# Standardized catch rates of blacktip sharks (Carcharhinus limbatus) 

 collected during bottom longline surveys in Mississippi, Louisiana, and Alabama coastal waters from 2004 to 2010Eric Hoffmayer, Jill Hendon, Marcus Drymon, Sean Powers, Adam Pollack, and John Carlson

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# STANDARDIZED CATCH RATES OF BLACKTIP SHARKS (CARCHARHINUS LIMBATUS) COLLECTED DURING BOTTOM LONGLINE SURVEYS IN MISSISSIPPI, LOUISIANA, AND ALABAMA COASTAL WATERS FROM 2004 TO 2010. 

Eric Hoffmayer ${ }^{1}$, Jill Hendon ${ }^{2}$, Marcus Drymon ${ }^{3}$, Sean Powers ${ }^{3}$, Adam Pollack ${ }^{1}$, and John Carlson ${ }^{4}$

Originally three separate indices were created to detail bottom longline survey blacktip shark catches in the Alabama, Mississippi, and Louisiana coastal waters. Detailed information about the three surveys is found within the following documents: SEDAR29-WP-11 for the Alabama index, SEDAR29-WP14 for the inshore Mississippi index, and SEDAR29-WP-15 for the Louisiana/Mississippi index. The SEDAR 29 panel decided that this catch information would be most valuable if an index was created using the data from all three surveys combined. The combined index extended from 2004 to 2010, and resulted in 893 sets and 1,379 blacktip sharks. Standardized catch rates were estimated using a generalized linear mixed modeling approach assuming a delta-lognormal error distribution and negative binomial regression.

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## INTRODUCTION

During the SEDAR 29 data workshop, the panel decided that it would be most appropriate to combine the indices from three regional bottom longline surveys in the north central Gulf of Mexico to develop a more temporally and spatially robust index. Details for the three surveys are found within the following documents: SEDAR29-WP-11 for the Alabama survey, SEDAR29-WP14 for the inshore Mississippi survey, and SEDAR29-WP-15 for the Louisiana/Mississippi survey. As a result, the following index was generated using all catch data for age 1+ blacktip sharks from the three indices.

## METHODOLOGY

## Alabama Survey

The sampling protocol and equipment follows the procedures established by the NOAA Fisheries Mississippi Laboratories bottom longline survey (Grace and Henwood 1997). The longline gear consisted of a 1.6 km ( 426 kg test) monofilament mainline and 100, 3.7 m gangions ( 332 kg test monofilament) outfitted with \#15/0 circle hooks and baited with Atlantic mackerel (Scomber scombrus). The longline fished for one hour from the time of last high-flier deployment to the time of first high-flier retrieval. Bottom longline sampling for the Alabama nearshore survey began in May 2006 and employed a random stratified block design. Blocks were established both in the Mississippi Sound/Mobile Bay and waters south of Dauphin Island. Each month (January to December), stations were randomly selected within the blocks, and effort was allocated across three depth strata ( $0-5 \mathrm{~m}, 5-10 \mathrm{~m}$, and $10-20 \mathrm{~m}$ ). For additional details see SEDAR29-WP-11.

## Mississippi Inshore Survey

Sampling was conducted with a 152.4 m bottom longline that consisted of 50 hooks (\#12/0 circle), 1.0 m gangions ( 2.0 mm monofilament), with menhaden (Brevoortia patronus) as bait. The longline was typically fished between the hours of 0800 and 2000, and was allowed to soak for one hour prior to retrieval. The bottom longline sampling employed a random stratified block design, with twelve $10.6 \mathrm{~km}^{2}$ blocks select throughout the Mississippi Sound region. Each month from March to October, stations were randomly selected within each block. For additional details see SEDAR29-WP-14.

## Mississippi/Louisiana Survey

The sampling protocol and equipment follows the procedures established by the NOAA Fisheries Mississippi Laboratories bottom longline survey (Grace and Henwood 1997). The longline gear consisted of a 1.6 km ( 426 kg test) monofilament mainline and 100, 3.7 m gangions ( 332 kg test monofilament) outfitted with \#15/0 circle hooks and baited with Atlantic mackerel, (Scomber scombrus). The longline fished for one hour from the time of last high-flier deployment to the time of first high-flier retrieval. The bottom longline sampling employed a random stratified block design with effort within each block allocated across three depth strata ( $0-5 \mathrm{~m}, 5-10 \mathrm{~m}$, and $10-20 \mathrm{~m}$ ). The study area was broken into three regions: Mississippi Sound, South of barrier

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islands, and Chandeleur Sound. Each month from March to October, three stations were sampled from each region. For additional details see SEDAR29-WP-15.

## Combined Survey Modifications

The study area for the Alabama, Mississippi inshore, and Mississippi/Louisiana surveys was approximately $1,450,190$, and $1,050 \mathrm{~km}^{2}$, respectively. Due to the spatial overlap in the three surveys, the entire study area was divided into eleven $26 \times 6 \mathrm{~km}$ blocks (blocks 1-6, 8-12), and one 17 x 18 km block (Chandeleur Sound; block 7) (Figure 1). Each station sampled by the individual surveys was defined as being within one of these 12 blocks. Soak time was calculated differently between the three surveys. However, as all three surveys allowed the gear to fish for one hour prior to retrieval, one hour was chosen to use as the soak time in the combined index.

The three surveys also utilized different model input factors. To aggregate all three surveys for analysis, the Mississippi inshore and Mississippi/Louisiana datasets removed the 2011 catch data, as well as the monthly rainfall, previous month rainfall, temperature, salinity, and dissolved oxygen factors. The factors that remained in the combined dataset included survey, year, month, area, depth, set time, soak time, and hook size. Finally, because there was variability in the sample size among the surveys, the three indices were weighted by the spatial area covered by the survey.

## Index Construction

Delta-lognormal modeling methods were used to estimate relative abundance indices for blacktip sharks (Lo et al. 1992). The main advantage of using this method is the allowance for the probability of zero catch (Ortiz et al. 2000). The index computed by this method is a mathematical combination of yearly abundance estimates from two distinct generalized linear models: a binomial (logistic) model which describes the proportion of positive abundance values (i.e. presence/absence), and a lognormal model which describes variability in only the non-zero abundance data (Lo et al. 1992).

The delta-lognormal index of relative abundance $\left(I_{y}\right)$ as described by Lo et al. (1992) was estimated as:
(1) $\quad I_{y}=c_{y} p_{y}$,
where $c_{y}$ is the estimate of mean CPUE for positive catches only for year $y$, and $p_{y}$ is the estimate of mean probability of occurrence during year $y$. Both $c_{y}$ and $p_{y}$ were estimated using generalized linear models. Data used to estimate abundance for positive catches (c) and probability of occurrence ( $p$ ) were assumed to have a lognormal distribution and a binomial distribution, respectively, and modeled using the following equations:
(2) $\ln (c)=X \beta+\varepsilon$
and

$$
\begin{equation*}
p=\frac{e^{\mathrm{X}_{\beta}+\varepsilon}}{1+e^{\mathrm{X} \beta+\varepsilon}}, \tag{3}
\end{equation*}
$$

respectively, where $c$ is a vector of the positive catch data, $p$ is a vector of the presence/absence data, $X$ is the design matrix for main effects, $\beta$ is the parameter vector for main effects, and $\varepsilon$ is a vector of independent normally distributed errors with expectation zero and variance $\sigma^{2}$. Therefore, $c_{y}$ and $p_{y}$ were estimated as least-squares means for each year along with their corresponding standard errors, $\mathrm{SE}\left(c_{y}\right)$ and $\mathrm{SE}\left(p_{y}\right)$, respectively. From these estimates, $I_{y}$ was calculated, as in equation (1), and its variance calculated as:

$$
\begin{equation*}
V\left(I_{y}\right) \approx V\left(c_{y}\right) p_{y}^{2}+c_{y}^{2} V\left(p_{y}\right)+2 c_{y} p_{y} \operatorname{Cov}(c, p), \tag{4}
\end{equation*}
$$

where:
(5) $\left.\quad \operatorname{Cov}(c, p) \approx \rho_{c, \mathrm{p}} \operatorname{SE}\left(c_{y}\right) \operatorname{SE}\left(p_{y}\right)\right]$,
and $\rho_{\mathrm{c}, \mathrm{p}}$ denotes correlation of $c$ and $p$ among years.
The submodels of the delta-lognormal model were built using a backward selection procedure based on type 3 analyses with an inclusion level of significance of $\alpha=0.10$. Binomial submodel performance was evaluated using AIC, while the performance of the lognormal submodel was evaluated based on analyses of residual scatter and QQ plots in addition to AIC. Due to differences in area surveyed among the data sets, the time series was weighed by sample size or area surveyed.

## RESULTS

From 2004 to 2010, 893 sites were sampled resulting in the catch of 1,379 blacktip sharks. The number of sites sampled varied across surveys with Alabama (406) having the highest, followed by Mississippi inshore (276) and Mississippi/Louisiana (211). The total number of blacktip sharks captured each year ranged from 61 to 340 sharks. Approximately $37 \%$ of the stations sampled contained positive catches of blacktip sharks, with Mississippi/Louisiana (46.4\%) having the highest, followed by Alabama (38.4\%), and Mississippi inshore (28.0\%).

The model outputs for the series are in Table 1. Table 2 contains the standardized abundance series for both weighing schemes and Figure 2 illustrates the time series.

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Table 1. Analysis of deviance of explanatory variables for the binomial and lognormal generalized linear and mixed model formulations of the proportion of positive and positive catches for blacktip sharks.

| Proportion positive-Binomial error distribution |  |  |  |  |  |
| :--- | :---: | :---: | :---: | :---: | :---: |
| FACTOR | DEVIANCE/DF | \%DIFF | DELTA\% | CHISQUARE | PR>CHI |
| NULL | 1.7452 |  |  |  |  |
| YEAR | 1.7251 | 1.152 | 1.15 | 17.28 | 0.0083 |
|  |  |  |  |  |  |
| YEAR+ |  |  |  | 72.71 | $<.0001$ |
| AREA | 1.5656 | 10.291 | 9.14 | 43.49 | $<.0001$ |
| SEASON | 1.6107 | 7.707 |  | 28.87 | $<.0001$ |
| SURVEY | 1.6494 | 5.489 |  | 26.04 | $<.0001$ |
| DEPTH | 1.6579 | 5.002 |  | 20.26 | $<.0001$ |
| HOOK | 1.6701 | 4.303 |  |  |  |
|  |  |  |  | 52.11 | $<.0001$ |
| YEAR+AREA+ | 1.4188 | 18.703 | 8.41 | 13.66 | 0.0011 |
| SEASON | 1.5331 | 12.153 |  | 5.01 | 0.0253 |
| DEPTH | 1.555 | 10.898 |  | 5.42 | 0.0664 |
| HOOK | 1.5585 | 10.698 |  |  |  |
| SURVEY |  |  |  |  |  |
|  |  |  |  | 13 | 0.0015 |
| YEAR+AREA+SEASON | 1.3871 | 20.519 | 1.82 |  | 0.07 |
| DEPTH | 1.4018 | 19.677 |  | 6.92 | 0.032 |
| HOOK | 1.406 | 19.436 |  |  |  |
| SURVEY |  |  |  |  |  |


| Proportion positive-Lognormal error distribution |  |  |  |  |  |
| :--- | :--- | :--- | :--- | :--- | :---: |
| FACTOR | DEVIANCE/DF | \%DIFF | DELTA\% | CHISQUARE | PR>CHI |
| NULL | 0.7458 |  |  |  |  |
| YEAR | 0.7359 | 1.327 | 1.327 | 10.43 | 0.1077 |
|  |  |  |  |  |  |
| YEAR+ |  |  |  |  |  |
| SEASON | 0.7296 | 2.172 | 0.845 | 4.86 | 0.0879 |
| DEPTH | 0.7311 | 1.971 |  | 13.21 | 0.122 |
| AREA | 0.7312 | 1.958 |  | 3.9 | 0.2586 |
| SURVEY | 0.7318 | 1.877 |  | 2.77 | 0.1421 |
| HOOK | 0.732 | 1.850 |  | 0.0962 |  |

Table 2. The standardized index (number of sharks per set) of absolute abundance and coefficients of variation (CV) for all blacktip sharks. Indices are provided for the time series weighed by sample size or area.

| Year | Sets | Sample <br> size <br> index | CV |  | Area <br> index | CV |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
| 2004 | 44 | 2.23 | 0.26 |  | 2.49 | 0.27 |
| 2005 | 29 | 2.46 | 0.19 |  | 2.59 | 0.20 |
| 2006 | 127 | 2.11 | 0.11 |  | 2.18 | 0.12 |
| 2007 | 176 | 1.35 | 0.10 |  | 1.39 | 0.11 |
| 2008 | 210 | 1.34 | 0.09 |  | 1.30 | 0.10 |
| 2009 | 131 | 1.28 | 0.14 |  | 1.21 | 0.16 |
| 2010 | 176 | 1.89 | 0.10 |  | 1.86 | 0.10 |



Figure 1. Sampling universe for the combined Louisiana/Mississippi/Alabama bottom longline index. The study area was divided into 12 blocks: 11 blocks were the same size ( $156 \mathrm{~km}^{2}$ ), and one block (7) was larger ( $306 \mathrm{~km}^{2}$ ). Monthly sampling sites were randomly selected within each of the blocks.

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Figure 2. Standardized indices of abundance for all blacktip sharks. Time series are provided for indices weighed by sample size or area sampled. Each index has been divided by the maximum of the index


[^0]:    ${ }^{1}$ NOAA, National Marine Fisheries Service, Southeast Fisheries Science Center, Mississippi Laboratories, Pascagoula, Mississippi 39567; ${ }^{2}$ Center for Fisheries Research and Development, The University of Southern Mississippi, Gulf Coast Research Laboratory. 703 East Beach Drive. Ocean Springs, MS 39564; ${ }^{3}$ Dauphin Island Sea Lab, Center for Ecosystem Based Fishery Management, 101 Bienville Blvd, Dauphin Island, Alabama 36528; ${ }^{4}$ National Marine Fisheries Service, Southeast Fisheries Science Center, 3500 Delwood Beach Rd. Panama City, FL 32408

