |
| Fmsy proxy for Linear Model | 0.68 | 0.2439 | 0 | 1 |
| $\mathrm{~F}=0.0$ To estimate maximum growth rate | 0.68 | 0 | 0.067364 | 1.07 |
| $75 \%$ of Fmsy Proxy | 0.68 | 0.182925 | 0.015151 | 1.01527 |
| 50\% of Fmsy Proxy | 0.68 | 0.12195 | 0.031274 | 1.03177 |
| Old Fmsy Proxy based on Life History model | 0.68 | 0.325 | -0.01891 | 0.98127 |
| Old Target F based on Life History Model | 0.68 | 0.207 | 0.008989 | 1.00892 |

Table 2. Summary statistics for population biomass, total catch and estimated USA landings in 2150 for a population stabilized at Bmsy and $\mathrm{F}=$ Fmsyproxy. A fishing mortality rate of 0.235 from 2010 to 2129 and followed by the threshold $\mathrm{F}=0.2439$ from 2130 to 2259.

|  | USA Landings(mt) | Total Catch(mt) | SSB (mt) |
| :---: | :---: | :---: | :---: |
| Minimum | 6,597 | 14,972 | 98,451 |
| Maximum | 15,908 | 33,445 | 219,921 |
| Median | 11,220 | 24,146 | 158,770 |
| Mean | 11,224 | 24,152 | 158,815 |
| Standard Dev | 2,088 | 4,143 | 27,243 |
| Percentile |  |  |  |
| 1\% | 6,890 | 15,552 | 102,266 |
| 5\% | 7,741 | 17,242 | 113,375 |
| 10\% | 8,412 | 18,574 | 122,131 |
| 20\% | 9,332 | 20,399 | 134,135 |
| 25\% | 9,699 | 21,127 | 138,921 |
| 30\% | 10,033 | 21,790 | 143,278 |
| 40\% | 10,646 | 23,005 | 151,270 |
| 50\% | 11,220 | 24,146 | 158,770 |
| 60\% | 11,796 | 25,287 | 166,276 |
| 70\% | 12,408 | 26,503 | 174,269 |
| 75\% | 12,744 | 27,168 | 178,642 |
| 80\% | 13,111 | 27,897 | 183,437 |
| 90\% | 14,041 | 29,741 | 195,566 |
| 95\% | 14,721 | 31,092 | 204,445 |
| 99\% | 15,599 | 32,832 | 215,890 |

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Table. 3. Summary of predicted catch statistics and SSB in 2011 given F=Fmsyproxy

|  | USA Landings | Total Catch | SSB |
| :---: | :---: | :---: | :---: |
| Minimum | 6,513 | 16,274 | 113,608 |
| Maximum | 14,638 | 31,266 | 259,245 |
| Median | 10,547 | 23,718 | 185,921 |
| Mean | 10,550 | 23,724 | 185,976 |
| Standard Dev | 1,823 | 3,363 | 32,664 |
| Percentile |  |  |  |
| 1\% | 6,768 | 16,745 | 118,179 |
| 5\% | 7,510 | 18,115 | 131,495 |
| 10\% | 8,096 | 19,196 | 141,991 |
| 20\% | 8,899 | 20,677 | 156,382 |
| 25\% | 9,219 | 21,268 | 162,119 |
| 30\% | 9,510 | 21,805 | 167,342 |
| 40\% | 10,046 | 22,793 | 176,933 |
| 50\% | 10,547 | 23,718 | 185,921 |
| 60\% | 11,049 | 24,644 | 194,918 |
| 70\% | 11,584 | 25,631 | 204,505 |
| 75\% | 11,877 | 26,171 | 209,748 |
| 80\% | 12,197 | 26,762 | 215,492 |
| 90\% | 13,009 | 28,261 | 230,050 |
| 95\% | 13,603 | 29,356 | 240,691 |
| 99\% | 14,368 | 30,768 | 254,408 |

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Table 4. Summary of predicted catch statistics and SSB in 2011 given $\mathrm{F}=75 \%$ Fmsyproxy

|  | USA Landings | Total Catch | SSB |
| :---: | :---: | :---: | :---: |
| Minimum | 4,711 | 12,327 | 113,608 |
| Maximum | 10,886 | 23,720 | 259,245 |
| Median | 7,777 | 17,984 | 185,921 |
| Mean | 7,779 | 17,988 | 185,976 |
| Standard Dev | 1,385 | 2,555 | 32,664 |
| Percentile |  |  |  |
| 1\% | 4,905 | 12,685 | 118,179 |
| 5\% | 5,469 | 13,726 | 131,495 |
| 10\% | 5,914 | 14,547 | 141,991 |
| 20\% | 6,524 | 15,673 | 156,382 |
| 25\% | 6,768 | 16,122 | 162,119 |
| 30\% | 6,989 | 16,530 | 167,342 |
| 40\% | 7,396 | 17,281 | 176,933 |
| 50\% | 7,777 | 17,984 | 185,921 |
| 60\% | 8,158 | 18,688 | 194,918 |
| 70\% | 8,565 | 19,437 | 204,505 |
| 75\% | 8,787 | 19,848 | 209,748 |
| 80\% | 9,031 | 20,297 | 215,492 |
| 90\% | 9,648 | 21,436 | 230,050 |
| 95\% | 10,099 | 22,268 | 240,691 |
| 99\% | 10,681 | 23,341 | 254,408 |

Draft Working Paper for Predissemination Peer Review Only

## APPENDIX 1. Sampling Distribution of Biomass and Fishing Mortality

The sampling distributions of biomass and fishing mortality are approximated by integrating over the factors which constitute the primary sources of uncertainty. These factors include the sampling variability in the NEFSC spring bottom trawl survey, the size of the area swept by an average tow, and the uncertainty in the discard estimates. The sample means and variances for each of these factors were used to parameterize their respective normal distributions. Sampling theory suggests that the survey means should be asymptotically normal. We exploit this feature to simplify the estimation of the sampling distribution of biomass and fishing mortality.

The sampling distribution of each of the Fs described above was evaluated by integrating over each of the normal distributions for average number $\bar{X}$, survey footprint $\boldsymbol{a}$, and total discards for gill nets and trawls, and recreational catch, $\boldsymbol{D}_{G}, \boldsymbol{D}_{\boldsymbol{T}}$, and $\boldsymbol{D}_{\boldsymbol{R}}$ respectively. The density $\bar{X}$ and footprint a parameters were evaluated over 500 equal probability intervals, while the sampling distribution $D_{G}, D_{T}$, and $D_{R}$ were evaluated over 20 equal probability intervals.

Let $\Phi=$ Normal cumulative distribution function. The inverse of $\Phi$, denoted as $\Phi^{-1}$ allows the evaluation of a set of values over a specified range, say $\alpha_{\min }$ and $\alpha_{\max }$, over equal probability intervals. The value of the random variable X associated with the $\alpha$ level is defined as:

$$
\begin{equation*}
X_{t, \alpha}^{\prime}=\Phi^{-1}\left(\alpha \mid \overline{\bar{X}}, \bar{S}_{t}^{2}\right) \tag{1.1}
\end{equation*}
$$

The step size between successive values of $\alpha$ was set as $\delta_{1}=1 / 500(0.975-0.025)$, where $\alpha_{\text {min }}$ $=0.025$ and $\alpha_{\max }=0.975$. An equivalent approach was used for evaluation of the footprint parameter a where $\mathrm{a} \sim \mathrm{N}\left(\mu_{\mathrm{a}}, \sigma_{\mathrm{a}}{ }^{2}\right)$. The total discard estimate $\mathrm{D} \sim \mathrm{N}\left(\mu_{\mathrm{D}}, \sigma_{\mathrm{D}}{ }^{2}\right)$ was approximated by dividing the $95 \%$ confidence interval into 20 equal probability values ( $\delta_{2}=1 / 20$ ). Discard means and variances were estimated for each gear and sex.

This property can be illustrated for the biomass estimates by rewriting Eq. 2 and 3 as

$$
\begin{equation*}
B^{\prime}(l)=N^{\prime}(l) W(l)=I^{\prime}(l)\left(\frac{A}{a^{\prime}}\right) W(l)=\bar{X}_{\alpha}^{\prime} p_{s}(l)\left(\frac{A}{a_{\alpha}^{\prime}}\right) W(l) \tag{1.2}
\end{equation*}
$$

where the prime mark indicates that value is drawn from a normal distribution (Eq. 1.1)
corresponding the $\alpha$ probability level. The expected value of $\boldsymbol{B} \boldsymbol{\prime}(\boldsymbol{l})$ is obtained by summing over the sampling distributions of $\boldsymbol{X}$ and $\boldsymbol{a}$ as follows

$$
\begin{equation*}
E\left[B^{\prime}(l)\right]=\sum_{\alpha=\alpha_{\text {min }}}^{\alpha_{\text {max }}} \sum_{\beta=\beta_{\text {min }}}^{\beta_{\text {max }}}\left(\bar{X}_{\alpha}^{\prime} p_{s}(l)\left(\frac{A}{a_{\beta}^{\prime}}\right) W(l) \delta_{\alpha} \delta_{\beta}\right) \tag{1.3}
\end{equation*}
$$

The sampling distribution of $\boldsymbol{B}^{\prime}(\mathbf{l})$ can be constructed by noting that the each element within the brackets of the rhs of Eq. 1.3 has a probability weight of $\delta_{\alpha} \delta_{\beta}=(1 / 500)(1 / 500)$.

The sampling distribution of F in Eq. 5 is a straightforward extansion of Eq. 1.3 with summation terms corresponding to average number $\bar{X}$, survey footprint $\boldsymbol{a}$, and total discards for gill nets and trawls, and recreational catch, $\boldsymbol{D}_{\boldsymbol{G}}, \boldsymbol{D}_{\boldsymbol{T}}$, and $\boldsymbol{D}_{\boldsymbol{R}}$ respectively. This approximation of the multidimensional integration provides reasonable assurance that the sampling distribution of the F will be appropriately estimated.

## Appendix 1 Extra: Selectivity function for 2006-2008

Comparison of size composition of commercial catch (landings + Discards) for male and female spiny dogfish with the NEFSC spring survey for 2006-2008. Both catch and survey frequencies represent $3-\mathrm{yr}$ moving averages. Summary of estimated selectivity pattern for male and female spiny dogfish. Selectivity at length $L$ is modeled as $\operatorname{sel}(L)=1 /(1+\exp (a+b$ $\mathrm{L})$ ) where $\operatorname{sel}(\mathrm{F})$ is the fraction of the spiny dogfish population vulnerable to the commercial fishery (both landings and discards). Size composition of the commercial fishery is based on analyses of port sampling and at-sea observer sampling, 1989-2008. Selectivity blocks are based on a 3-yr moving average, eg. 2006-2008.


Appendix 2. Summary of model projections using the Ricker model estimates in Rago and Sosebee (2010).

As an exploratory exercise, the Ricker stock recruitment model was used to predict recruitment. The relative catchability of the pups compared to adult spiny dogfish must ultimately be estimated via a dynamic model. For the purposes of this exercise, the expected female SSB and yield were estimated for a variety of scaling factors on recruits. To initially rescale the population it was assumed that the Fmsy proxy for the linear model would be approximate Fmsy estimate from the Ricker model. The predicted Ricker model recruitment was adjusted by a factor of 0.74829 (rather than the expected 0.5 for a $50: 50$ sex ratio) to obtain a predicted equilibrium $\operatorname{SSB}$ of 157,791 (row 1 below). Thus an adjustment factor of 0.74829 resulted in an Bmsy estimate of 157,791 or approximately $159,288 \mathrm{mt}$ the proxy value of Bmsy (See Rago and Sosebee 2010). The following table summarizes the effects of further searches for maximum total yield. Note the high sensitivity of the total yield estimate and SSB to changes in F. In the vicinity of this rescaling the maximum yield of 24042 mt is obtained at a fishing mortality rate of 0.257925 and a predicted SSBmsy of $149,538 \mathrm{mt}$. This analysis is by means definitive but it does suggest that modest consistency between the Bmsy proxy from the linear model and a Bmsy estimate from the Ricker model.

| Adj Factor | F | SSB | Ytot |
| ---: | ---: | ---: | ---: |
| 0.74829 | 0.2439 | 157791 | 23994 |
| 0.74829 | 0.2 | 126194 | 23639 |
| 0.74829 | 0.3 | 185479 | 23225 |
| 0.74829 | 0.27195 | 141533 | 23995 |
| 0.74829 | 0.257925 | 149538 | 24042 |

## Longterm Projection at $\mathrm{F}=\mathrm{Fmsyproxy}$ with $\mathrm{S}-\mathrm{R}$ function at 0.5 X



Fig. A2.1 Predicted median trajectory for population subjected to $\mathrm{F}=0.2439$ and recruitment predicted by Ricker stock recruitment function. Predicted recruits are estimated to have a 50:50 sex ratio.

## Longterm Projection at F=Fmsyproxy with S-R function at 1.0X



Fig. A2.2 Predicted median trajectory for population subjected to $\mathrm{F}=0.2439$ and recruitment predicted by Ricker stock recruitment function. Predicted recruits are adjusted upward by a factor of 2. This rescales the predicted recruitment consistent with a relative catchability of juveniles to adults of $1 / 2$.

## Longterm Projection at F=Fmsyproxy with S-R function at 0.74829X



Fig. A2.3. Predicted median trajectory for population subjected to an estimated Fmsy=0.258 and recruitment predicted by Ricker stock recruitment function. Predicted recruits are adjusted upward by a factor of 1.496 . Predicted USA landings would be $24,042 \mathrm{mt}$. This rescales the predicted recruitment consistent with a relative catchability of juveniles to adults of 0.748.

