Assessing injury on GOM cetaceans

August 31, 2022

Preamble

This is an Electronic Supplement to the manuscript Marques et al. "Quantifying Deepwater Horizon oil spill induced injury on pelagic cetaceans" submitted to Marine Ecology Progress Series (MEPS).

There are 7 Electronic Supplements to the paper. The master file containing links to all the other 6 additional Electronic Supplements related to this paper is ESO_ElectronicSupplements.

You might be reading this file as a pdf or as an html. The links on this file only work if you are using the html version of it, available via the github repository or if you compiled it yourself as html and you have all the 7 html files in the same folder. Otherwise, as a pdf distributed as an Electronic Supplement to the MEPS paper, the links might not work. They might work. If it is possible, we can work with the MEPS Editorial Office such that we can add links below that will link to actual files, say the pdfs of each of these 7 files, on the publisher server.

Version history

This section details the version history for static pdf files submitted as Electronic Supplement pdfs:

• 1.0 [12 Aug 2022] Version included as a pdf Electronic Supplement in the MEPS original submission

Introduction

In this document we present the results of the age, sex and class structured model implemented for all species/stock in the GOM.

We report here the injury metrics originally considered in Schwacke et al. (2017): (1) lost cetacean years (LCY), the difference between the baseline and injured population sizes, summed over the entire modeled time period; (2) years to recovery (YTR), the number of years required before the injured population trajectory reaches 95% of the baseline population trajectory; and (3) maximum proportional decrease (MPD), the difference between the two population trajectories when the injured trajectory is at its lowest point, divided by the baseline. Note that LCY is intuitively the metric that is most dependent on initial population size.

Running the simulations

The code below runs the simulations for all species, and stores the results in appropriate files, stored in appropriate species specific folders.

This code is included for completeness and does not need to be run as part of compiling this dynamic report as the results are already available in relevant files and it takes a considerably log time to run.

Creating objects to hold summary results:

The taxonomic units and the corresponding codes considered in this document are:

- Bwsp beaked whales, Beaked whales spp
- Fatt pygmy killer whale, Feresa attenuata
- Ggri Risso's dolphin, Grampus griseus

- Gmac short-finned pilot whale, Globicephala macrorhynchus
- Kosp Kogia species, Kogia sp.
- Pele melon-headed whale, Peponocephala electra
- Pmac sperm whale, Physeter macrocephalus
- Satt pantropical spotted dolphin, Stenella attenuata
- Sbre rough-toothed dolphin, Steno bredanensis
- Scly Clymene dolphin, Stenella clymene
- Scoe striped dolphin, Stenella coeruleoalba
- Sfro Atlantic spotted dolphin, Stenella frontalis
- Slon spinner dolphin, Stenella longirostris
- Ttro offshore bottlenose dolphins, Tursiops truncatus
- Ttrs shelf bottlenose dolphins, Tursiops truncatus

We re-order the results tables upfront so that results are organized by alphabetical order of the 4 letter code used to describe the taxonomic unit (this is also the order that the species are reported in tables 1 to 3 of the offshore paper).

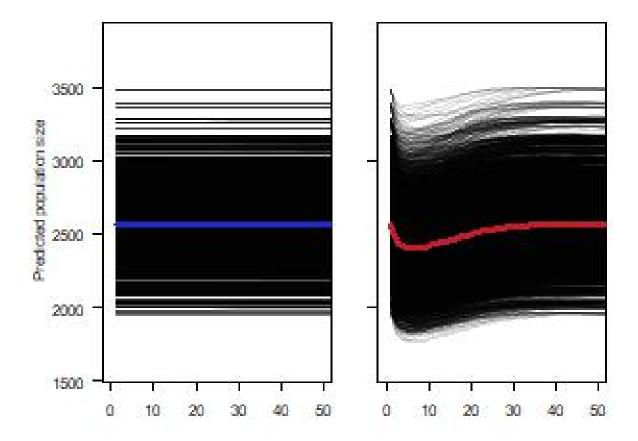
The data for initial population sizes and proportion exposed are imported from a txt file produced inside ES2 ElectronicSupplements.

Results by species

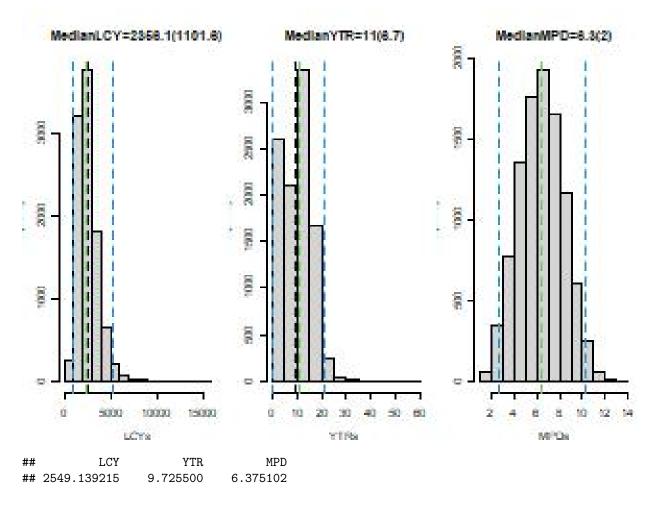
We represent in turn, for each species, the simulated population trajectories under the oil spill and under the baseline scenarios, as well as the histograms of the distributions of the 3 injury metrics for each species. To each histogram four dashed vertical lines were added:

- green median value for the injury metric
- black mean value for the injury metric
- blue 95% confidence interval for the injury metric

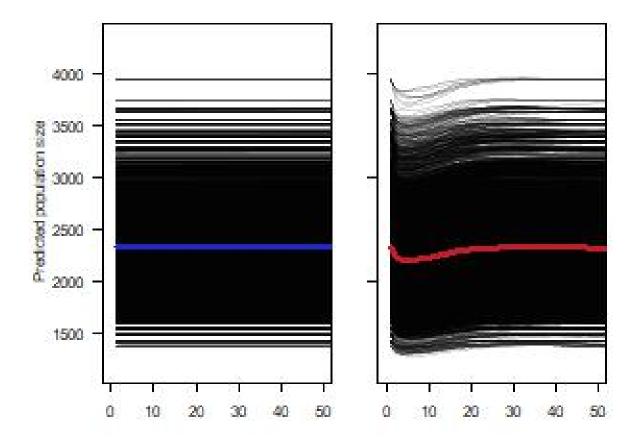
Physeter macrocephalus (Pmac, sperm whale)



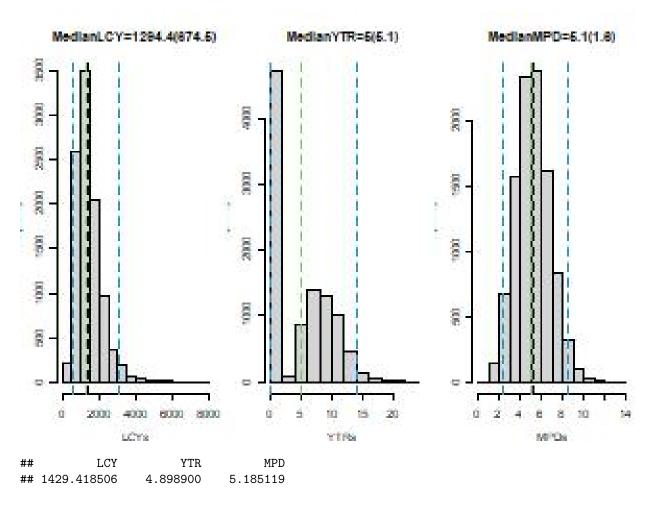
Mean for each of the injury metrics



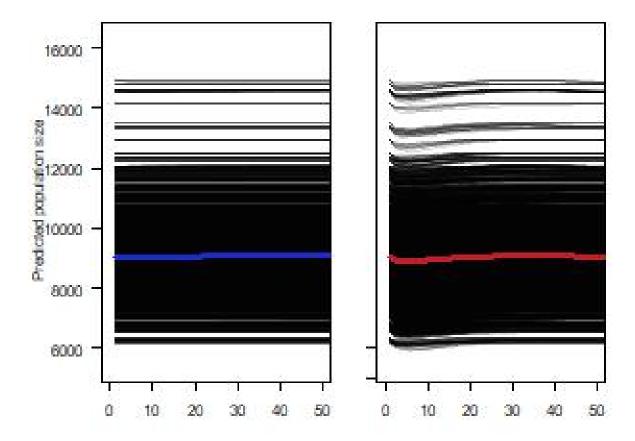
Kogia sp. (Kosp, kogia)



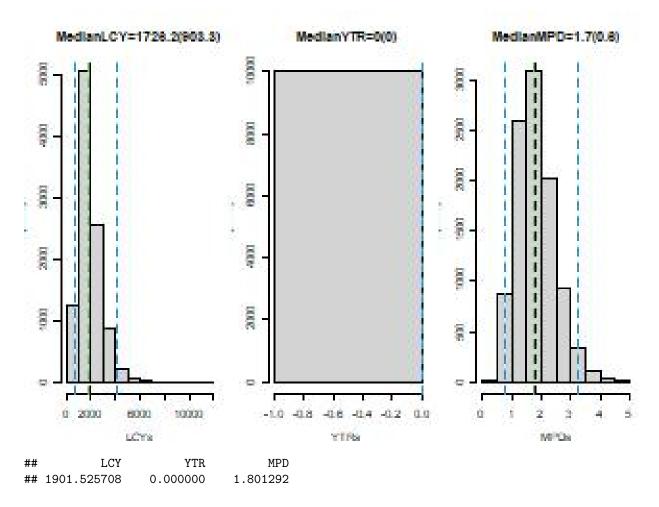
Mean for each of the injury metrics



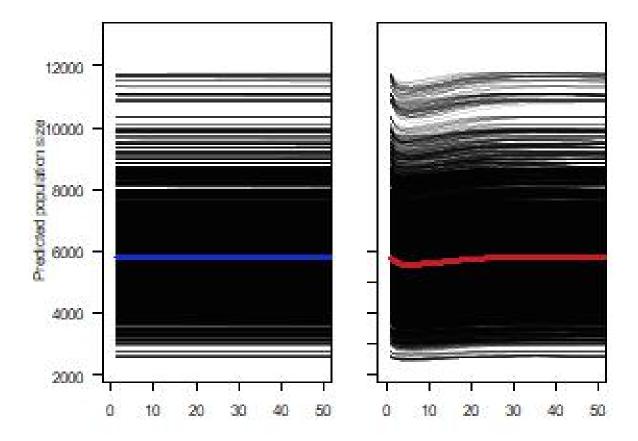
Stenella clymene (Scly, Clymene dolphin)



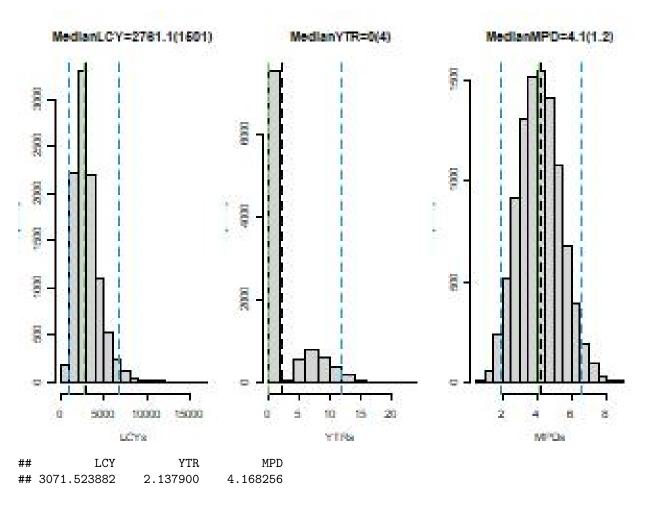
Mean for each of the injury metrics



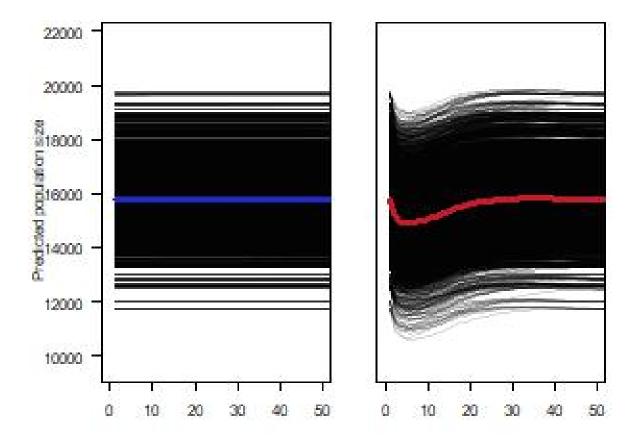
Peponocephala electra (Pele, melon-headed whale)



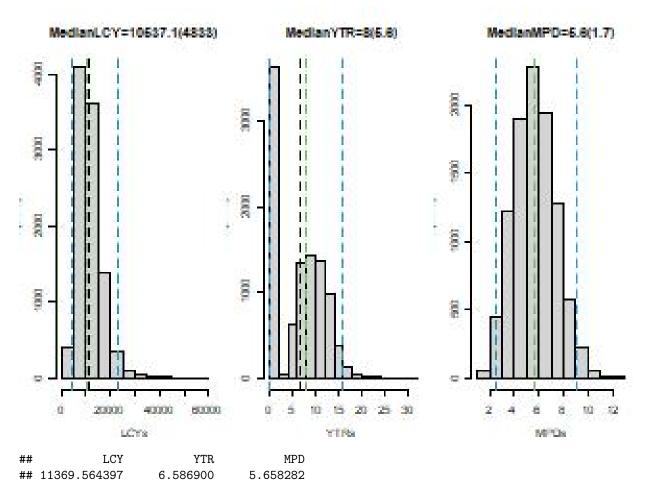
Mean for each of the injury metrics



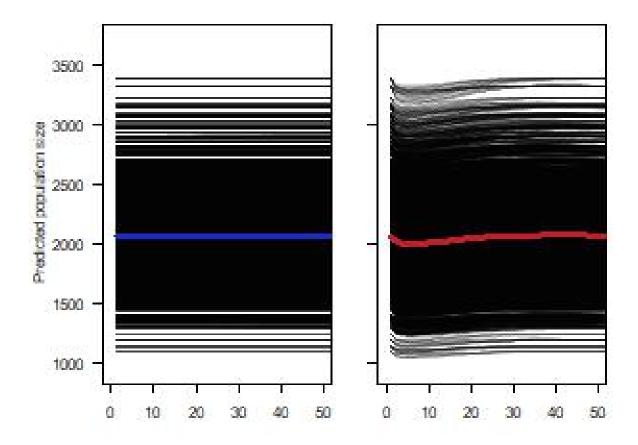
Tursiops truncatus (Ttro, offshore bottlenose dolphin)



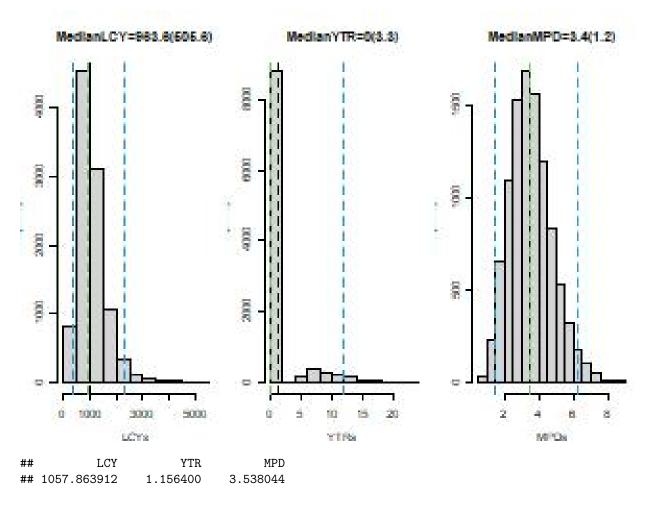
Mean for each of the injury metrics



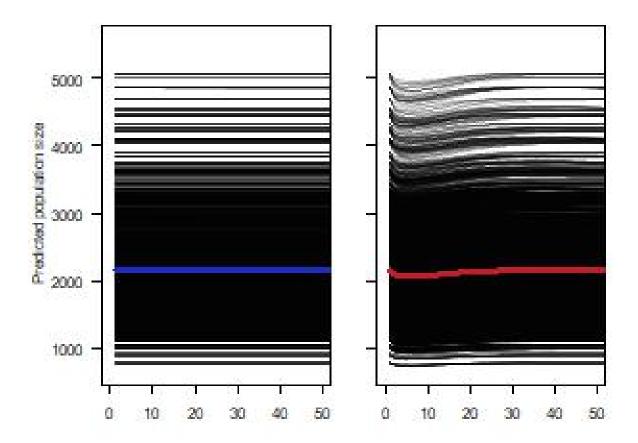
Globicephala macrorhynchus (Gmac, short finned pilot whale)



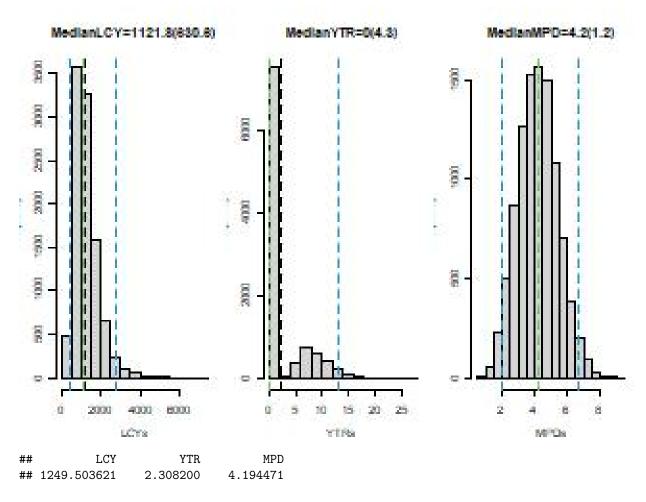
Mean for each of the injury metrics



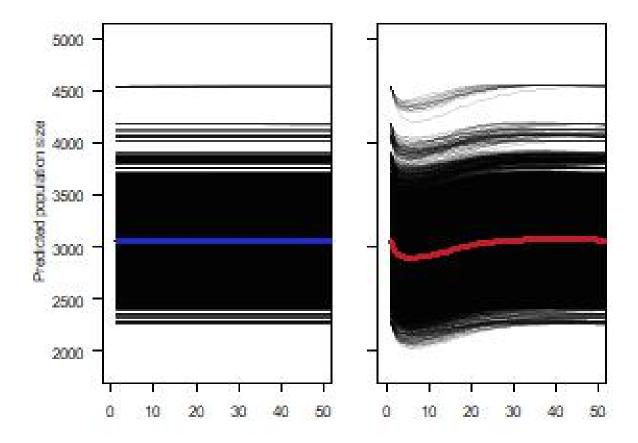
Feresa attenuata (Fatt, pygmy killer whale)



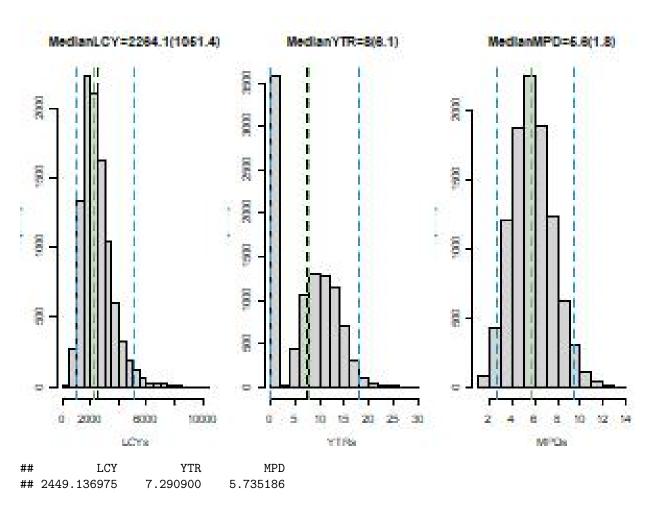
Mean for each of the injury metrics



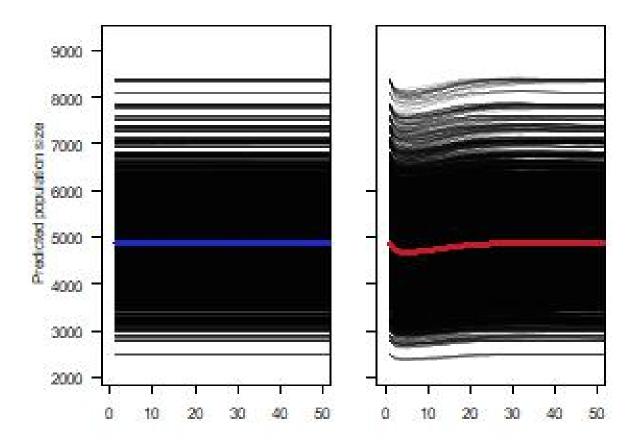
Grampus griseus (Ggri, Risso's dolphin)



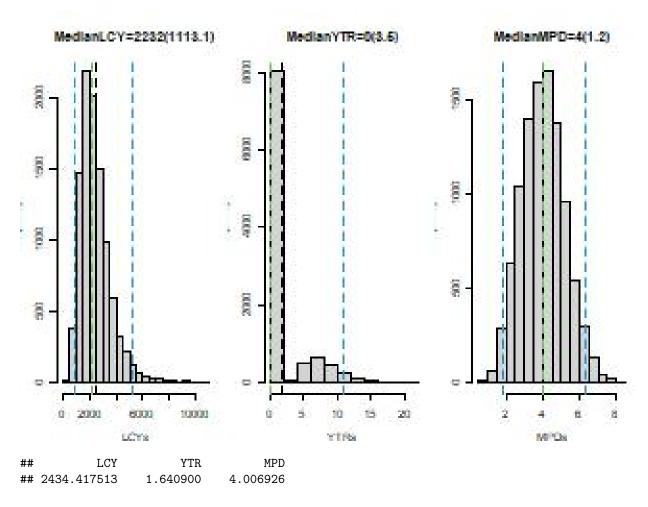
Mean for each of the injury metrics



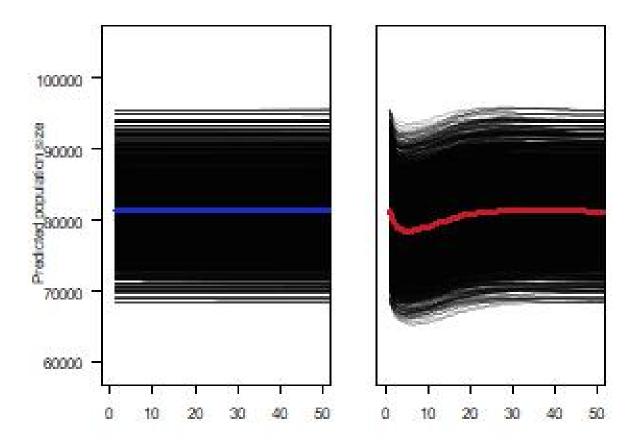
 $Steno\ bredanensis\ ({\bf Sbre},\ {\bf rough-toothed\ dolphin})$



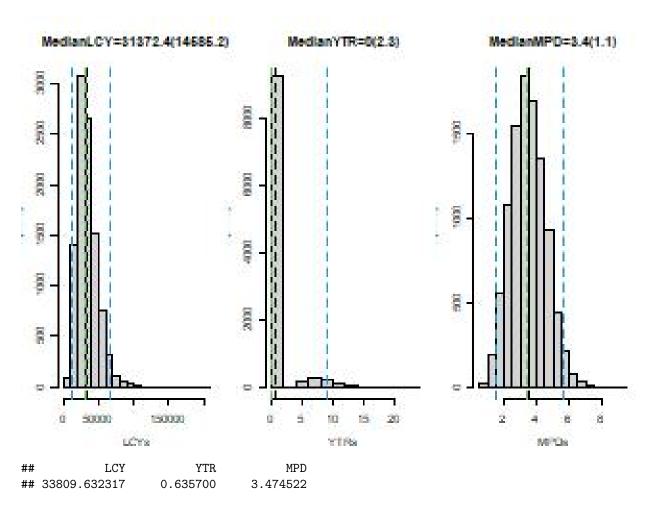
Mean for each of the injury metrics



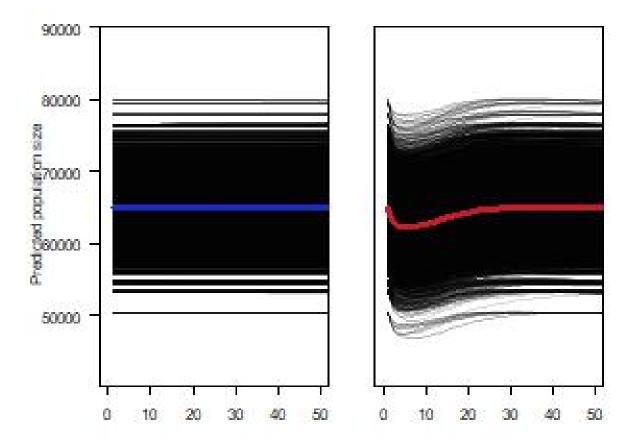
Stenella attenuata (Satt, pantropical spotted dolphin)



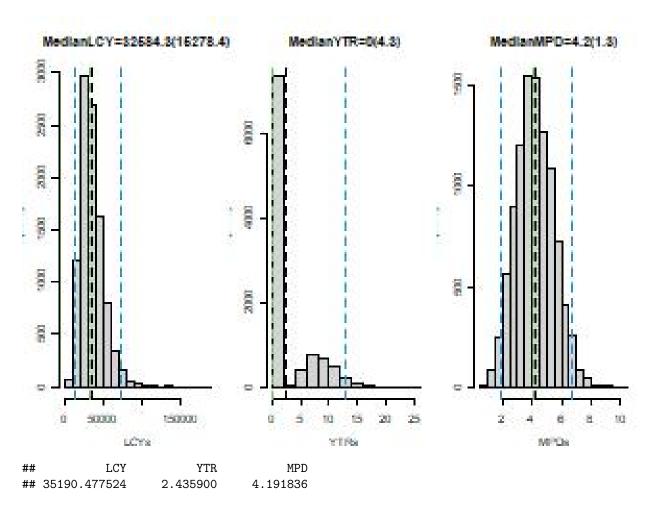
Mean for each of the injury metrics



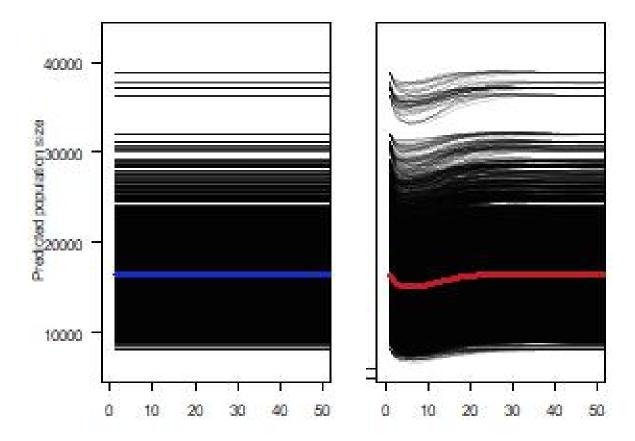
Tursiops truncatus (Ttrs, shelf bottlenose dolphin)



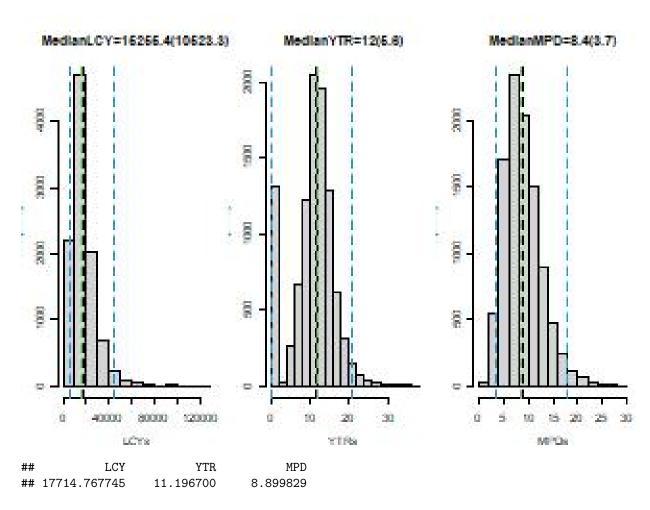
Mean for each of the injury metrics



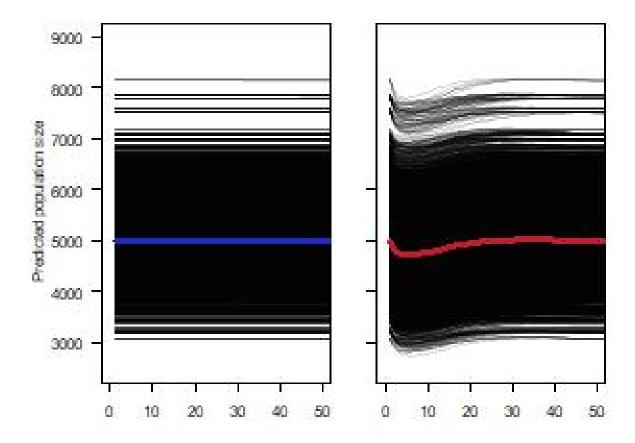
Stenella longirostris (Slon, spinner dolphin)



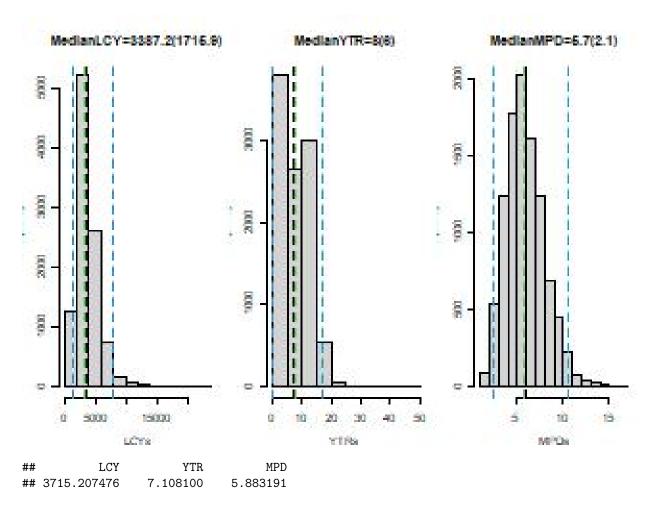
Mean for each of the injury metrics



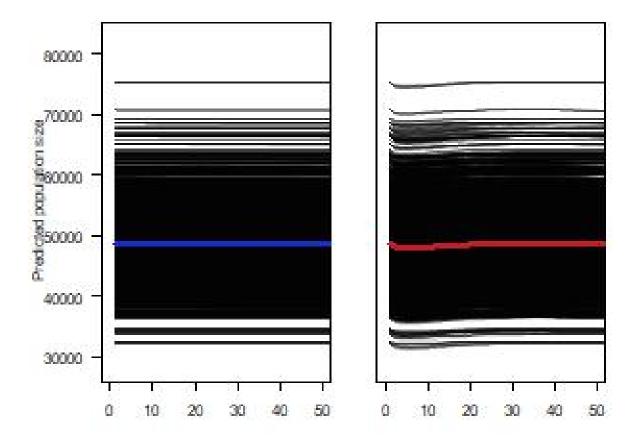
Stenella coeruleoalba (Scoe, striped dolphin)



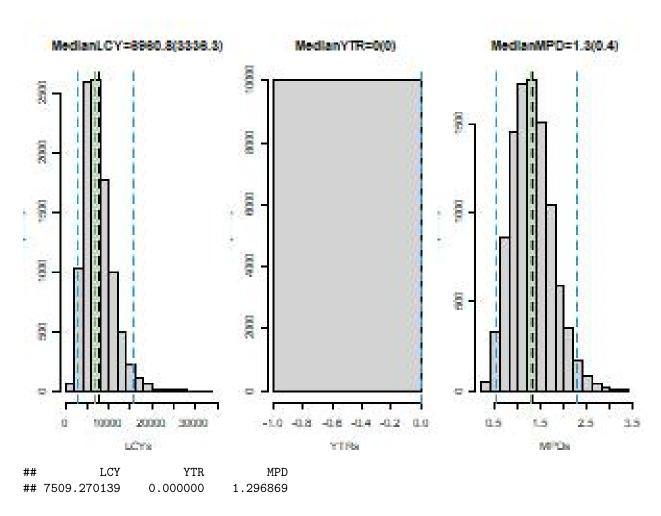
Mean for each of the injury metrics



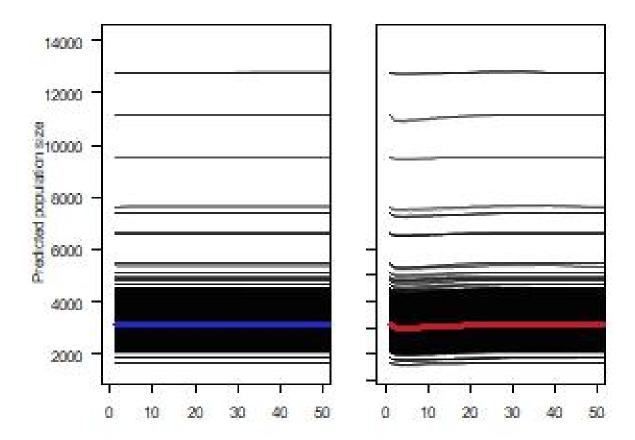
Stenella frontalis (Sfro, Atlantic spotted dolphin)



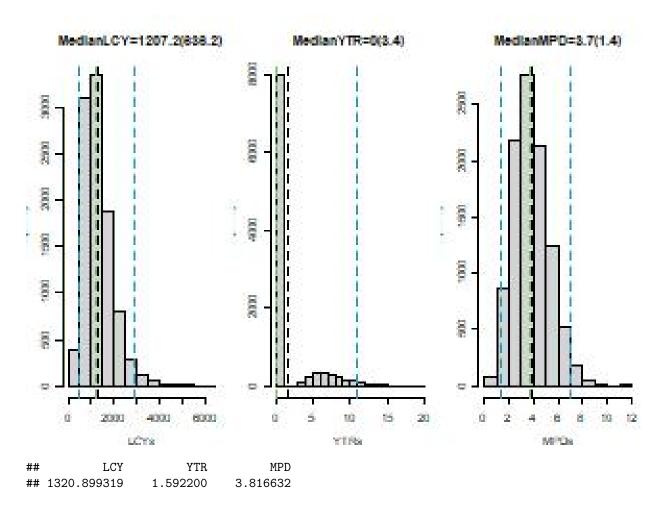
Mean for each of the injury metrics



Beaked whales spp (Bwsp)



Mean for each of the injury metrics



Comparisons across species

In this section we present some summary analysis of the results across the 15 taxonomic units considered.

Initial population and proportion exposed

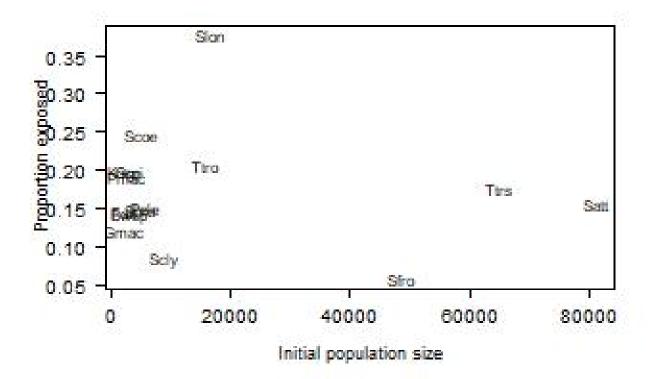
The population size and the proportion exposed are apriori expected to be key determinants of injury, since the population size will have a direct impact on LCY and the proportion exposed should have a direct impact on all 3 measures.

Table 1: Initial population size (N) and proportion exposed (Pexp) for the different stocks considered.

| Sp | Nstart | Pexp | | |
|------|--------|-------|--|--|
| Bwsp | 3098 | 0.140 | | |
| Fatt | 2152 | 0.145 | | |
| Ggri | 3063 | 0.197 | | |
| Gmac | 2065 | 0.120 | | |
| Kosp | 2322 | 0.197 | | |
| Pele | 5784 | 0.152 | | |
| Pmac | 2561 | 0.191 | | |
| Satt | 81233 | 0.155 | | |
| Sbre | 4867 | 0.148 | | |

| Sp | Nstart | Pexp |
|-----------------------|--------|-------|
| Scly | 9065 | 0.082 |
| Scoe | 5011 | 0.247 |
| Sfro | 48688 | 0.058 |
| Slon | 16501 | 0.377 |
| Ttro | 15791 | 0.206 |
| Ttrs | 64897 | 0.177 |

The proportion exposed against the initial population size is shown here for the 15 taxonomic units



Injury metrics

In the following table we present the means of the 3 injury metrics:

Table 2: Injury results for the different stocks considered.

| Sp | LCY | MPD | YTR |
|------|------|--------|-----|
| Bwsp | 1321 | -0.038 | 1.6 |
| Fatt | 1250 | -0.042 | 2.3 |
| Ggri | 2449 | -0.057 | 7.3 |
| Gmac | 1058 | -0.035 | 1.2 |
| Kosp | 1429 | -0.052 | 4.9 |
| Pele | 3072 | -0.042 | 2.1 |
| Pmac | 2549 | -0.064 | 9.7 |

| Sp | LCY | MPD | YTR |
|-----------------------|-------|--------|------|
| Satt | 33810 | -0.035 | 0.6 |
| Sbre | 2434 | -0.040 | 1.6 |
| Scly | 1902 | -0.018 | 0.0 |
| Scoe | 3715 | -0.059 | 7.1 |
| Sfro | 7509 | -0.013 | 0.0 |
| Slon | 17715 | -0.089 | 11.2 |
| Ttro | 11370 | -0.057 | 6.6 |
| Ttrs | 35190 | -0.042 | 2.4 |

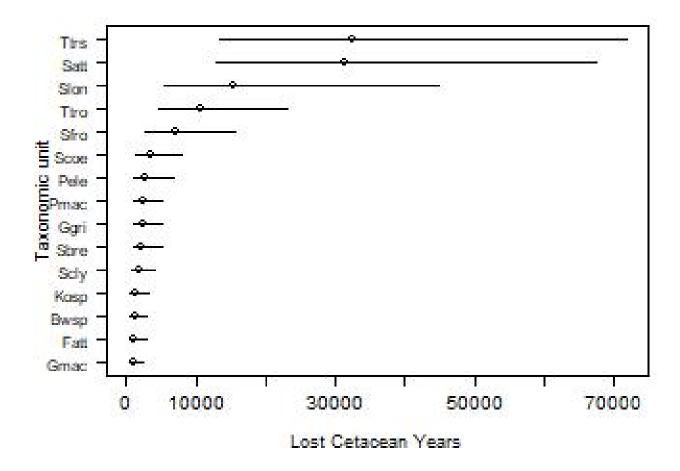
In the following table we present the medians and the 95% confidence intervals for our current estimates of injury. These correspond to the results in table 3 in the main paper.

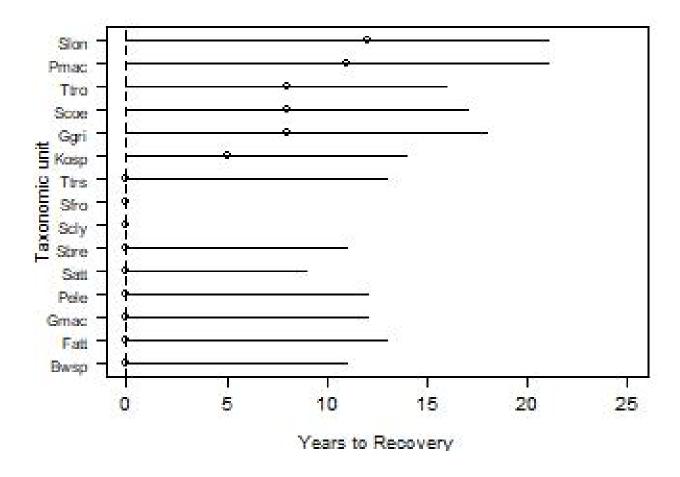
Table 3: Median injury results for the different stocks considered and respective 95% confidence intervals by the percentile method

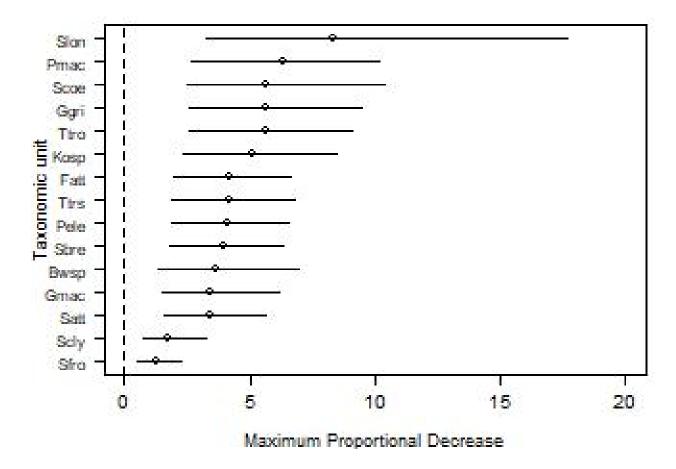
| Sp | LCYmed | LCYl | LCYu | MPDmed | MPDl | MPDu | YTRmed | YTRl | YTRu |
|-----------------------|--------|-------|-------|--------|-------|--------|--------|------|------|
| Bwsp | 1207 | 445 | 2899 | 3.683 | 1.386 | 6.951 | 0 | 0 | 11 |
| Fatt | 1122 | 437 | 2809 | 4.186 | 1.939 | 6.642 | 0 | 0 | 13 |
| Ggri | 2264 | 978 | 5049 | 5.623 | 2.569 | 9.494 | 8 | 0 | 18 |
| Gmac | 964 | 385 | 2291 | 3.435 | 1.486 | 6.216 | 0 | 0 | 12 |
| Kosp | 1294 | 519 | 3112 | 5.114 | 2.315 | 8.492 | 5 | 0 | 14 |
| Pele | 2761 | 1081 | 6767 | 4.140 | 1.925 | 6.601 | 0 | 0 | 12 |
| Pmac | 2356 | 996 | 5240 | 6.344 | 2.672 | 10.189 | 11 | 0 | 21 |
| Satt | 31372 | 12884 | 67606 | 3.433 | 1.563 | 5.615 | 0 | 0 | 9 |
| Sbre | 2232 | 901 | 5207 | 3.995 | 1.836 | 6.349 | 0 | 0 | 11 |
| Scly | 1726 | 664 | 4067 | 1.734 | 0.747 | 3.261 | 0 | 0 | 0 |
| Scoe | 3387 | 1369 | 7897 | 5.673 | 2.479 | 10.426 | 8 | 0 | 17 |
| Sfro | 6961 | 2722 | 15509 | 1.264 | 0.542 | 2.280 | 0 | 0 | 0 |
| Slon | 15255 | 5302 | 44903 | 8.363 | 3.277 | 17.716 | 12 | 0 | 21 |
| Ttro | 10537 | 4597 | 23220 | 5.610 | 2.546 | 9.125 | 8 | 0 | 16 |
| Ttrs | 32584 | 13377 | 71967 | 4.152 | 1.859 | 6.758 | 0 | 0 | 13 |

These are the values that are used in table 3 of the paper, and the code to do so is in the .Rmd.

Representing the above information visually, focusing on each of the 3 injury metrics:







Exploring the injury metric results

In this section we explore the results obtained in terms of the different injury metrics and how these injury metrics might be explained by the different input parameters. To do so we consider linear models to model the mean value of the injury per species as a function of the mean values of the input parameters. After a preliminary exploratory analysis, only the following four parameters seem to have a non-negligible potential relevant effect on the injury metrics:

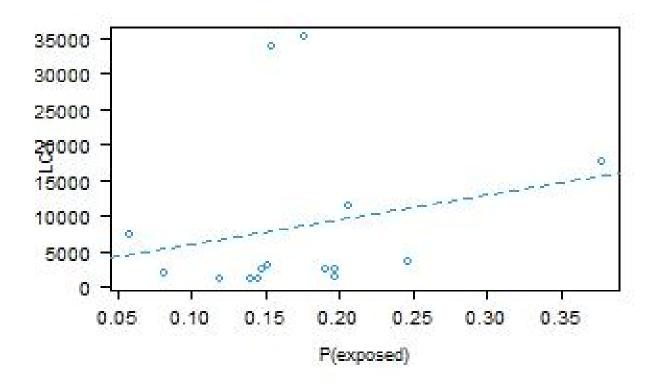
- Proportion Exposed
- Initial population size
- Gestation duration
- Survival reduction

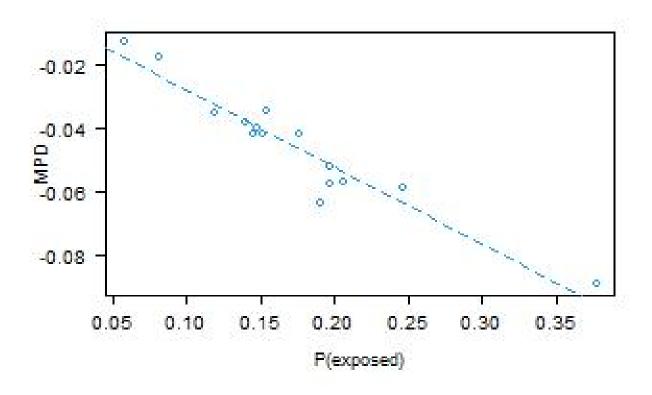
Exploratory analysis

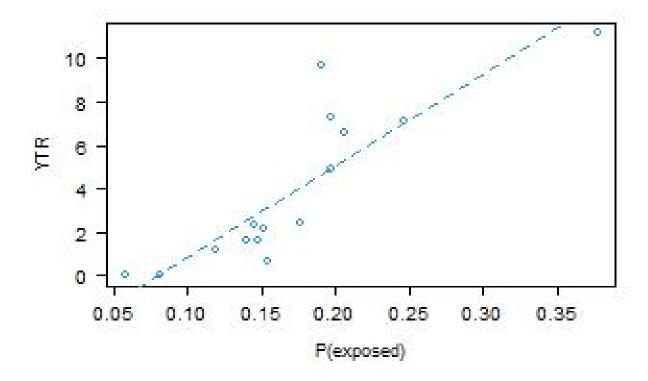
Here we plot the values of the injury metrics as a function of the four key parameters going into the model.

Proportion exposed

The next three plots describe the relationship between the proportion exposed and Lost Cetacean Years, the Maximum Proportion Decrease and the Years to recovery, respectively

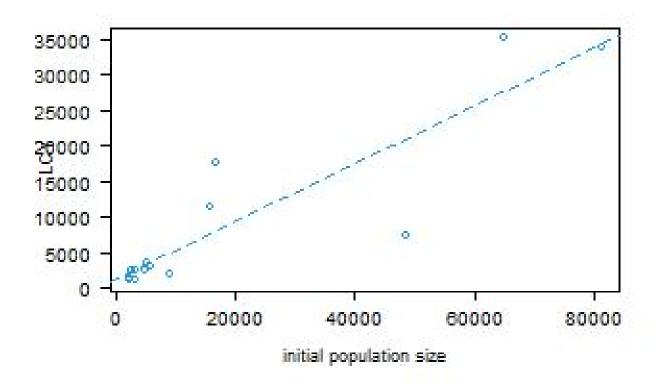


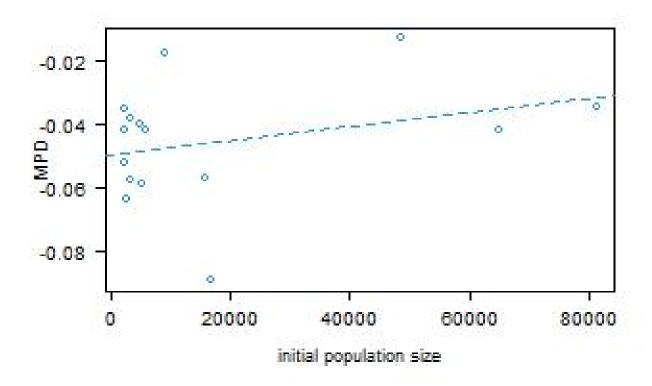


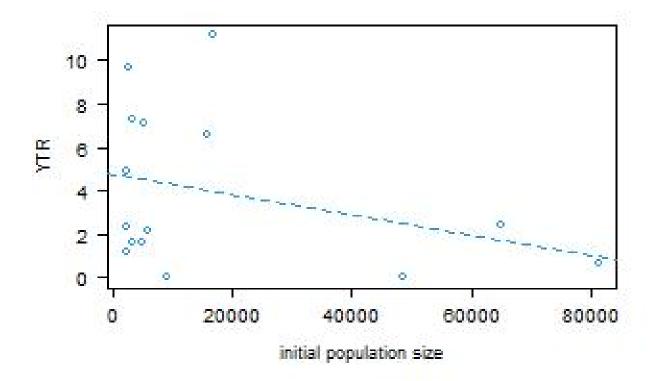


Initial population size

The next three plots describe the relationship between the initial population size and Lost Cetacean Years, the Maximum Proportion Decrease and the Years to recovery, respectively.

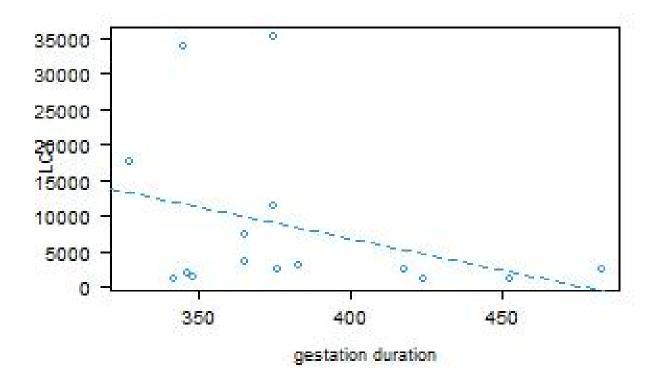


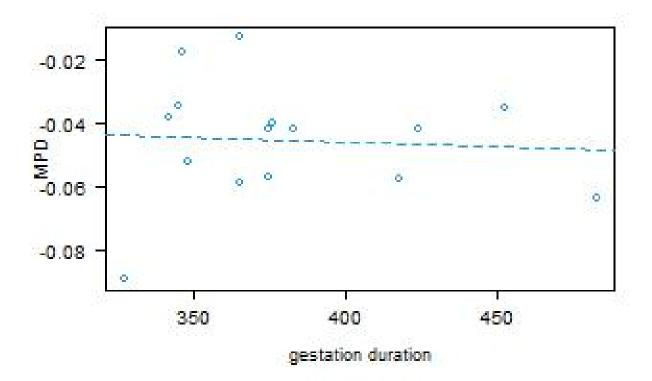


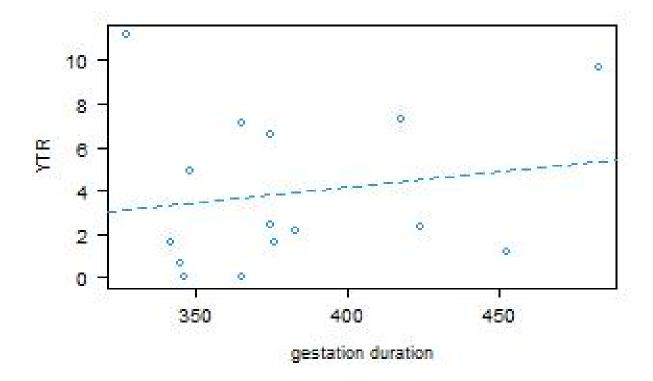


Gestation duration

The next three plots describe the relationship between the gestation duration and Lost Cetacean Years, the Maximum Proportion Decrease and the Years to recovery, respectively

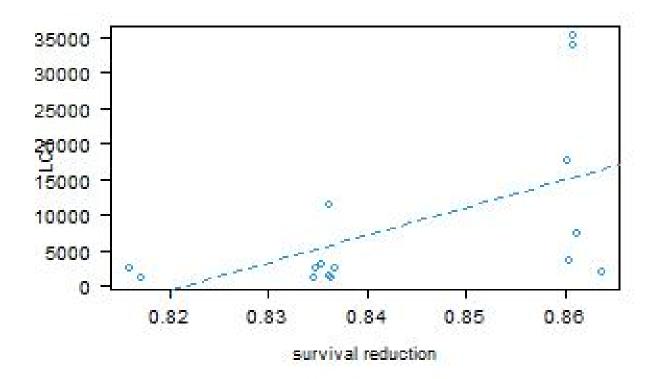


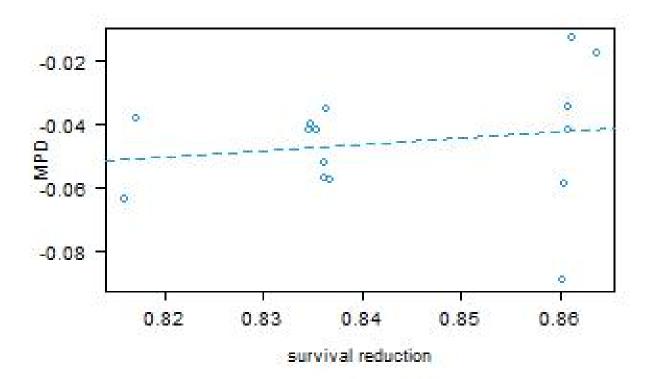


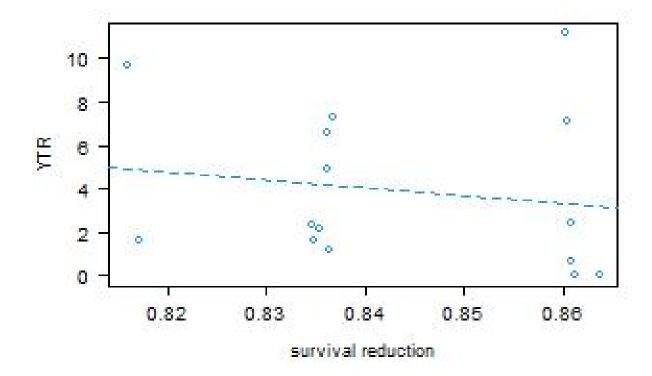


Survival reduction

The next three plots describe the relationship between the survival reduction and Lost Cetacean Years, the Maximum Proportion Decrease and the Years to recovery, respectively







Explaining injury metrics across species

Here we consider standard linear models to explain the injury metrics as a function of the input parameters. To obtain what we considered the most parsimonious model we conducted a traditional model selection approach using AIC, selecting the model with the lowest AIC.

We considered only the 4 explanatory variables which, given the exploratory data analysis above, could explain the injury metrics. Given only 4 explanatory variables, we conducted a full search over the possible models, ignoring interactions, using function bestglm from the package bestglm (McLeod & Lai, 2020).

LCY

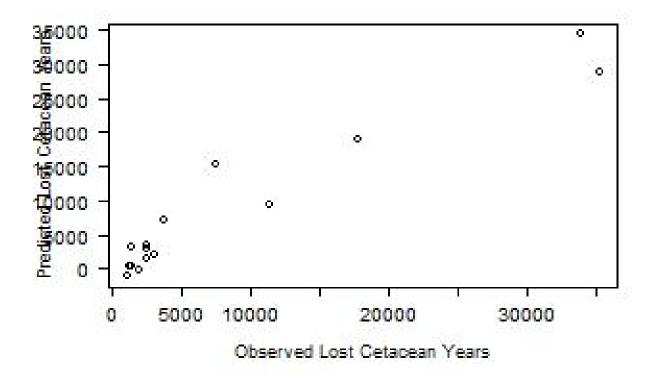
The summary of the best linear model follows.

```
## BICq equivalent for q in (0.00257603089392322, 0.727813974603144)
## Best Model:
##
                    Estimate
                                 Std. Error
                                               t value
                                                               Pr(>|t|)
## (Intercept) -8585.8086023
                              2457.09226124 -3.494296 0.00442815340683
                                 0.03567448 11.952412 0.00000005054783
## N
                   0.4263961
## p
               54656.6056191 12278.22336920 4.451508 0.00079078410382
##
## Call:
## lm(formula = y ~ p + N, data = LCY4bestglm)
##
## Residuals:
                1Q Median
##
       Min
                                3Q
                                        Max
```

```
## -7835.4 -1189.7
                     855.8 1477.7 6430.2
##
## Coefficients:
##
                                                   Pr(>|t|)
                  Estimate
                           Std. Error t value
## (Intercept) -8585.80860
                            2457.09226
                                        -3.494
                                                   0.004428 **
              54656.60562 12278.22337
                                         4.452
                                                   0.000791 ***
## p
## N
                               0.03567 11.952 0.0000000505 ***
                   0.42640
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 3380 on 12 degrees of freedom
## Multiple R-squared: 0.9263, Adjusted R-squared:
## F-statistic: 75.45 on 2 and 12 DF, p-value: 0.0000001598
We can look at the ANOVA table for the most parsimonious model based on lowest AIC criteria.
## Analysis of Variance Table
##
## Response: y
##
             Df
                              Mean Sq F value
                                                      Pr(>F)
                    Sum Sq
## p
              1
                  91857453
                             91857453
                                        8.0395
                                                     0.01502 *
## N
              1 1632291730 1632291730 142.8602 0.00000005055 ***
## Residuals 12
                137109617
                             11425801
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

Using the most parsimonious model, the proportion of the variability explained by each input retained in the regression model and the remaining unexplained variation can be obtained from the ANOVA table of the linear regression (as shown above). The proportion of variance explained in Lost Cetacean Years by the initial population size is 87.7 %, while the proportion exposed accounts for 4.94 %. Only 7.37 of the variability remains unexplained.

We illustrate how well the model predicts the observed values in the plot below:



YTR

The summary of the best linear model follows.

```
## AIC
## BICq equivalent for q in (0.0565670801885912, 0.714907105110615)
## Best Model:
##
                   Estimate Std. Error
                                          t value
                                                        Pr(>|t|)
## (Intercept) -14.46579200 3.864204450 -3.743537 0.002803820209
                0.02778531 0.009303748 2.986464 0.011348598337
## p
               44.99620475 5.546928494 8.111914 0.000003260501
##
## Call:
## lm(formula = y ~ p + gd, data = YTR4bestglm)
## Residuals:
##
     Min
             1Q Median
## -2.364 -1.229 0.256 1.220
                               2.177
##
## Coefficients:
                Estimate Std. Error t value
                                               Pr(>|t|)
## (Intercept) -14.465792
                            3.864204
                                     -3.744
                                                 0.0028 **
               44.996205
                            5.546928
                                      8.112 0.00000326 ***
## p
## gd
                            0.009304
                0.027785
                                      2.986
                                                 0.0113 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

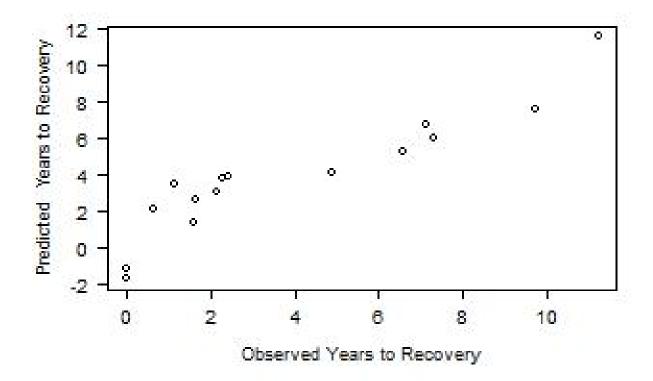
```
##
## Residual standard error: 1.515 on 12 degrees of freedom
## Multiple R-squared: 0.8502, Adjusted R-squared: 0.8253
## F-statistic: 34.06 on 2 and 12 DF, p-value: 0.00001129
```

We can look at the ANOVA table for the most parsimonious model based on lowest AIC criteria.

```
## Analysis of Variance Table
##
## Response: y
##
             \mathsf{Df}
                 Sum Sq Mean Sq F value
                                              Pr(>F)
## p
              1 135.955 135.955
                                 59.199 0.000005592 ***
                 20.483
                         20.483
                                   8.919
                                             0.01135 *
## gd
              1
## Residuals 12
                 27.559
                           2.297
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

The proportion of variance explained in Years to Recovery by the proportion exposed is 73.89 %, while the gestation duration accounts for 11.13 %. About 14.98 % of the total variability remains unexplained.

We illustrate how well the model predicts the observed values in the plot below:



MPD

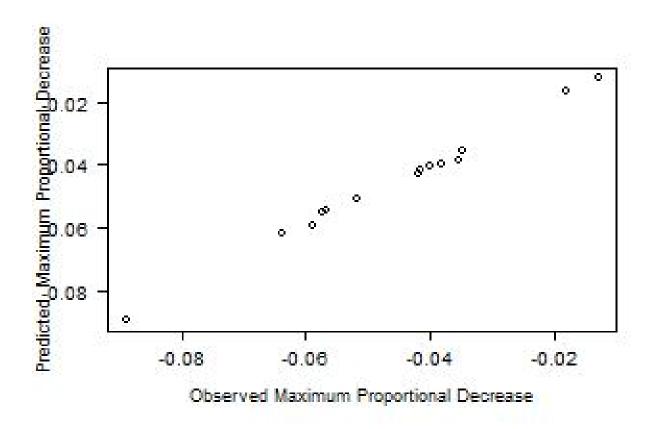
The summary of the best linear model follows.

```
## AIC
## BICq equivalent for q in (0.000181541676479879, 0.791379587004136)
## Best Model:
```

```
##
                                             t value
                                                                  Pr(>|t|)
                    Estimate
                                Std. Error
## (Intercept) -0.14174704099 0.02968258662 -4.775428 0.0005756315584552402
              -0.00006625358 0.00001198643 -5.527382 0.0001787223801420487
               0.19539414143 0.03189438029
                                            6.126287 0.0000745371892281724
## sr
## p
              -0.25025550911 0.00602766599 -41.517813 0.0000000000001925047
##
## Call:
## lm(formula = y ~ p + gd + sr, data = MPD4bestglm)
##
## Residuals:
##
                            Median
                                           3Q
                                                    Max
         Min
                     1Q
  -0.0021024 -0.0013630 0.0004400 0.0007655
##
## Coefficients:
##
                                                      Pr(>|t|)
                 Estimate
                          Std. Error t value
## (Intercept) -0.14174704
                           0.02968259
                                      -4.775
                                                      0.000576 ***
## p
              ## gd
              -0.00006625 0.00001199
                                      -5.527
                                                      0.000179 ***
               0.19539414 0.03189438
                                       6.126 0.000074537189228 ***
## sr
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 0.001646 on 11 degrees of freedom
## Multiple R-squared: 0.9939, Adjusted R-squared: 0.9922
## F-statistic: 594.3 on 3 and 11 DF, p-value: 0.00000000001911
We can look at the ANOVA table for the most parsimonious model based on lowest AIC criteria.
## Analysis of Variance Table
##
## Response: y
##
            Df
                  Sum Sq
                           Mean Sq F value
                                                       Pr(>F)
             1 0.0044337 0.0044337 1635.854 0.0000000000002563 ***
## p
## gd
             1 0.0002966 0.0002966 109.431 0.0000004704063385 ***
             1 0.0001017 0.0001017
                                     37.531 0.0000745371892282 ***
## sr
## Residuals 11 0.0000298 0.0000027
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

The proportion of variance explained in MPD by the proportion exposed is 91.19, while the gestation duration and the survival reduction only account for only 6.1~% and 2.09~%, respectively. About 0.61~% of the variability remains unexplained.

We illustrate how well the model predicts the observed values in the plot below:



References

DWH MMIQT 2015, Models and analyses for the quantification of injury to Gulf of Mexico cetaceans from the Deepwater Horizon Oil Spill, MM_TR.01_Schwacke_Quantification.of.lnjury.to.GOM.Cetaceans LINK.

McLeod A, Xu C, Lai Y (2020). bestglm: Best Subset GLM and Regression Utilities. R package version 0.37.3, https://CRAN.R-project.org/package=bestglm.

Schwacke, L.H., L. Thomas, R.S. Wells, W.E. McFee, A.A. Hahn, K.D. Mullin, E.S. Zolman, B.M. Quigley, T.K. Rowles and J.H. Schwacke. 2017. An age-, sex- and class-structured population model for estimating nearshore common bottlenose dolphin injury following the Deepwater Horizon oil spill. *Endangered Species Research* 33: 265-279. DOI: 10.3354/esr00777.