

Assessing injury on GOM cetaceans

August 11, 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 long 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).

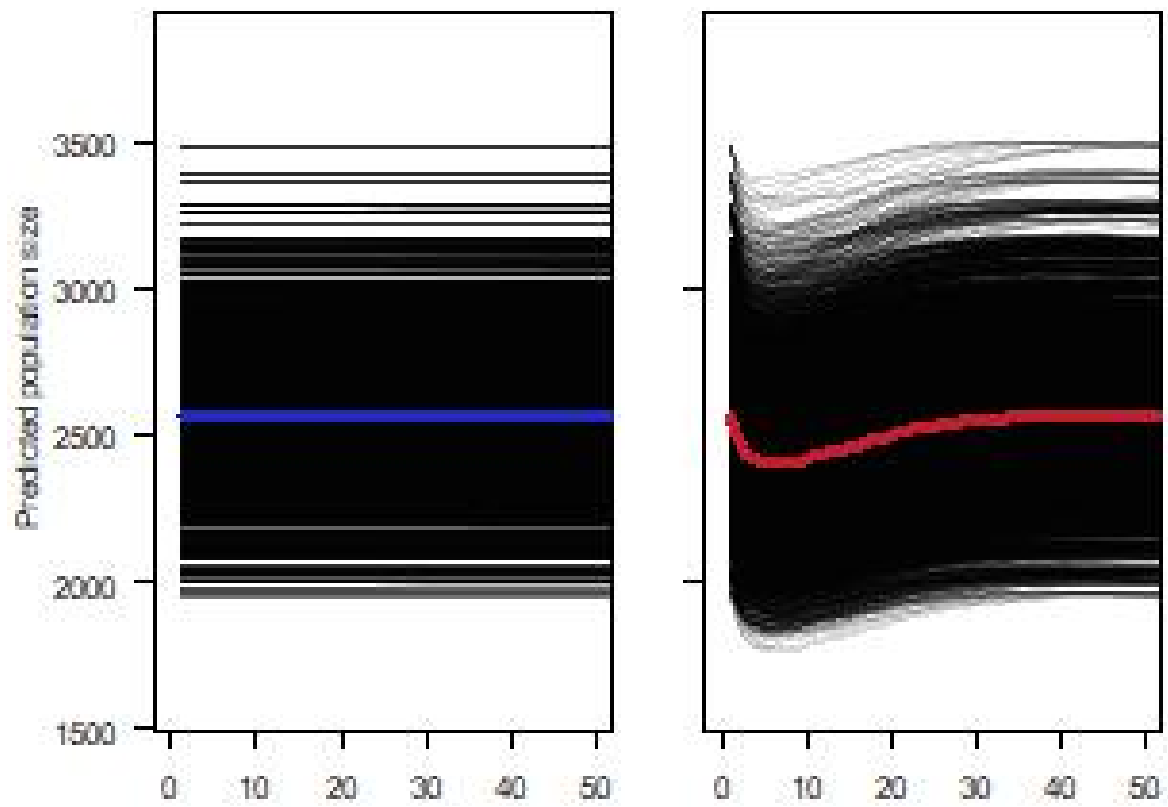
The data for initial population sizes and proportion exposed are imported from a `txt` file produced inside [ES2_ElectronicSupplements](#).

Results by species

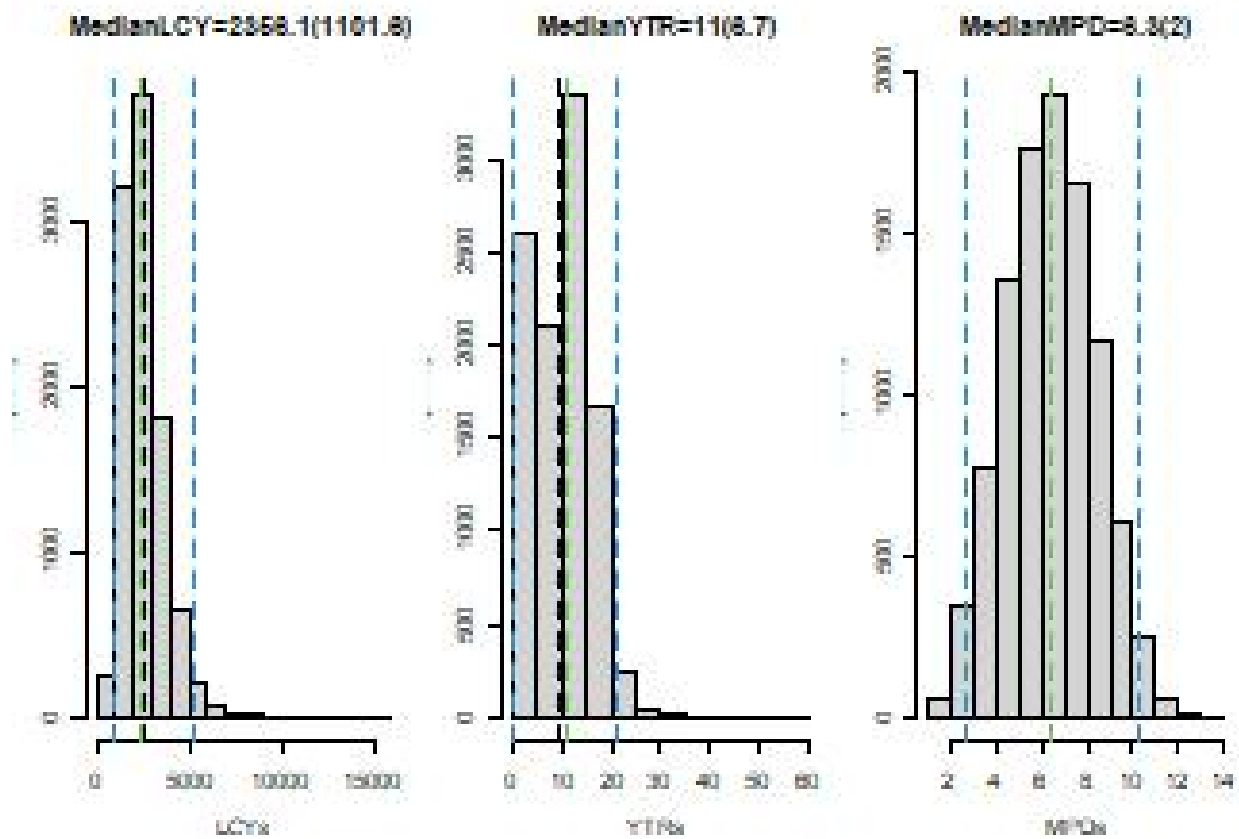
We represent in turn, for each species, the simulated population trajectories under the oil spill and under the baseline scenarios, as well as the histograms of the distributions of the 3 injury metrics for each species.

Physeter macrocephalus (Pmac, sperm whale)

Population trajectories



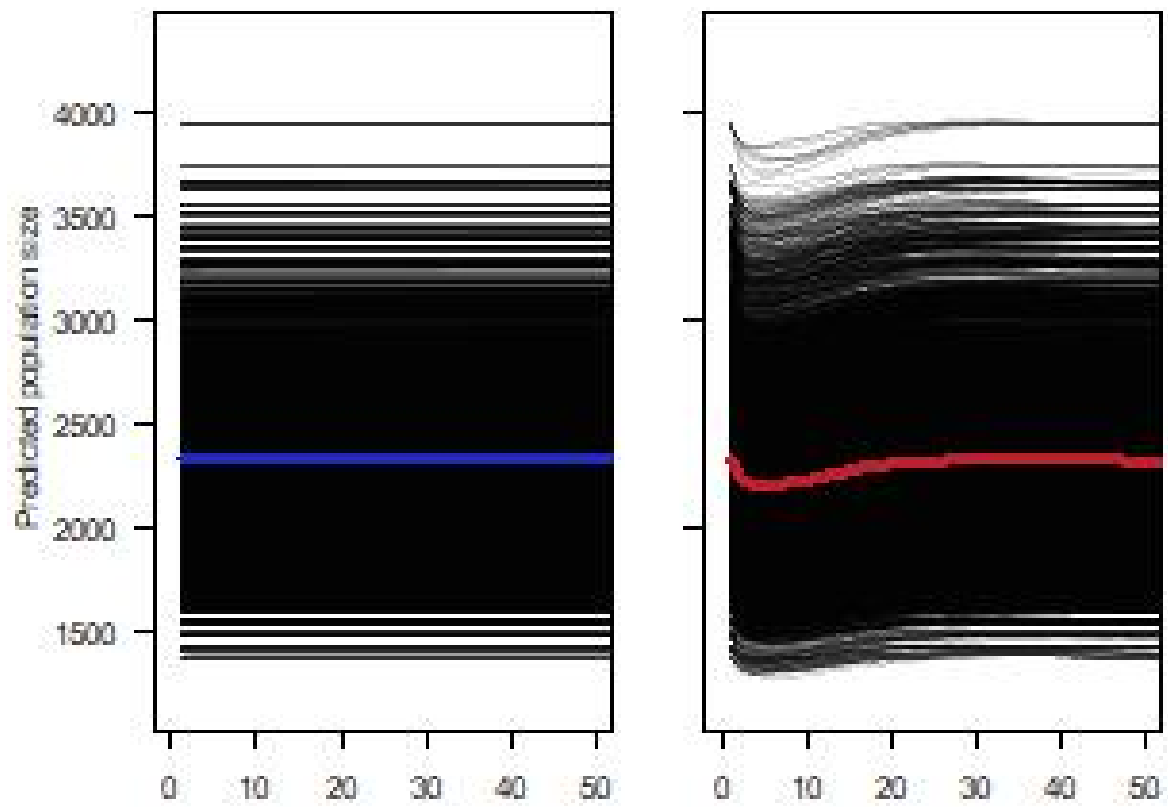
Mean for each of the injury metrics



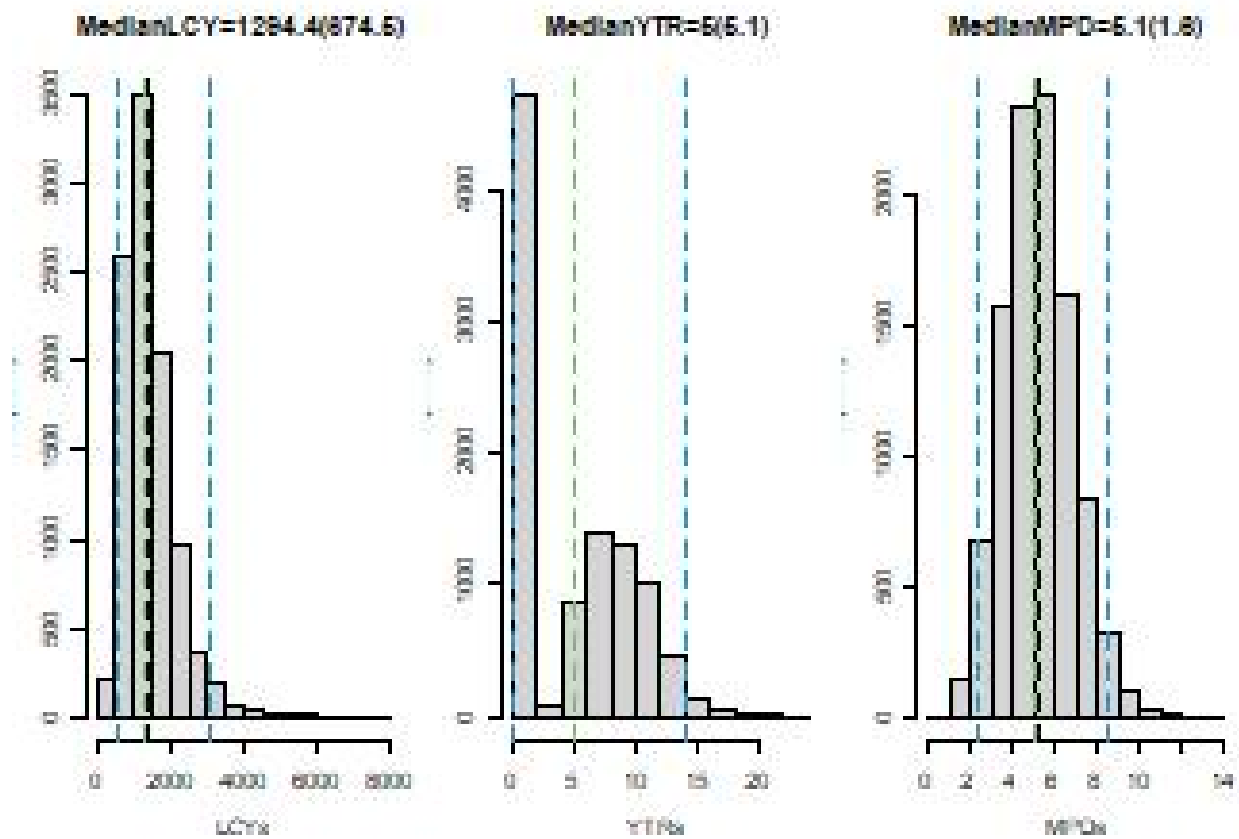
##	LCY	YTR	MPD
##	2549.139215	9.725500	6.375102

***Kogia* sp. (Kosp, kogia)**

Population trajectories



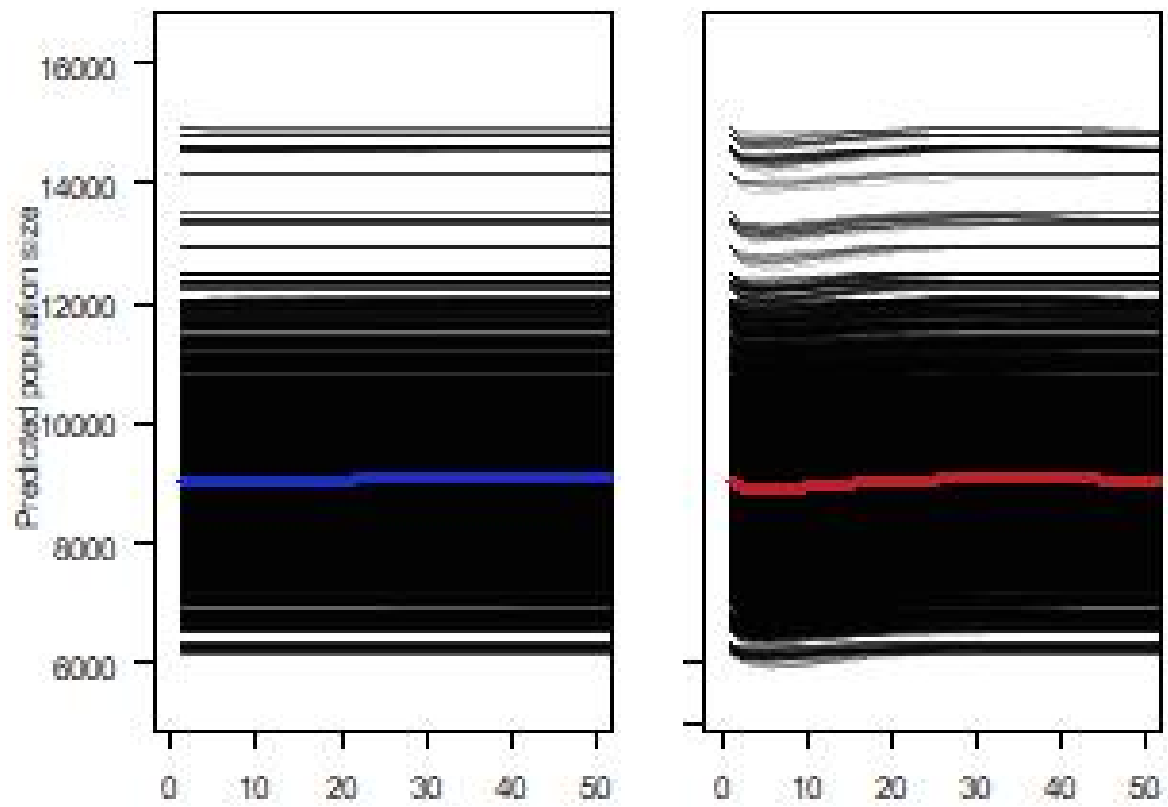
Mean for each of the injury metrics



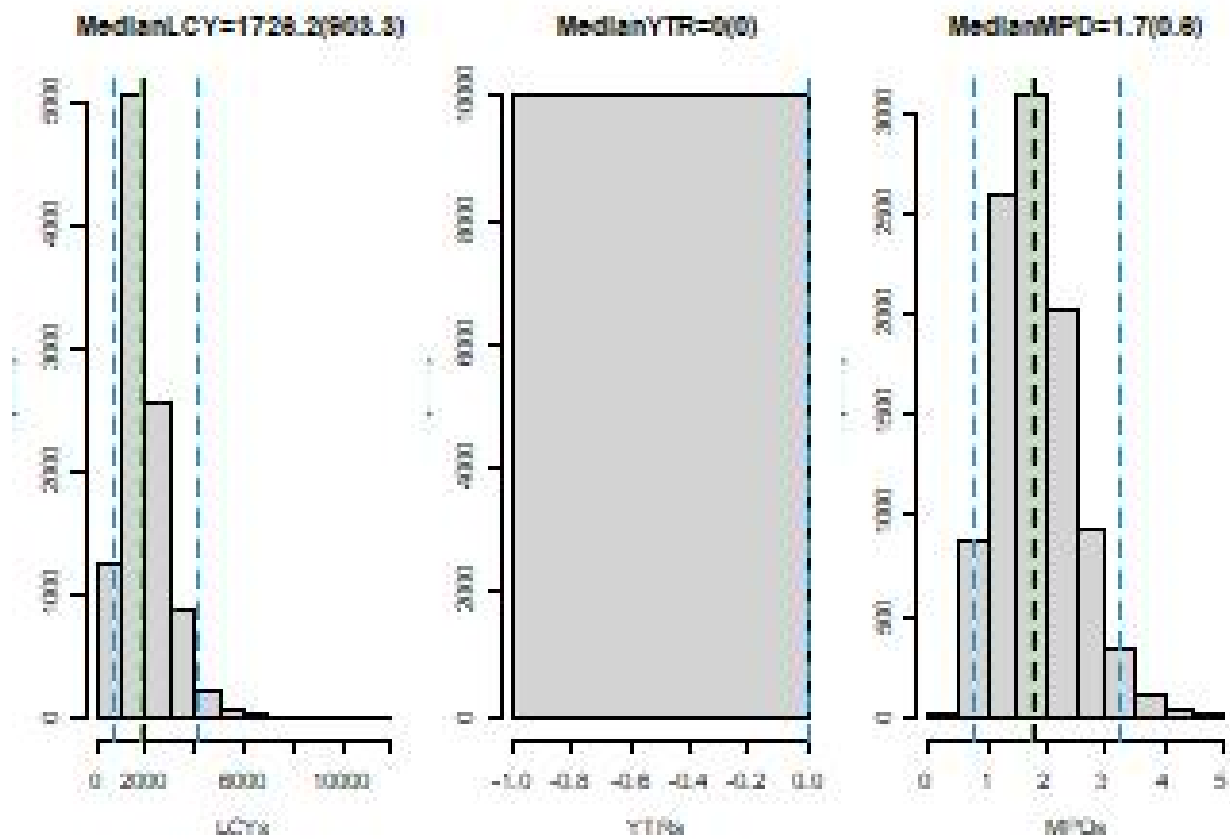
##	LCY	YTR	MPD
##	1429.418506	4.898900	5.185119

Stenella clymene (Scly, Clymene dolphin)

Population trajectories



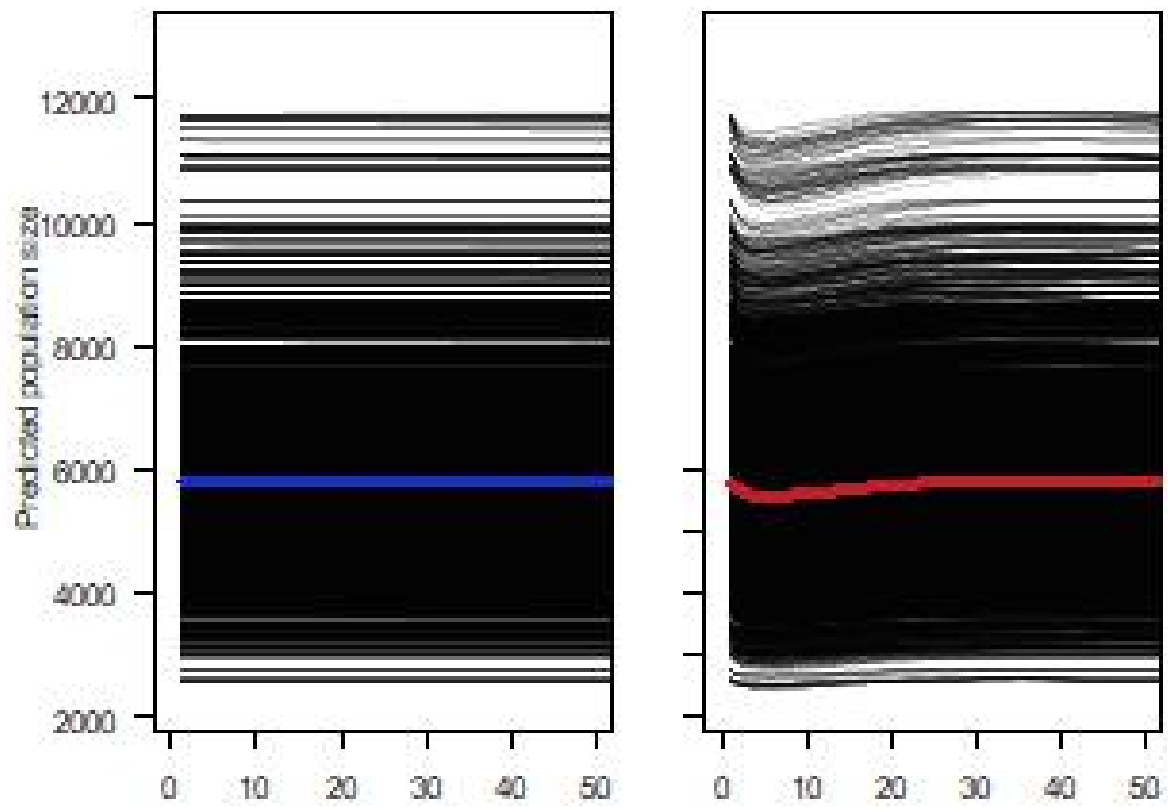
Mean for each of the injury metrics



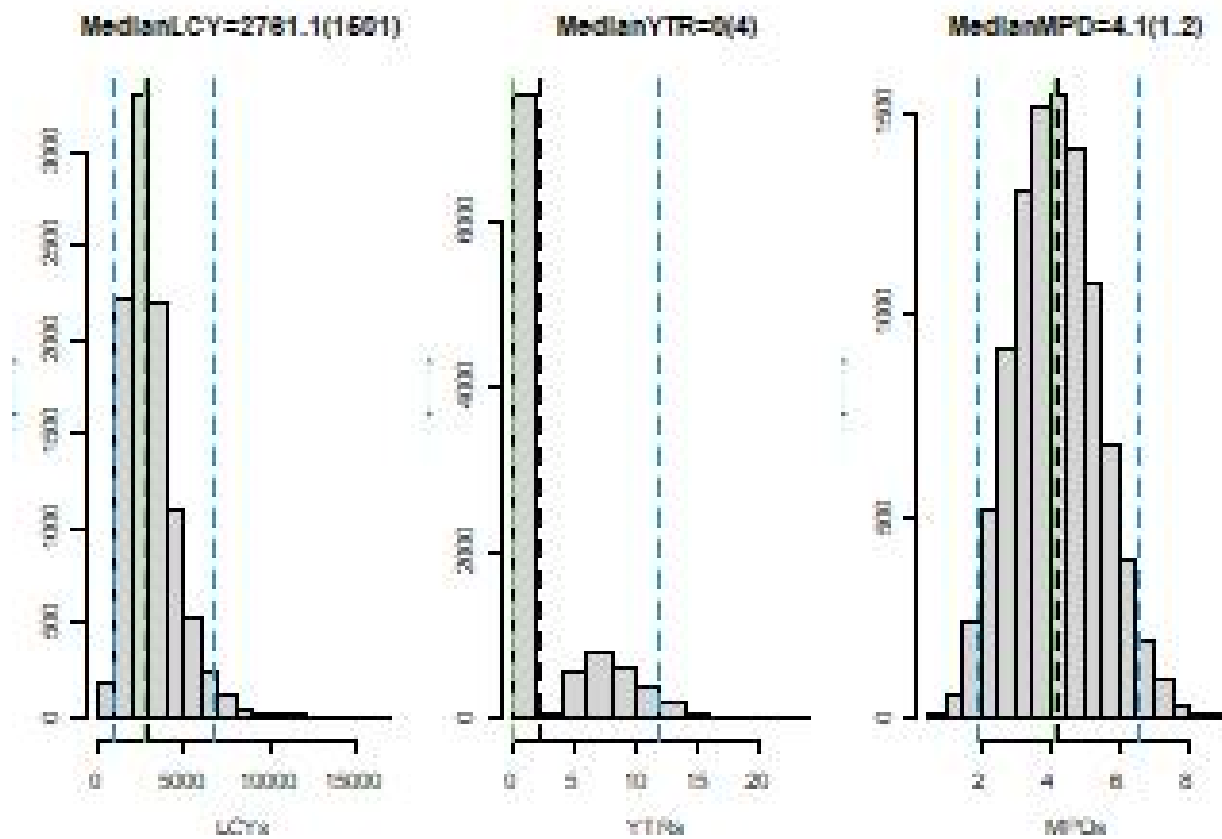
##	LCY	YTR	MPD
##	1901.525708	0.000000	1.801292

Peponocephala electra (Pele, melon-headed whale)

Population trajectories



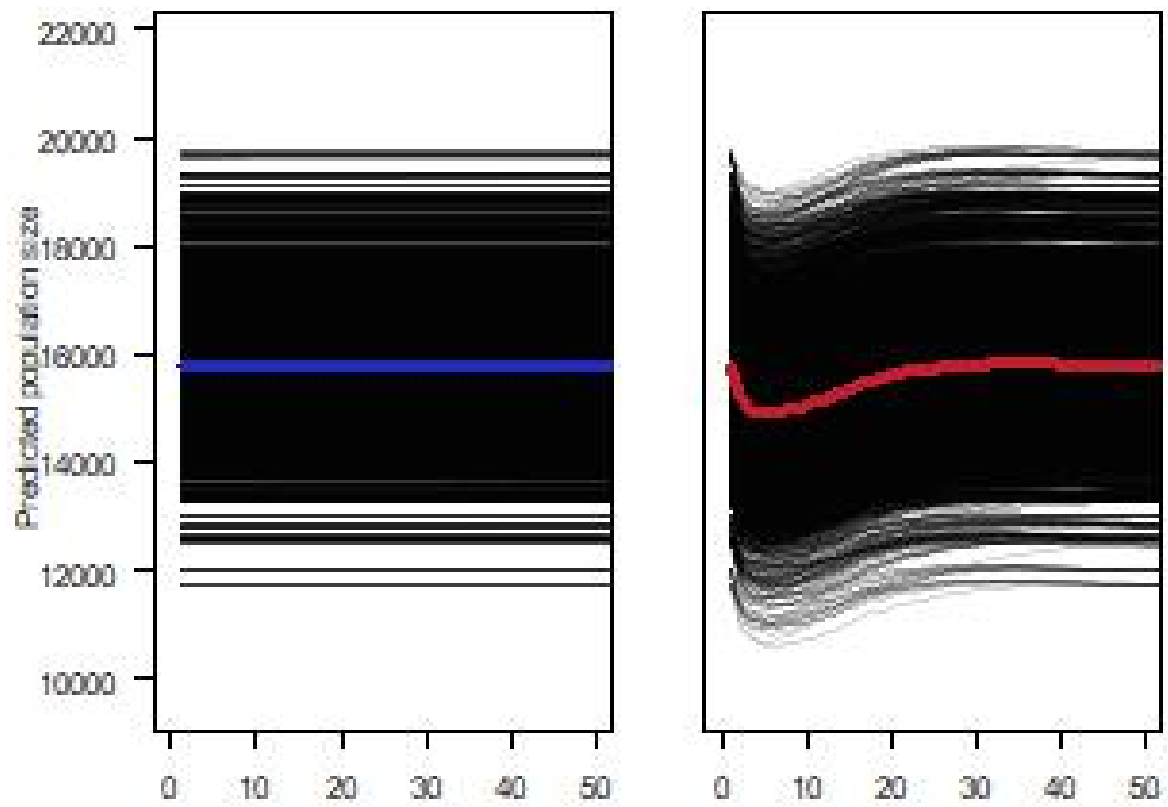
Mean for each of the injury metrics



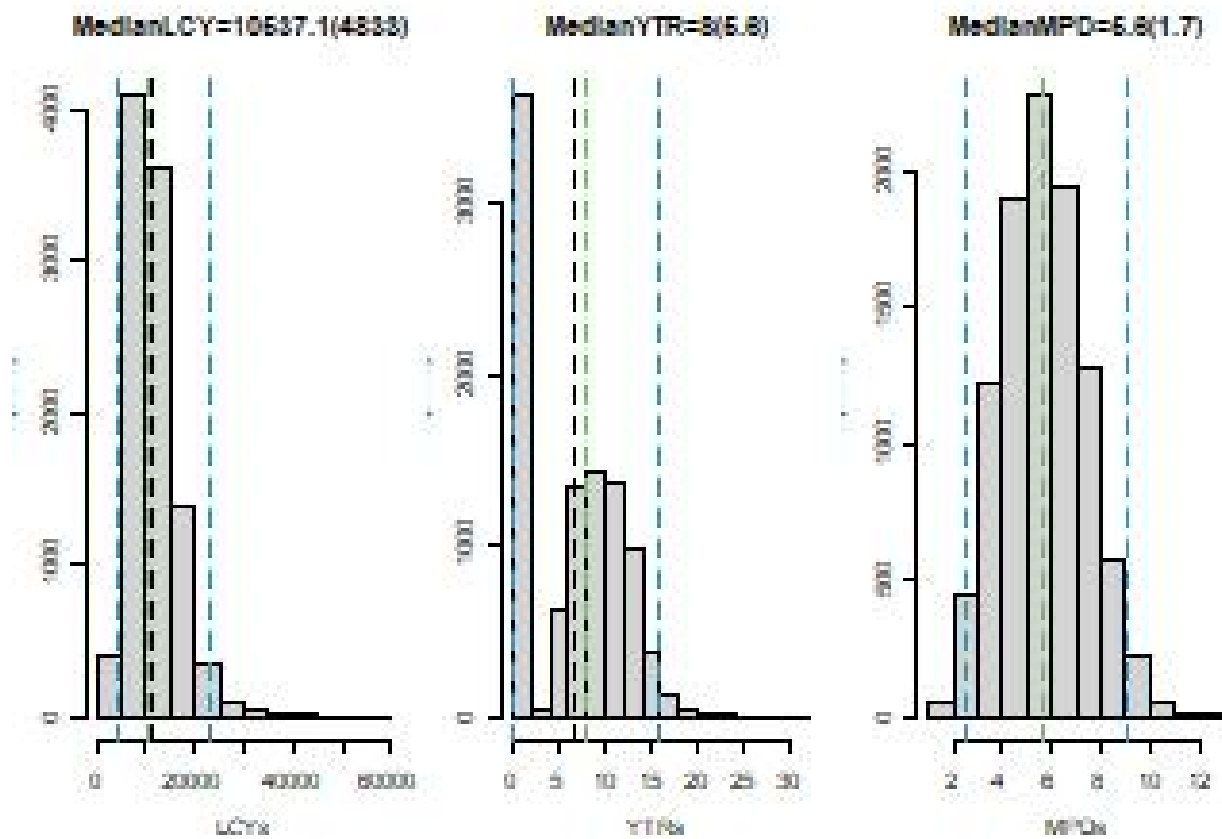
##	LCY	YTR	MPD
##	3071.523882	2.137900	4.168256

Tursiops truncatus (Ttro, offshore bottlenose dolphin)

Population trajectories



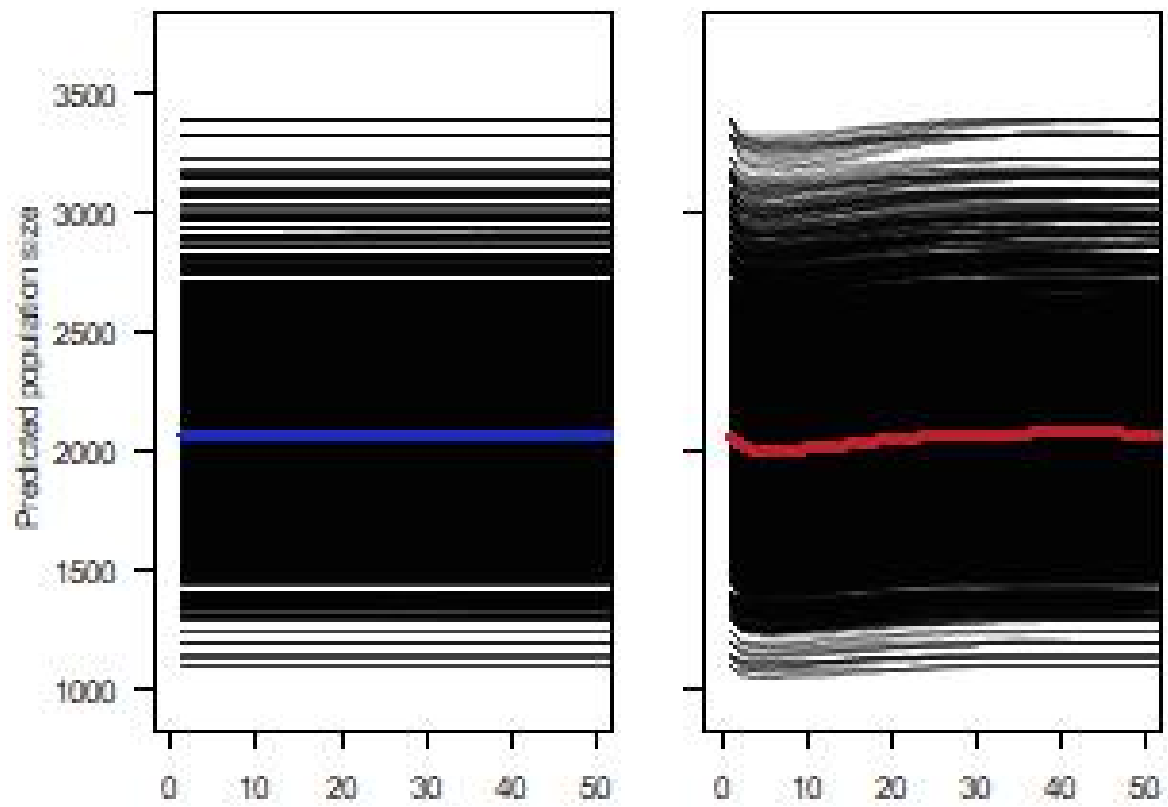
Mean for each of the injury metrics



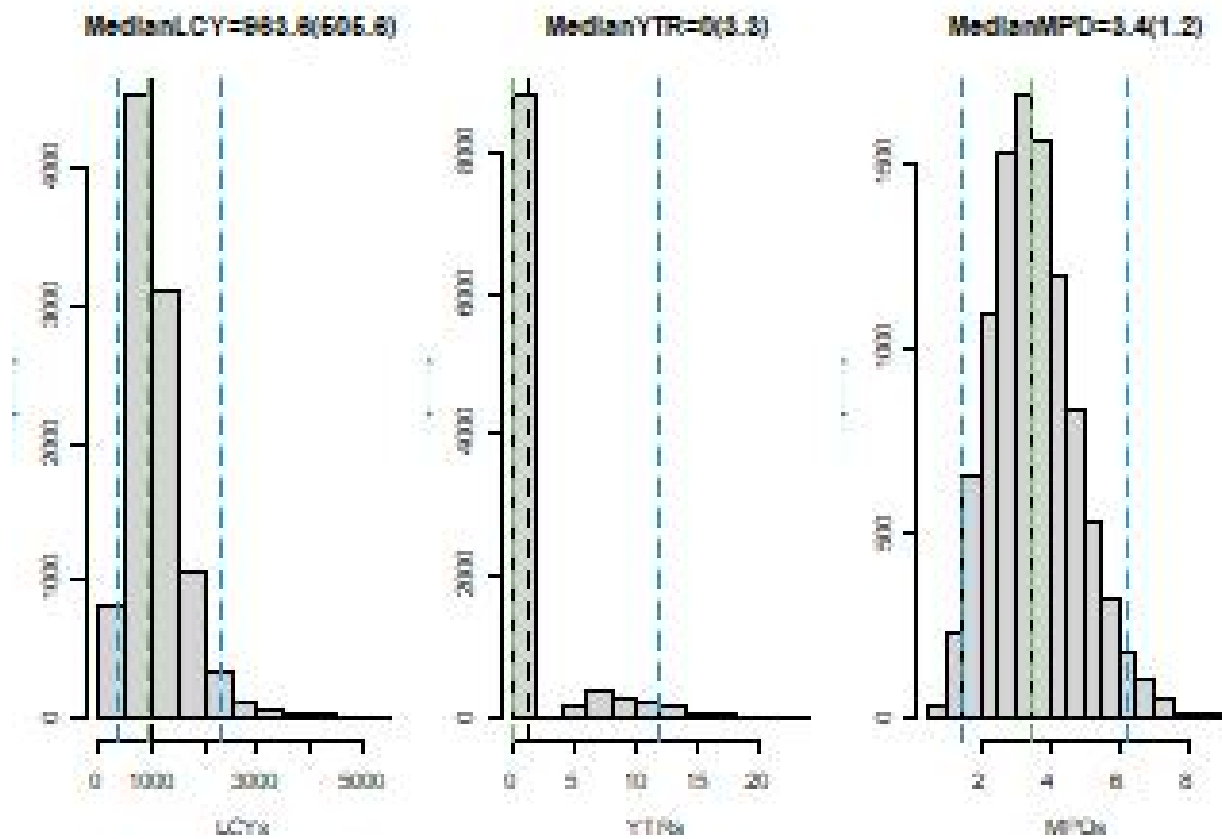
##	LCY	YTR	MPD
##	11369.564397	6.586900	5.658282

Globicephala macrorhynchus (Gmac, short finned pilot whale)

Population trajectories



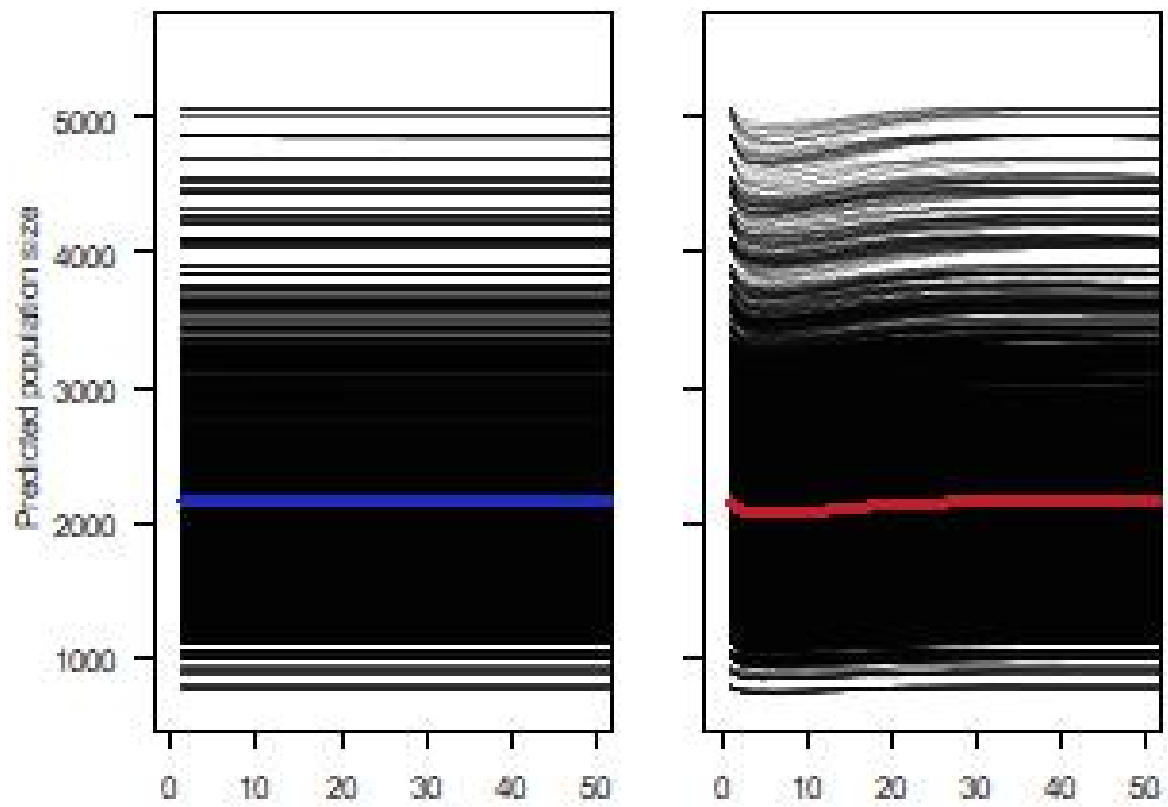
Mean for each of the injury metrics



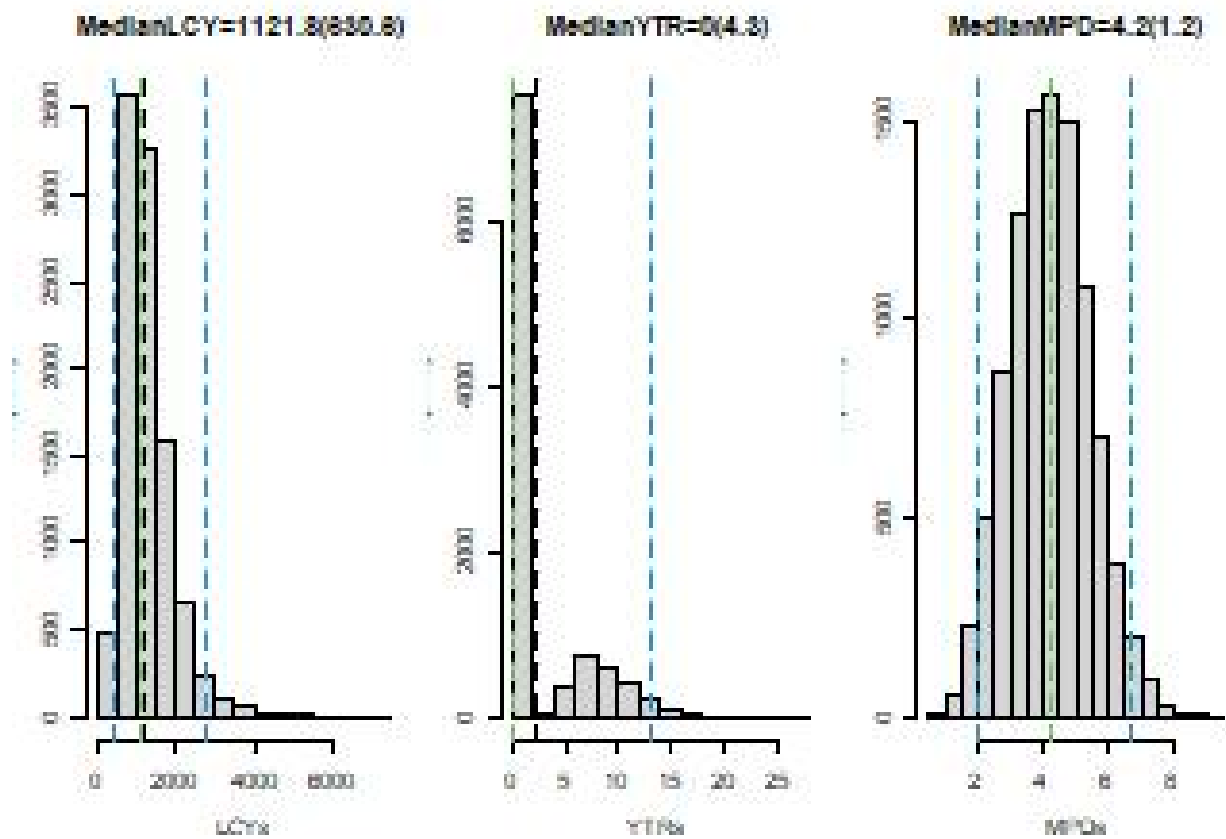
##	LCY	YTR	MPD
##	1057.863912	1.156400	3.538044

Feresa attenuata (Fatt, pygmy killer whale)

Population trajectories



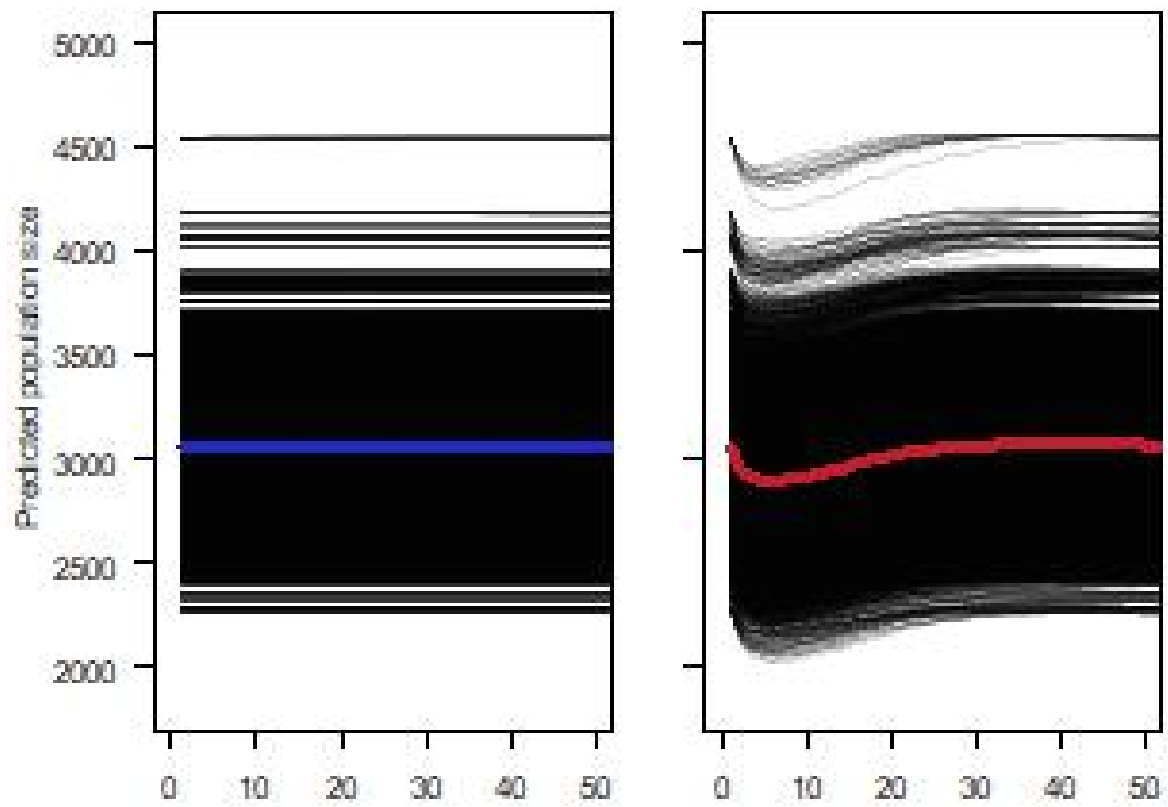
Mean for each of the injury metrics



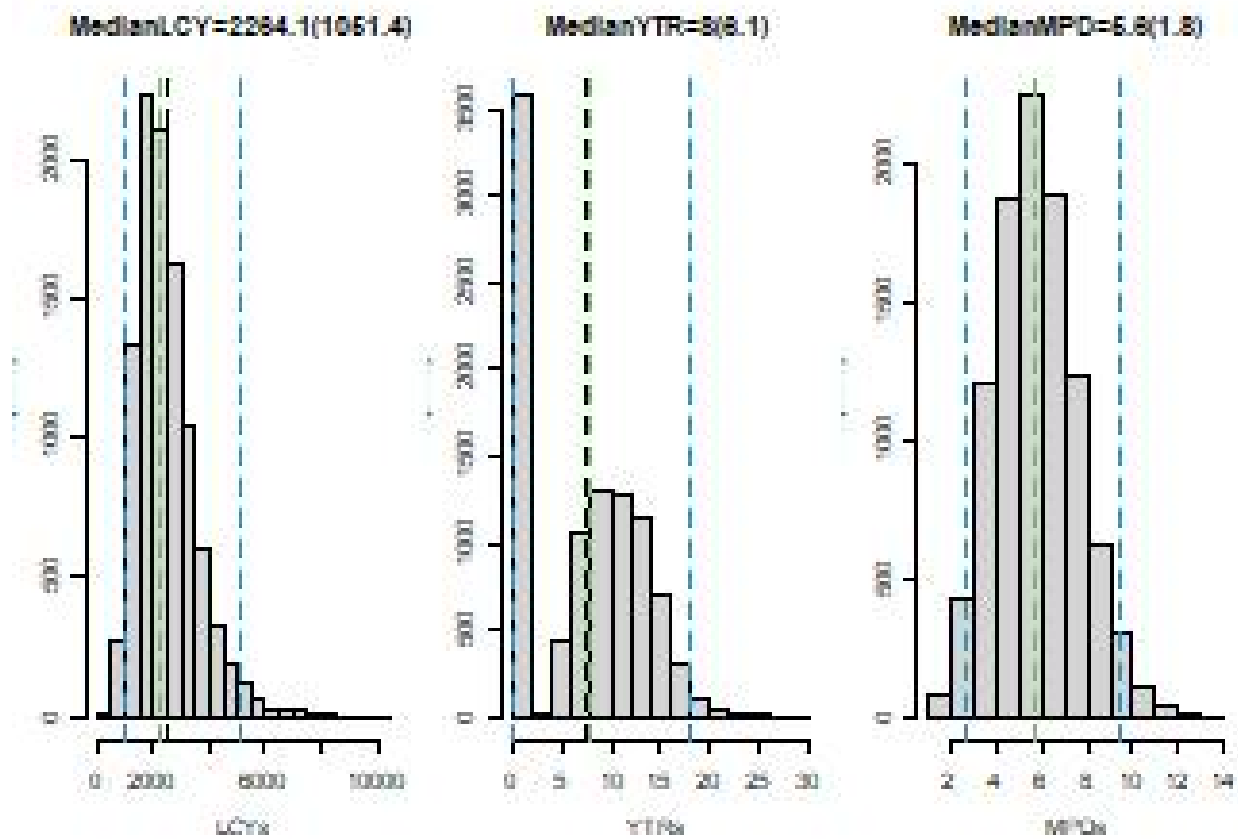
##	LCY	YTR	MPD
##	1249.503621	2.308200	4.194471

Grampus griseus (Ggri, Risso's dolphin)

Population trajectories



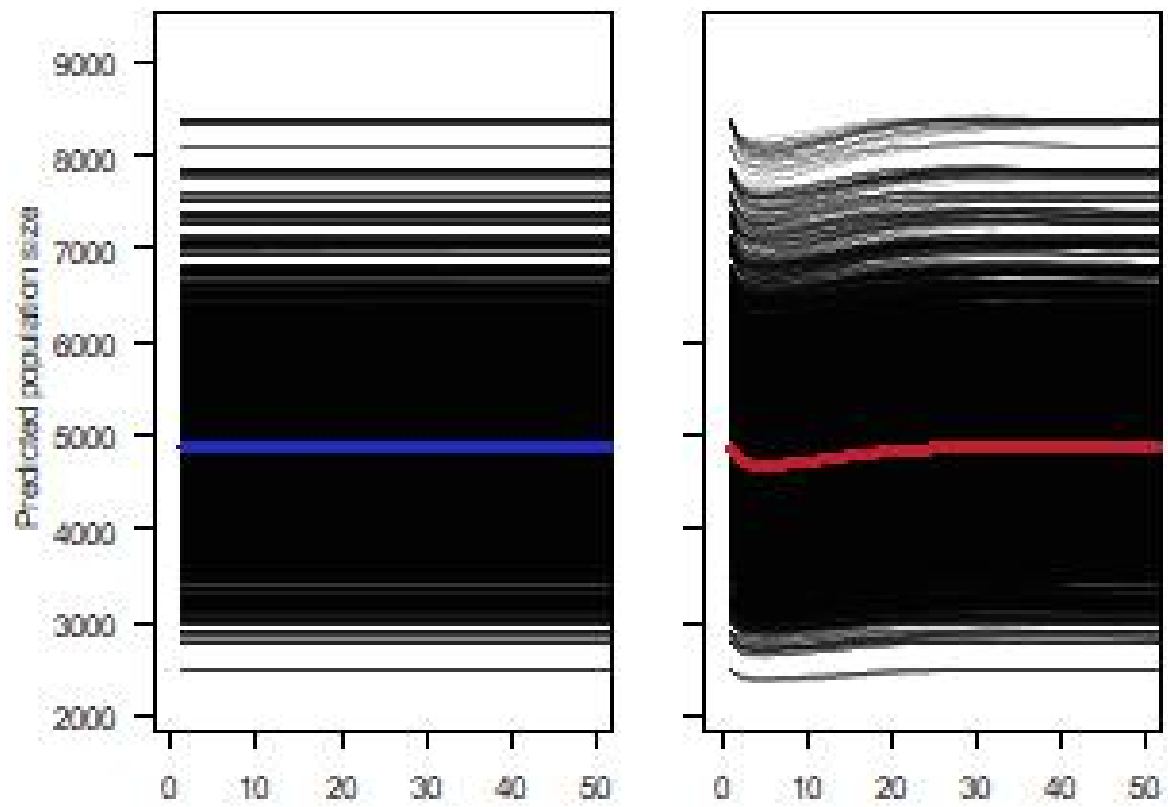
Mean for each of the injury metrics



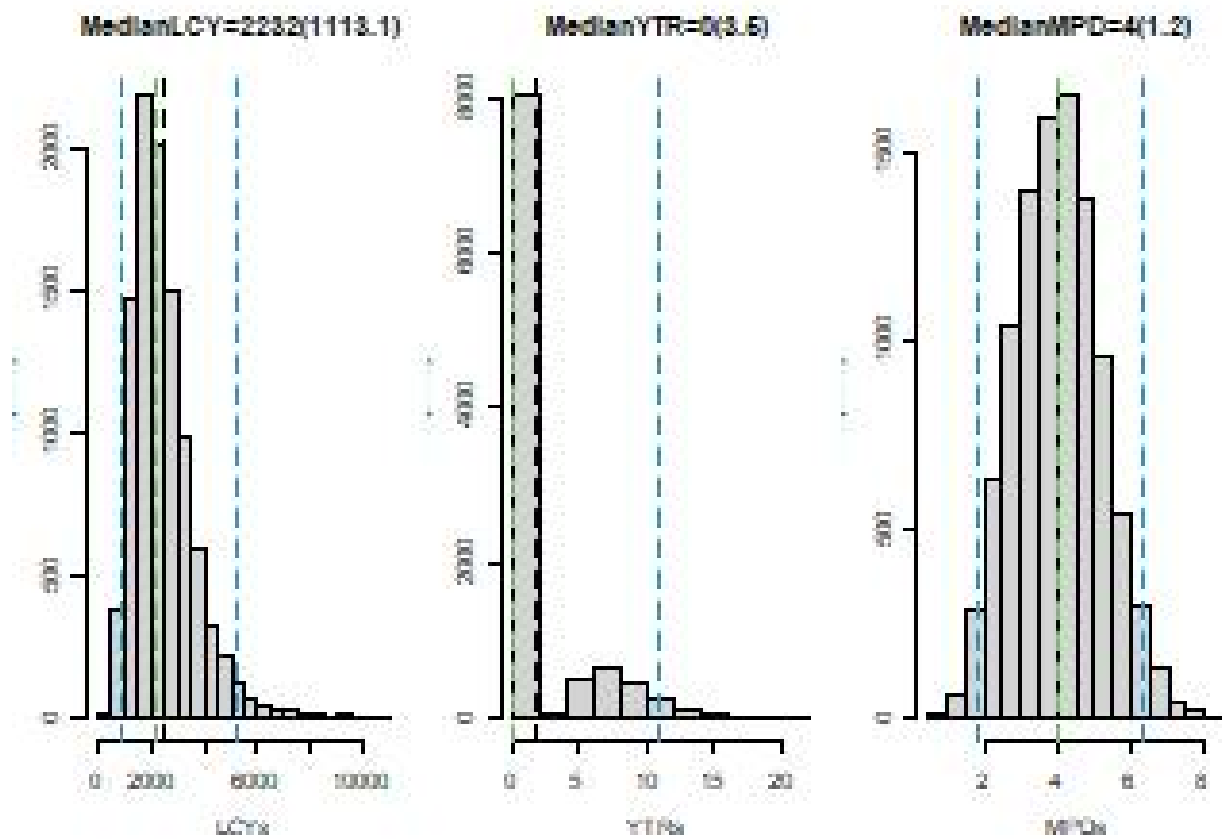
##	LCY	YTR	MPD
##	2449.136975	7.290900	5.735186

Steno bredanensis (Sbre, rough-toothed dolphin)

Population trajectories



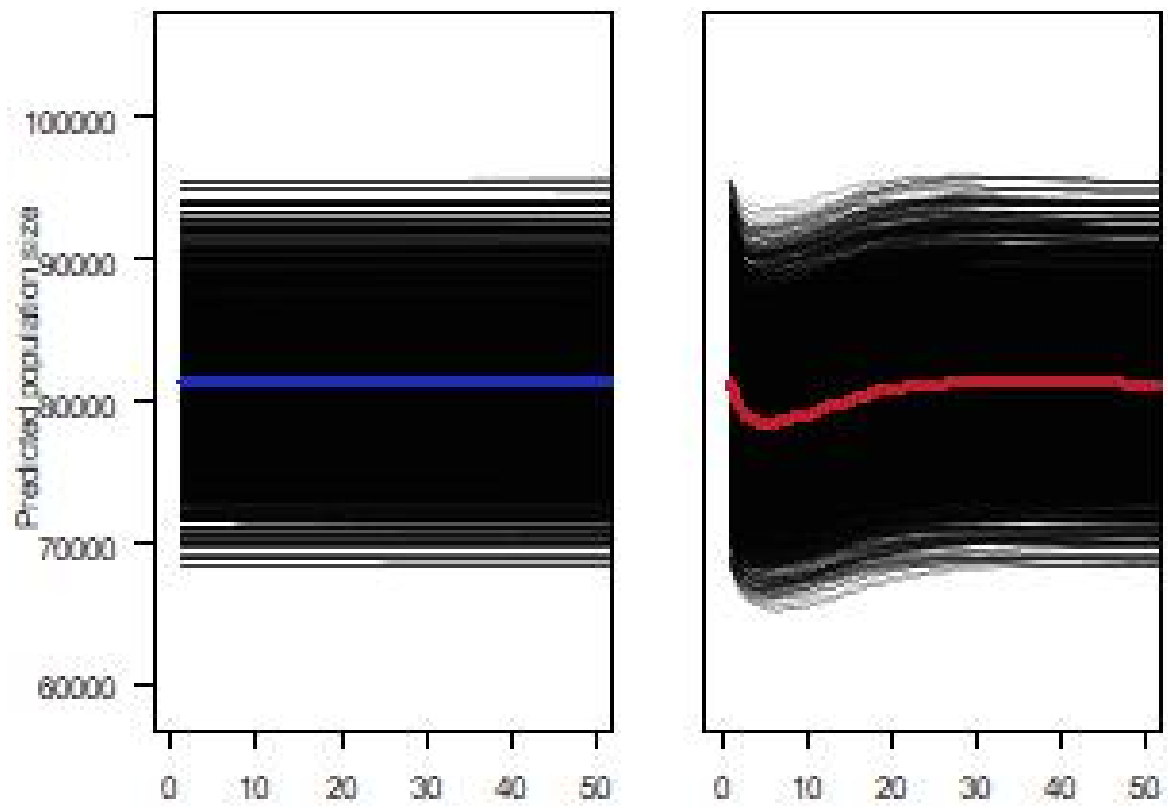
Mean for each of the injury metrics



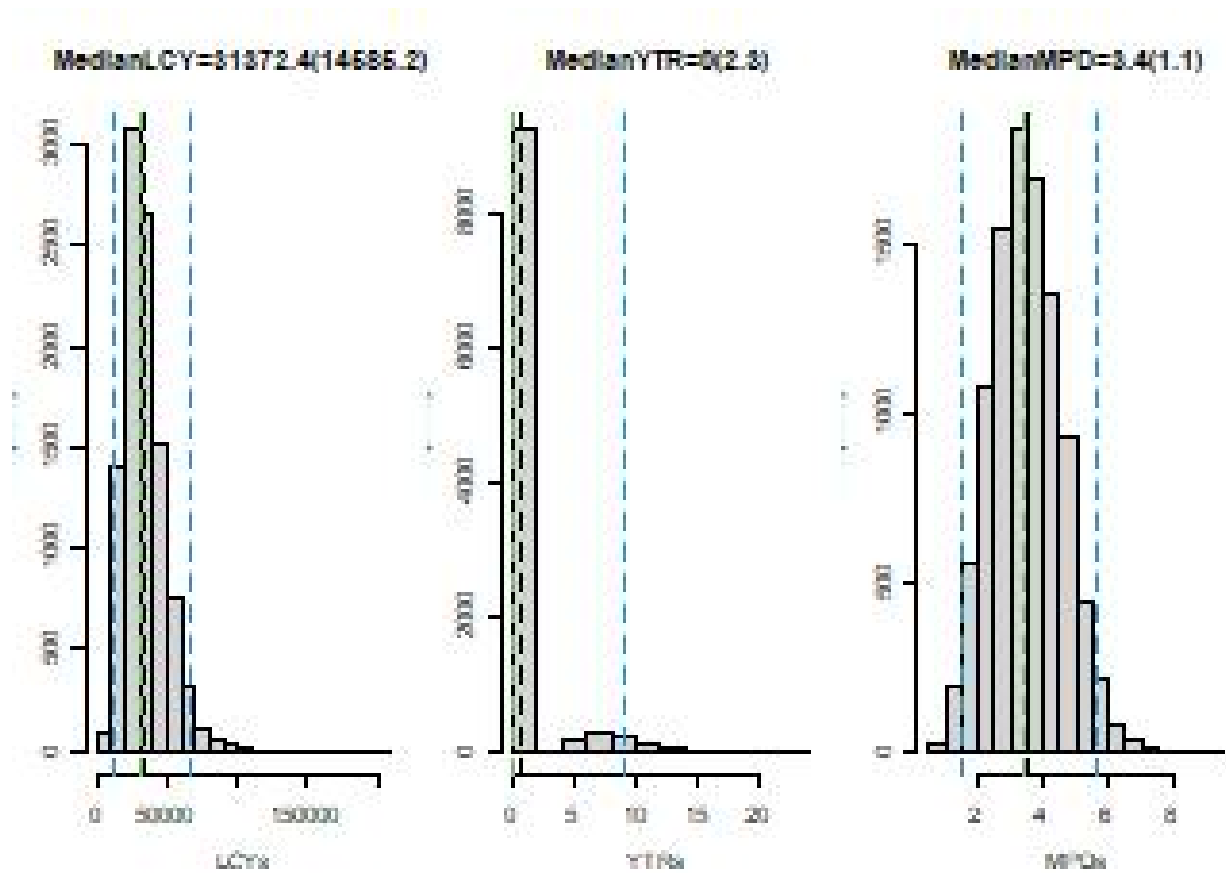
##	LCY	YTR	MPD
##	2434.417513	1.640900	4.006926

Stenella attenuata (Satt, pantropical spotted dolphin)

Population trajectories



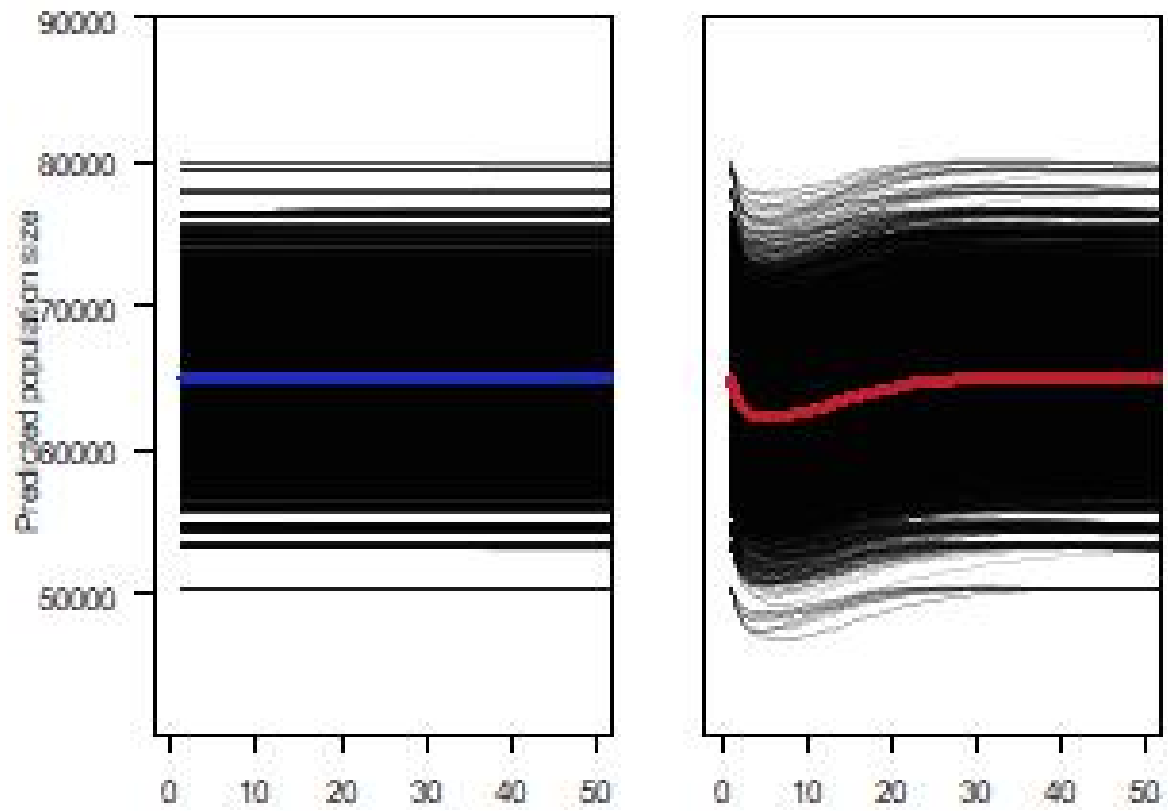
Mean for each of the injury metrics



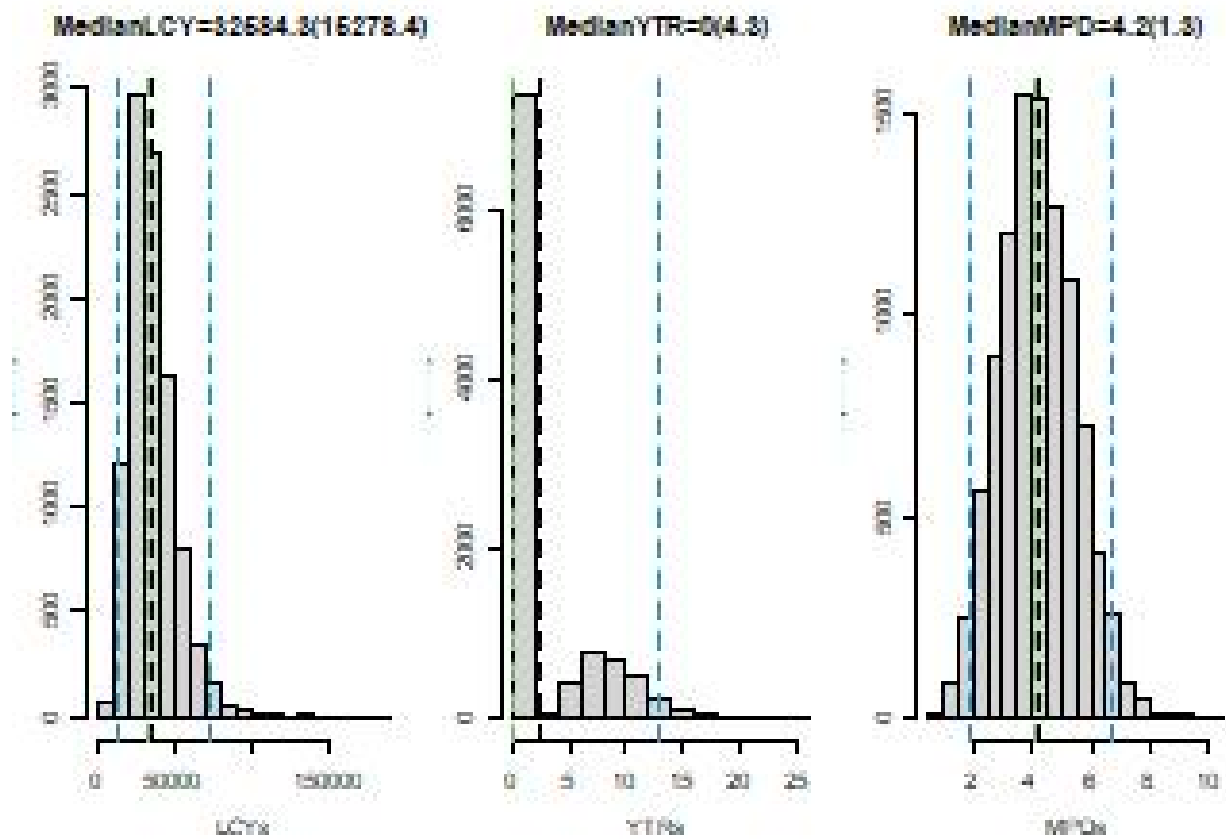
##	LCY	YTR	MPD
##	33809.632317	0.635700	3.474522

Tursiops truncatus (Ttrs, shelf bottlenose dolphin)

Population trajectories



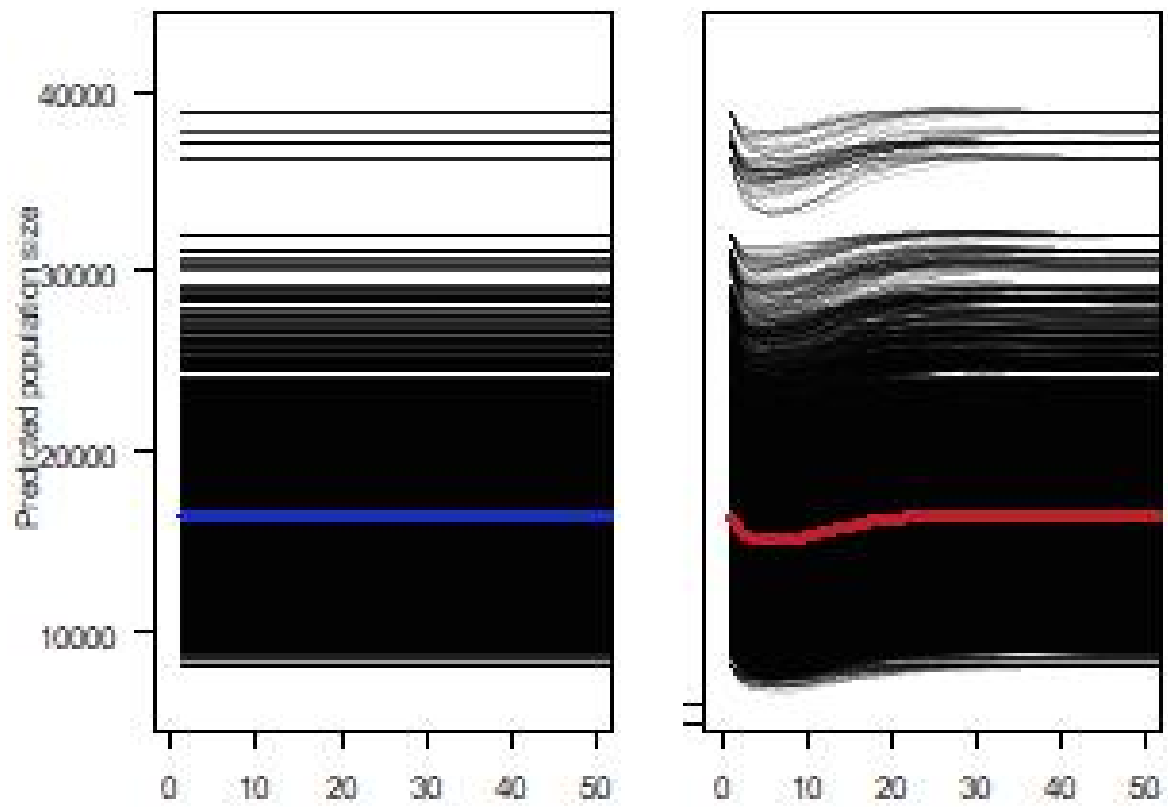
Mean for each of the injury metrics



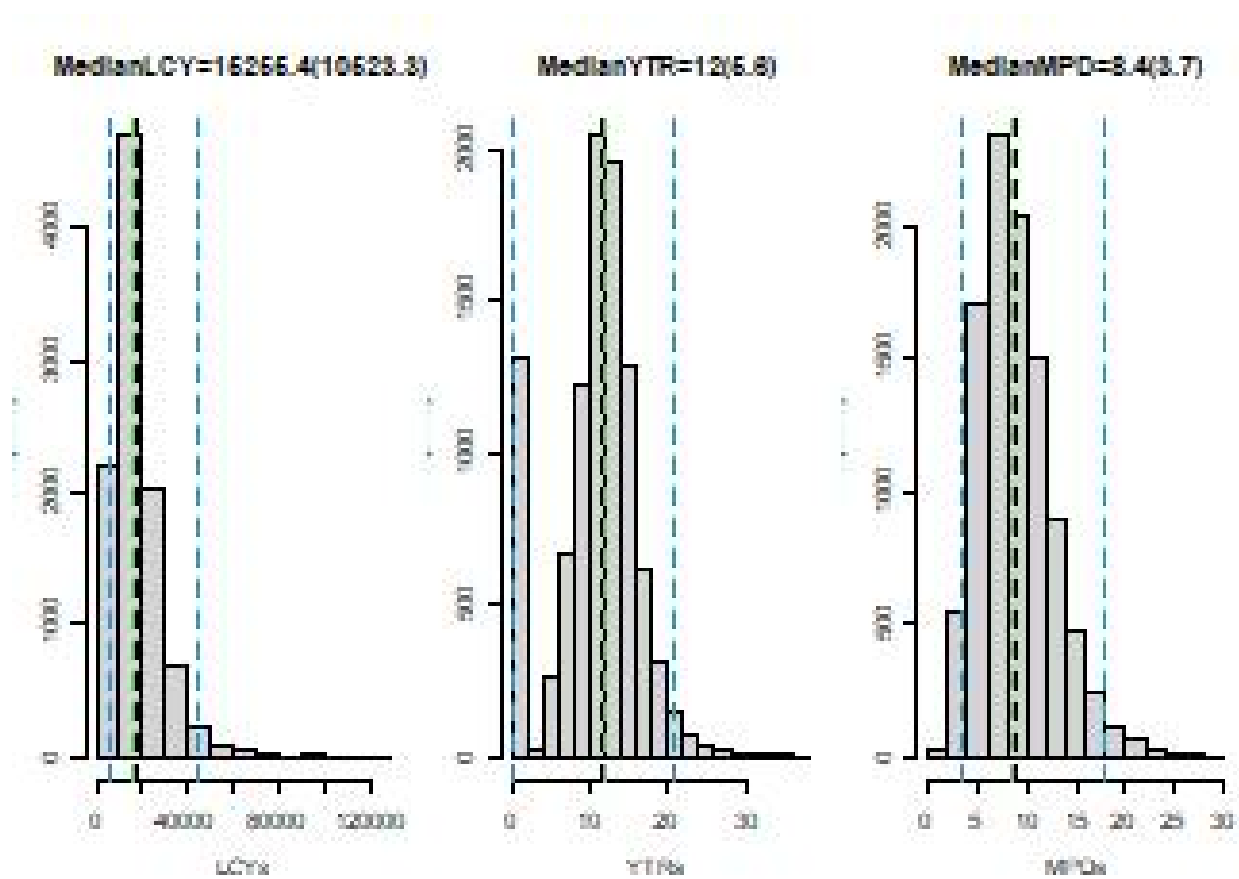
##	LCY	YTR	MPD
##	35190.477524	2.435900	4.191836

Stenella longirostris (Slon, spinner dolphin)

Population trajectories



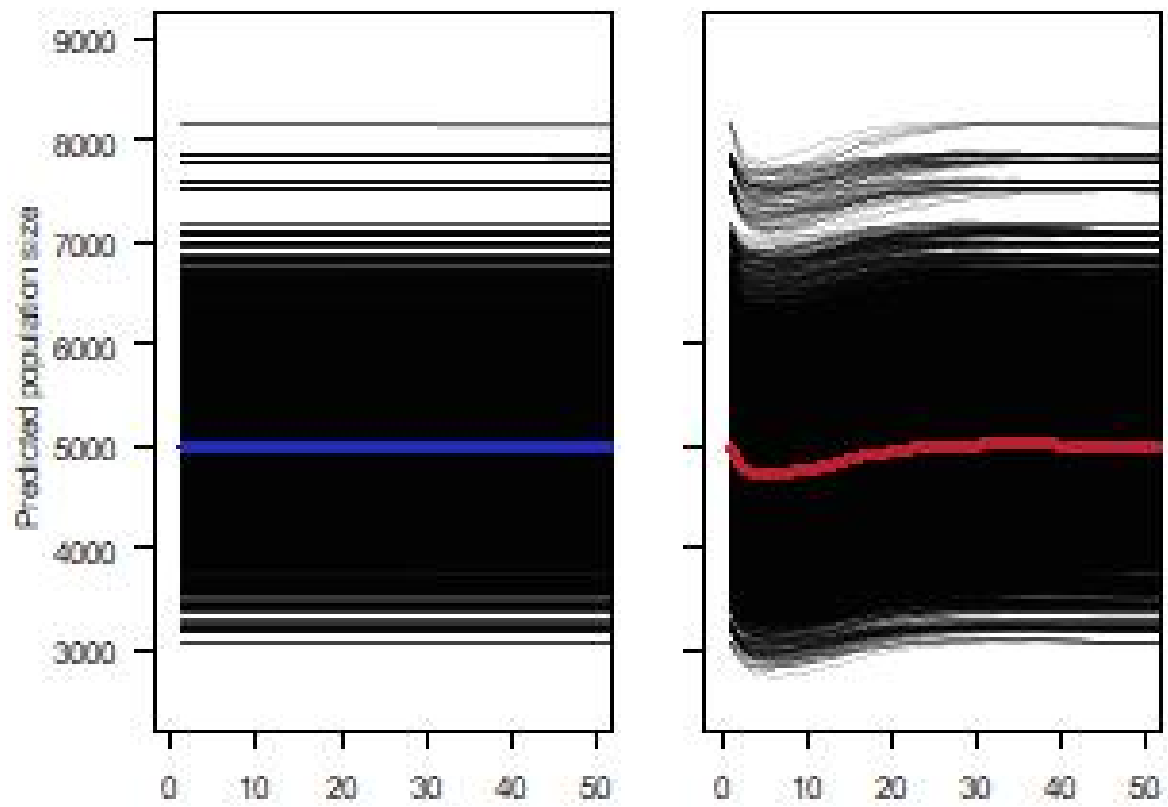
Mean for each of the injury metrics



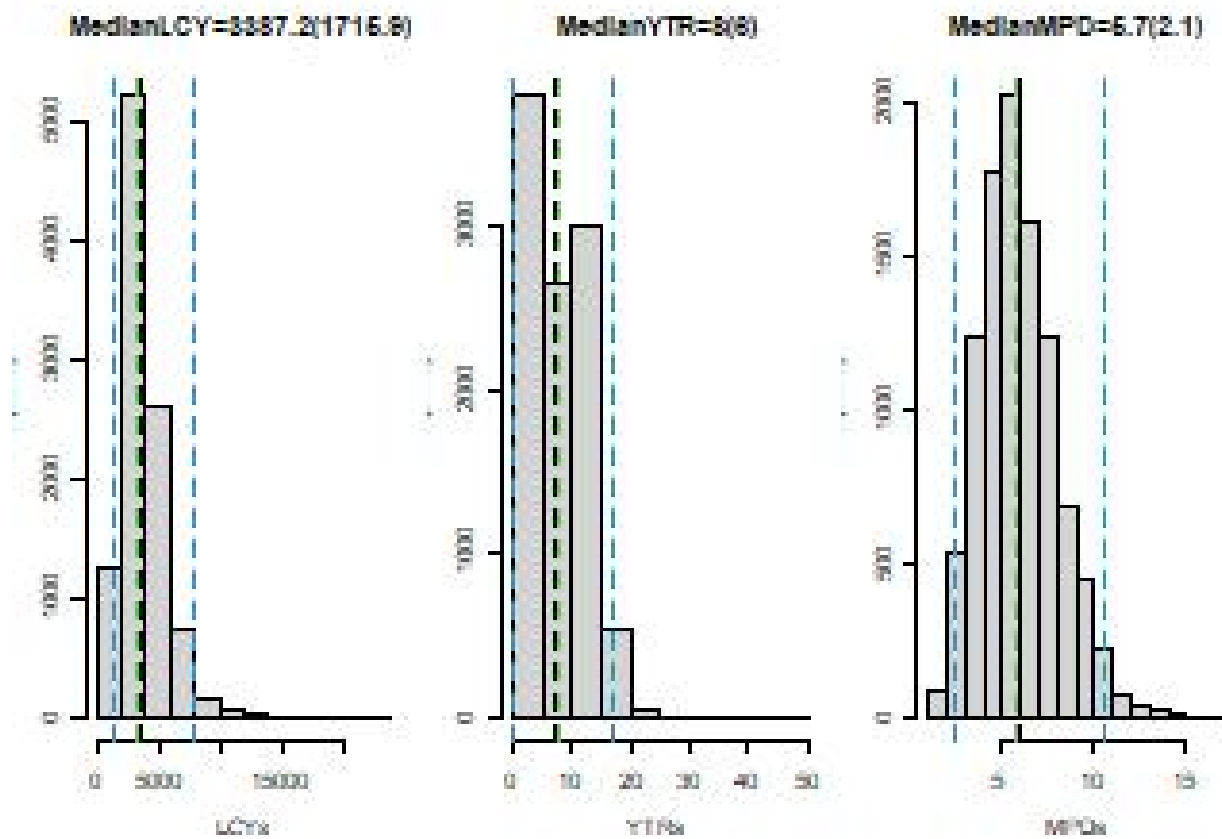
##	LCY	YTR	MPD
##	17714.767745	11.196700	8.899829

Stenella coeruleoalba (Scoe, striped dolphin)

Population trajectories



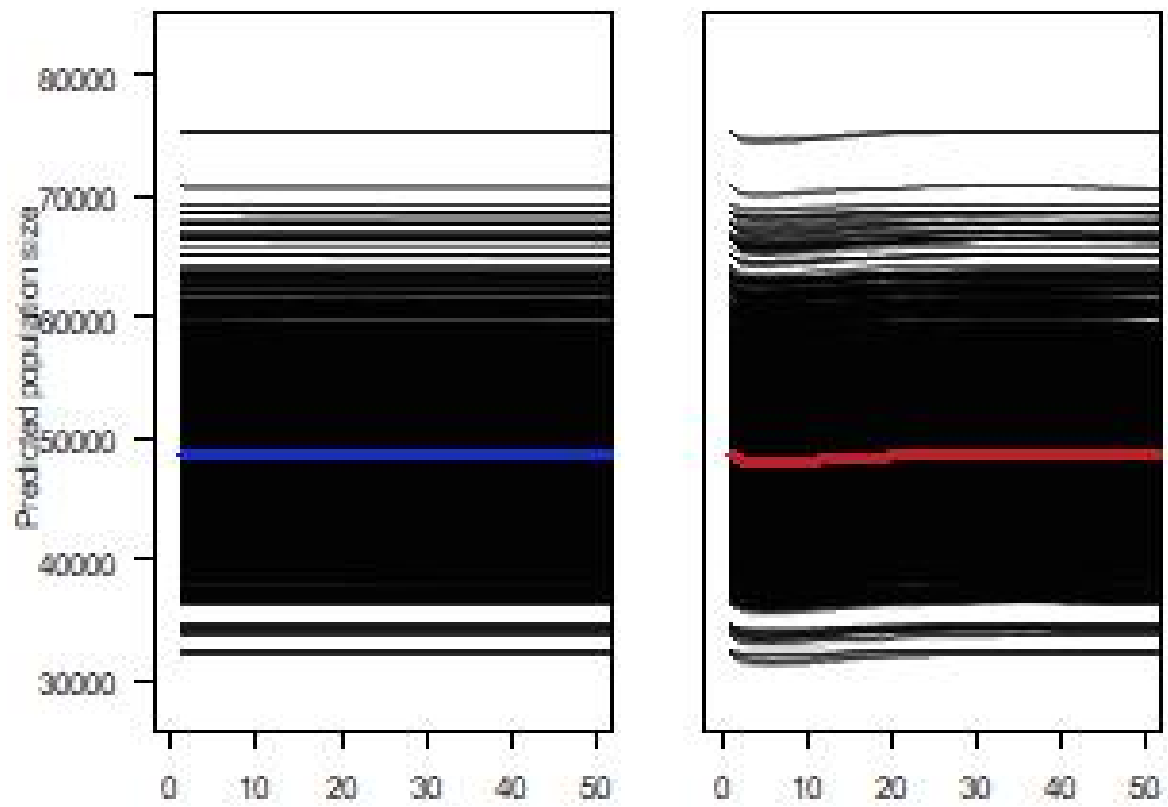
Mean for each of the injury metrics



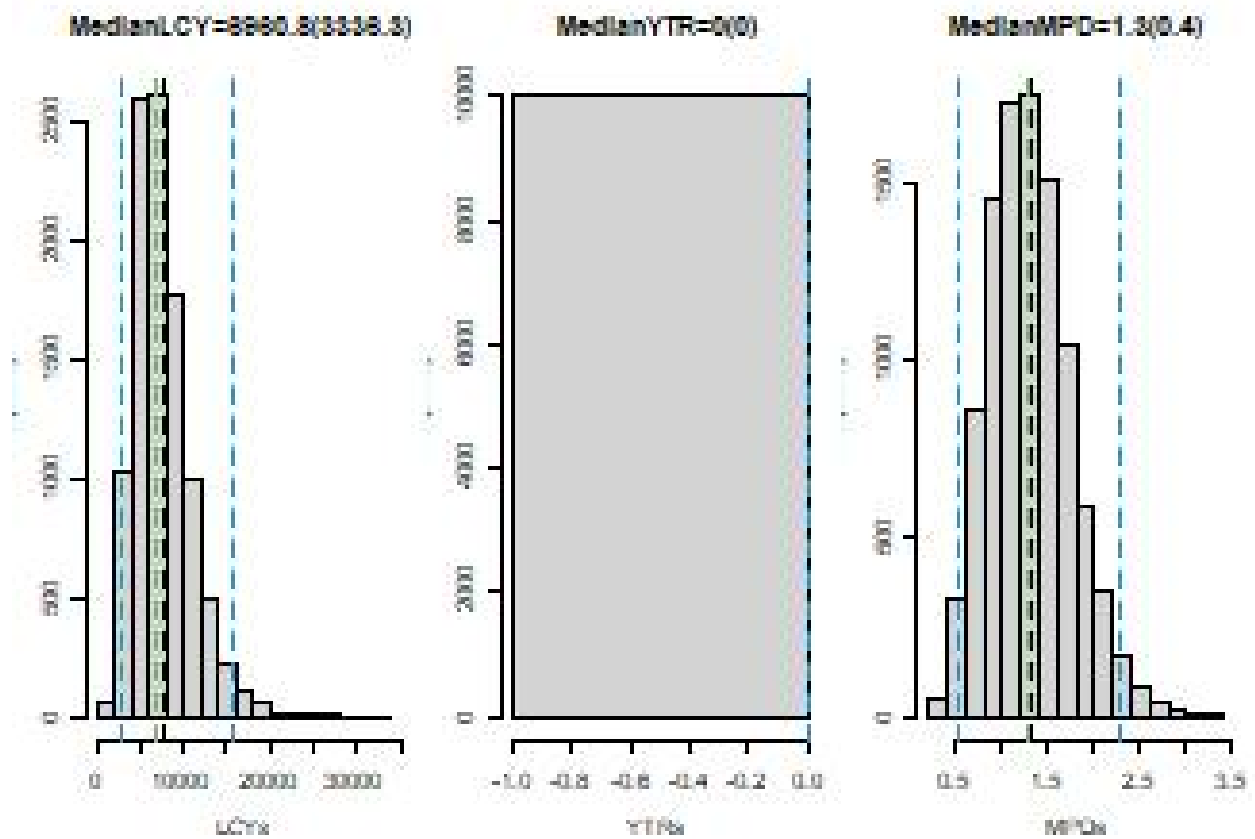
##	LCY	YTR	MPD
##	3715.207476	7.108100	5.883191

Stenella frontalis (Sfro, Atlantic spotted dolphin)

Population trajectories



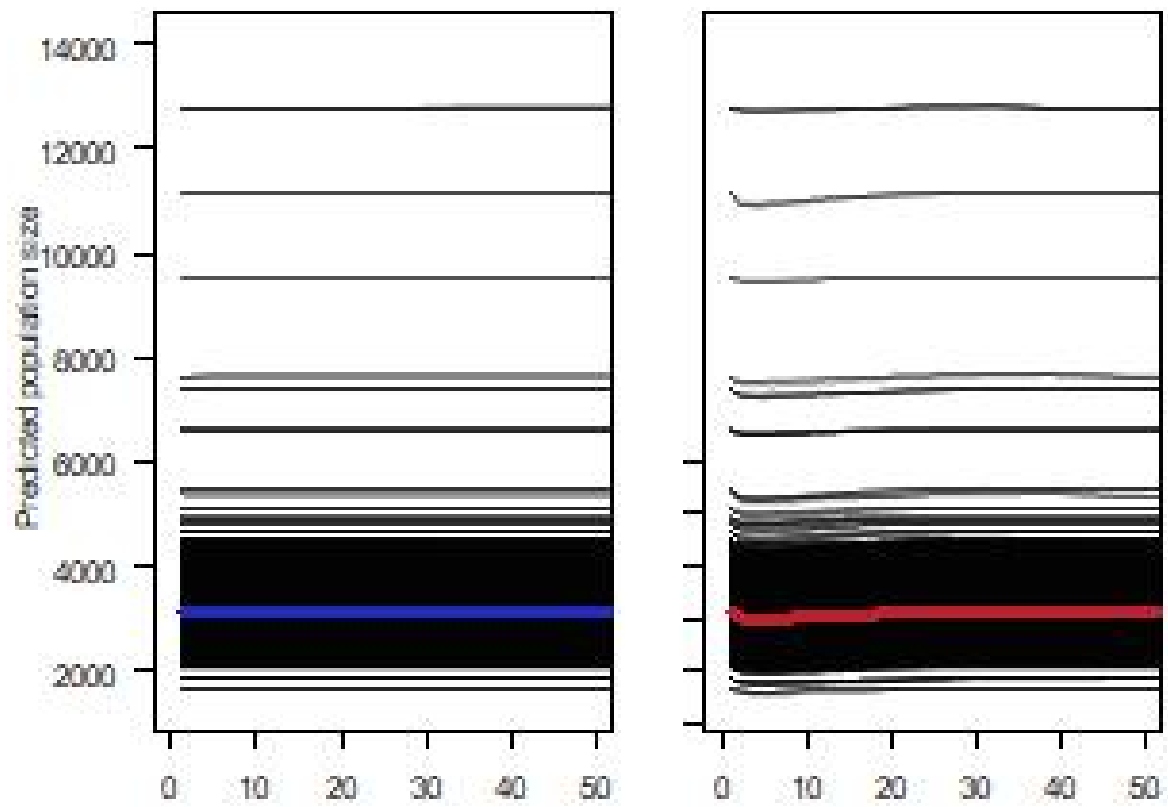
Mean for each of the injury metrics



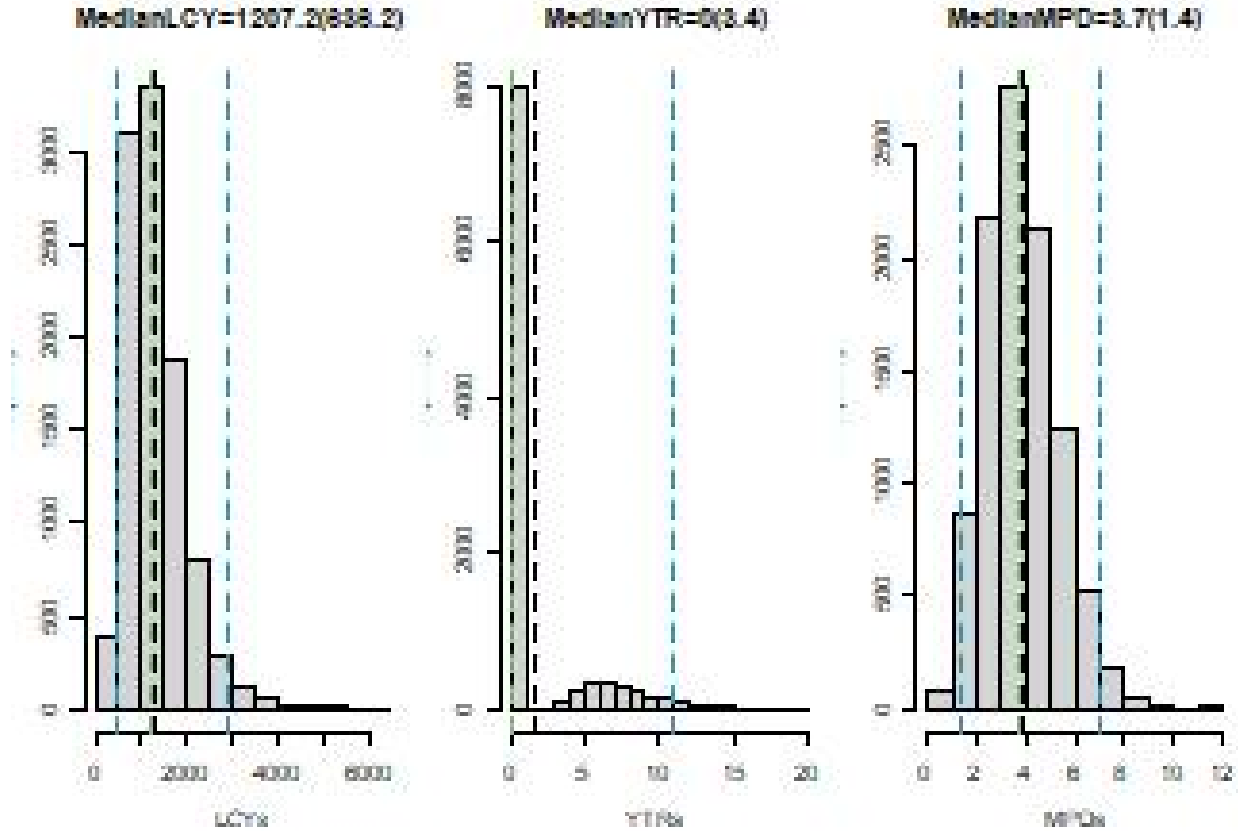
##	LCY	YTR	MPD
##	7509.270139	0.000000	1.296869

Beaked whales spp (Bwsp)

Population trajectories



Mean for each of the injury metrics



```
##          LCY          YTR          MPD
## 1320.899319    1.592200    3.816632
```

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

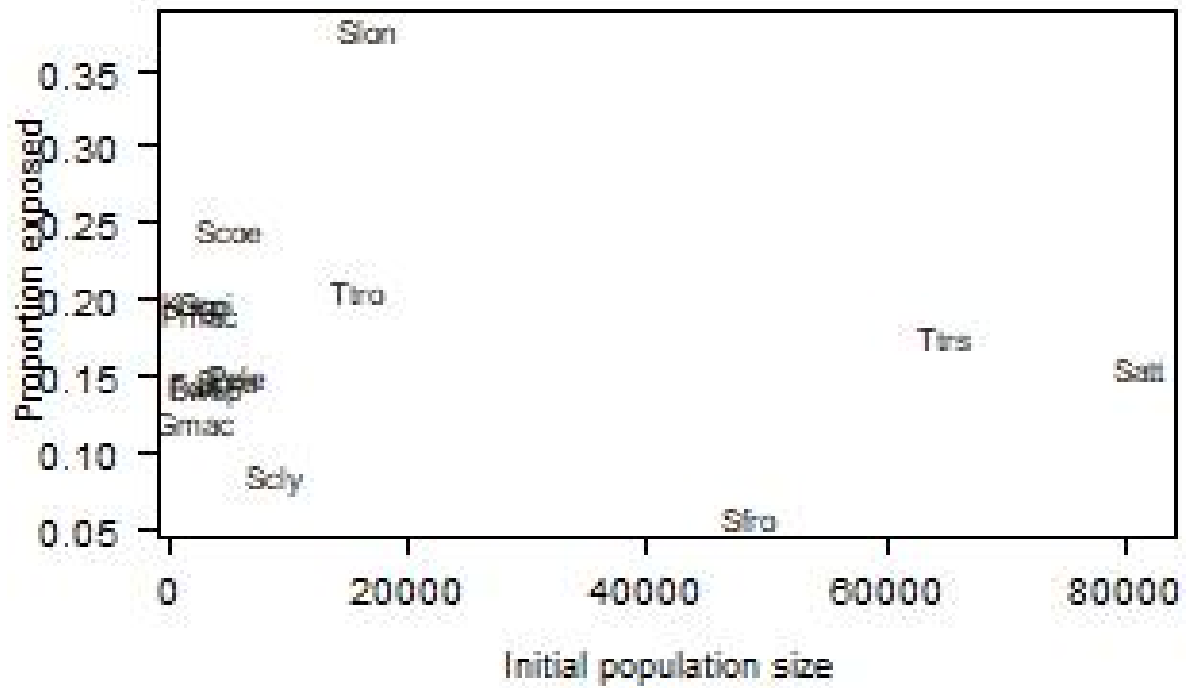
The population size and the proportion exposed are *a priori* expected to be key determinants of injury, since the population size will have a direct impact on LCY and the proportion exposed should have a direct impact on all 3 measures.

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

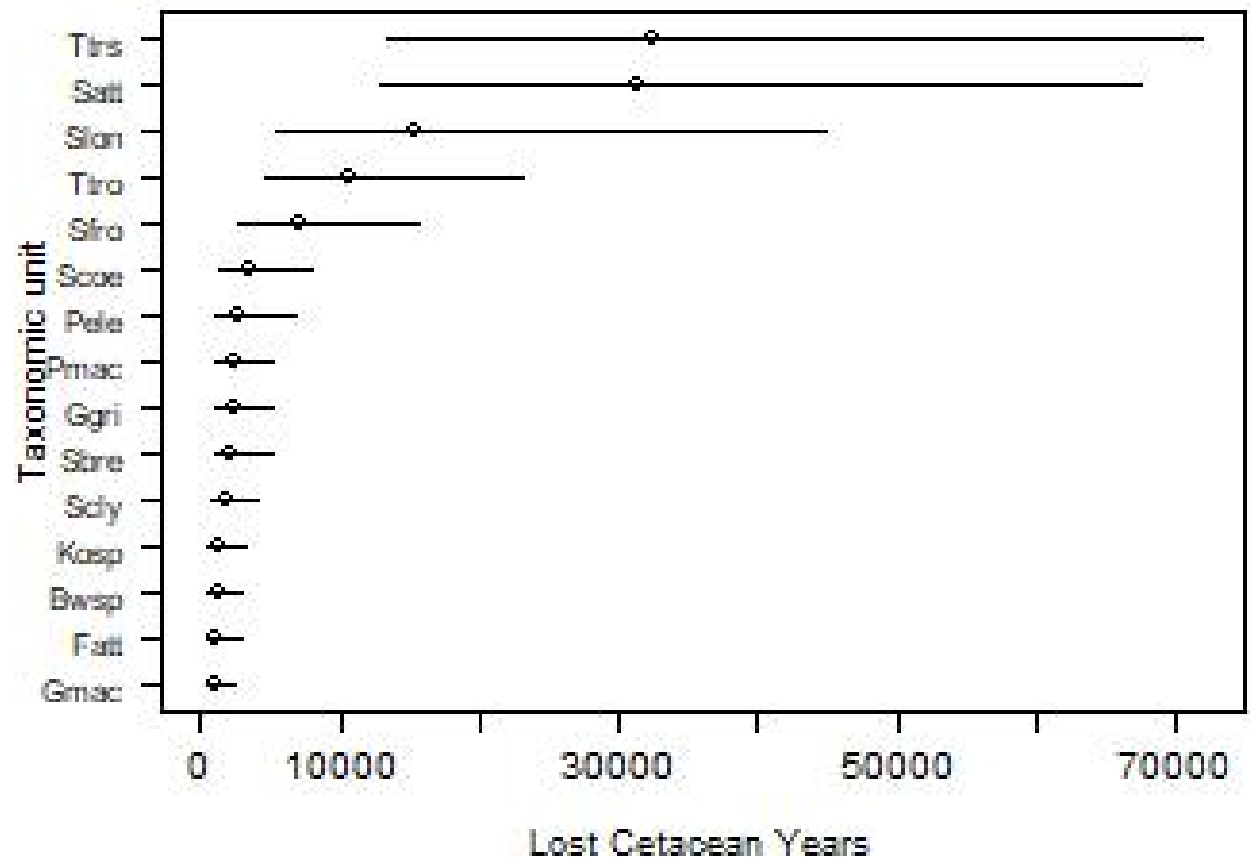
In the following table we present the medians and the 95% confidence intervals for our current estimates of injury. These correspond to the results in table 3 in the main paper.

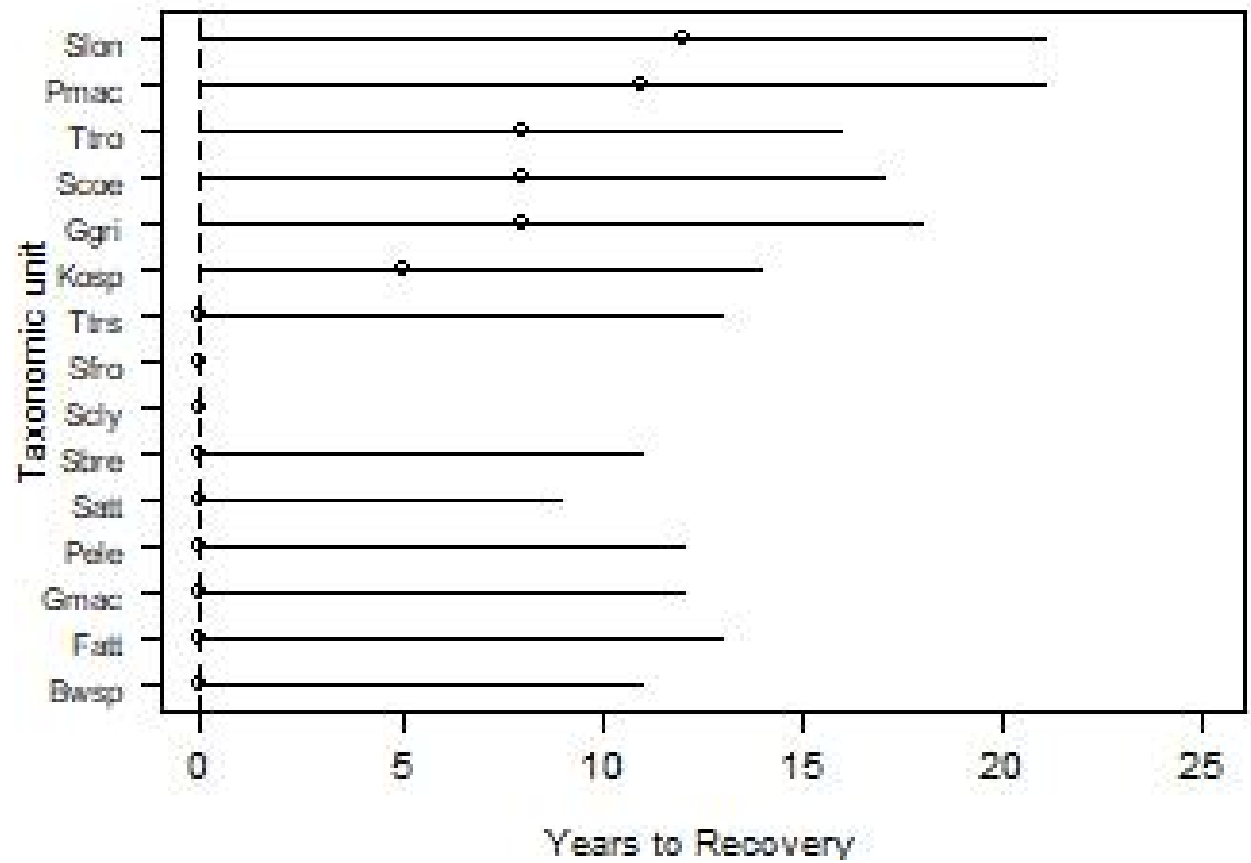
Table 3: Median injury results for the different stocks considered and respective 95% confidence intervals by the percentile method

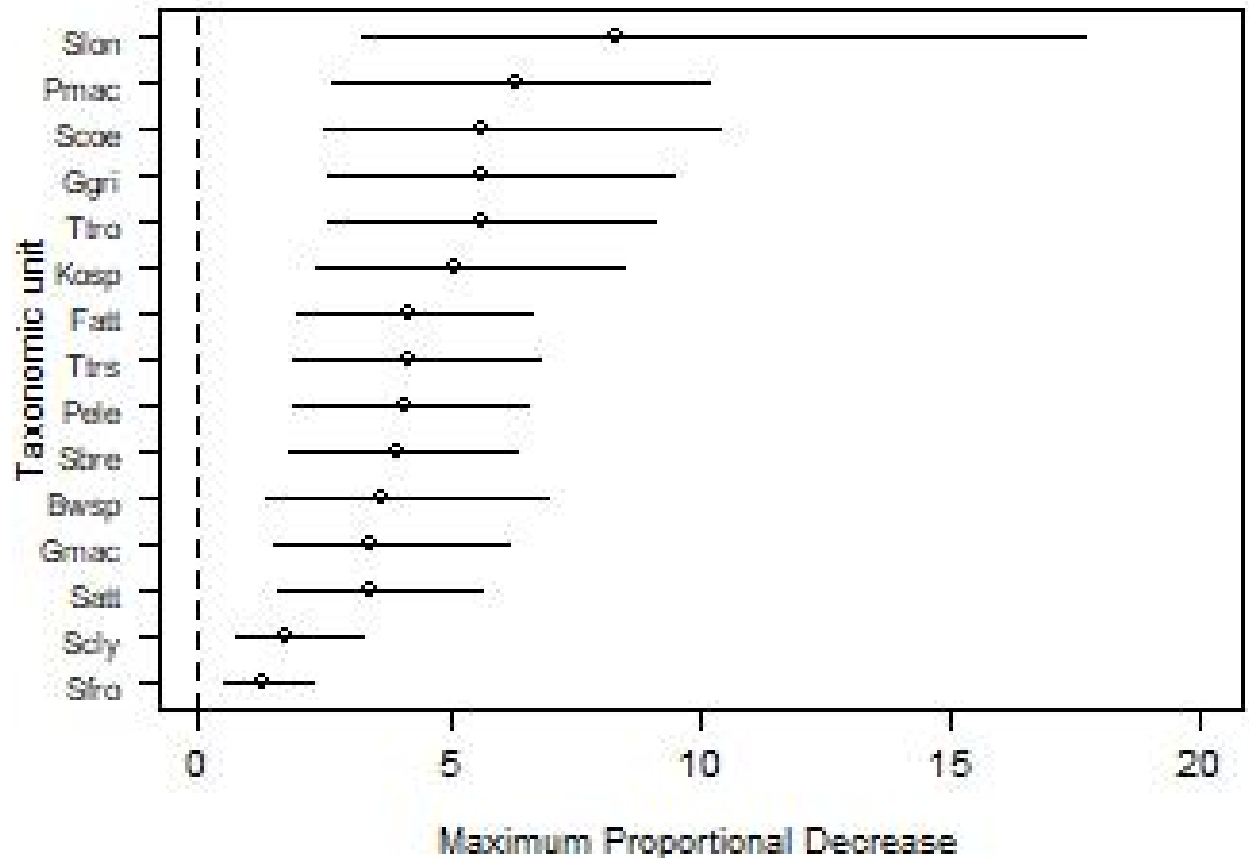
Sp	LCYmed	LCYl	LCYu	MPDmed	MPDl	MPDu	YTRmed	YTRl	YTRu
Bwsp	1207	445	2899	3.683	1.386	6.951	0	0	11
Fatt	1122	437	2809	4.186	1.939	6.642	0	0	13
Ggri	2264	978	5049	5.623	2.569	9.494	8	0	18
Gmac	964	385	2291	3.435	1.486	6.216	0	0	12
Kosp	1294	519	3112	5.114	2.315	8.492	5	0	14
Pele	2761	1081	6767	4.140	1.925	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 generalized linear models to model the mean value of the injury per species as a function of the mean values of the input parameters. Only the following four parameters seem to have a considerable potential relevant effect on the injury metrics:

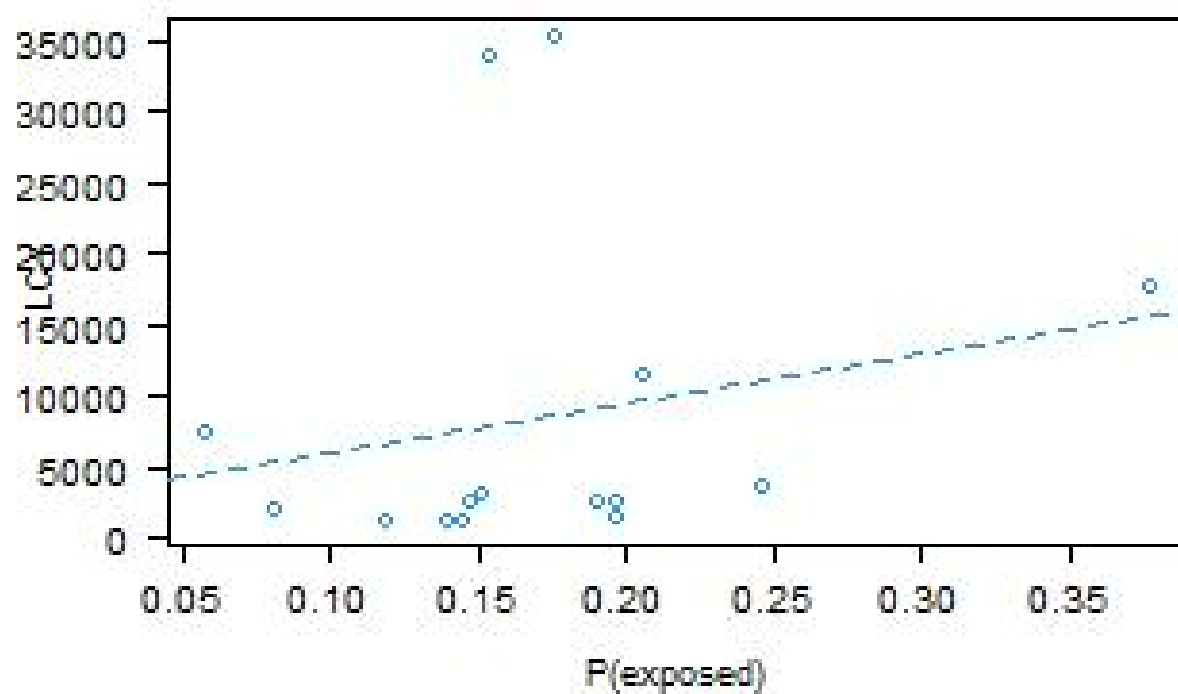
- Proportion Exposed
- Initial population size
- Gestation duration
- Survival reduction

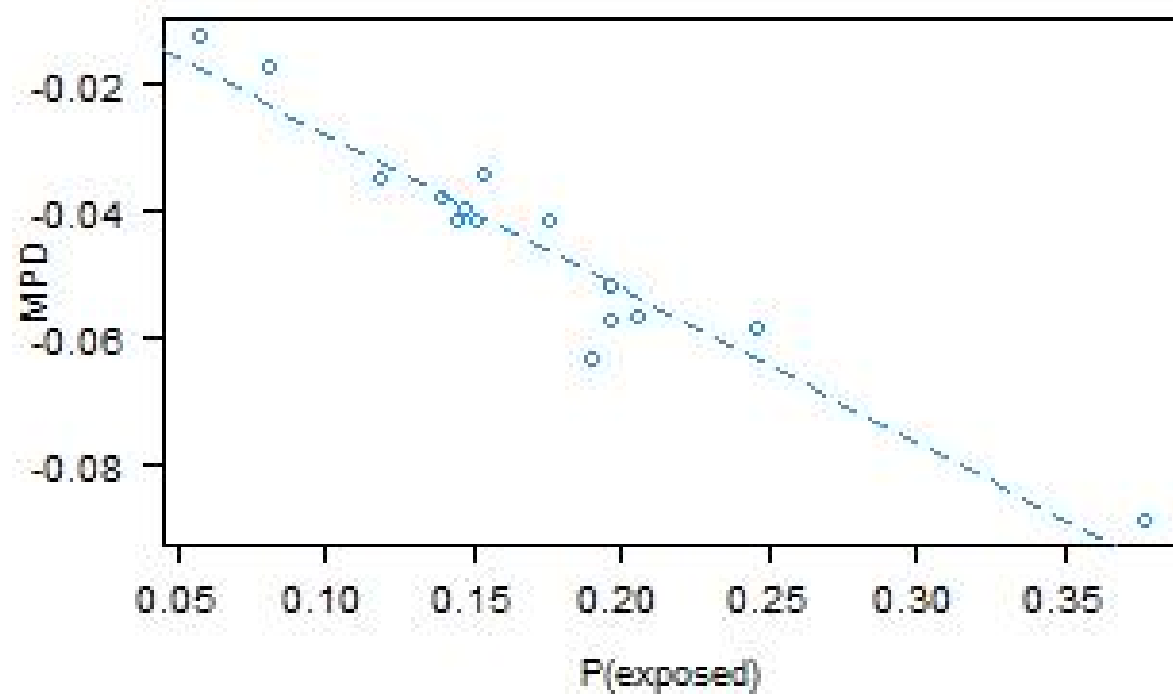
Exploratory analysis

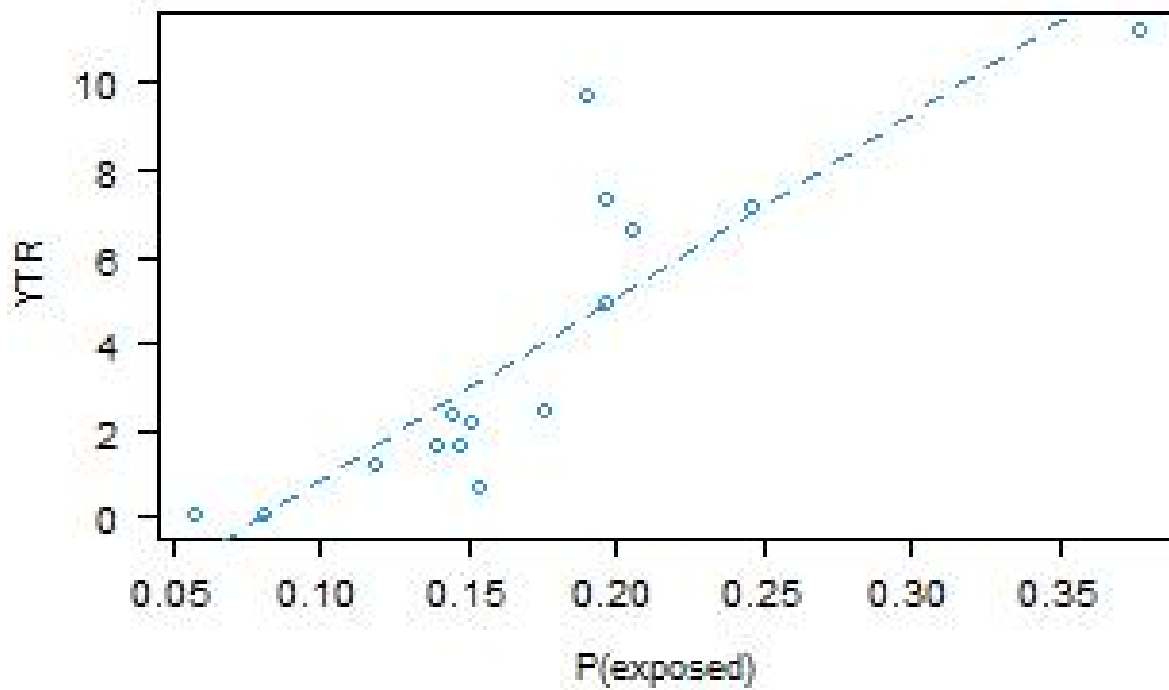
Here we plot the values of the injury metrics as a function of the four key parameters going into the model.

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

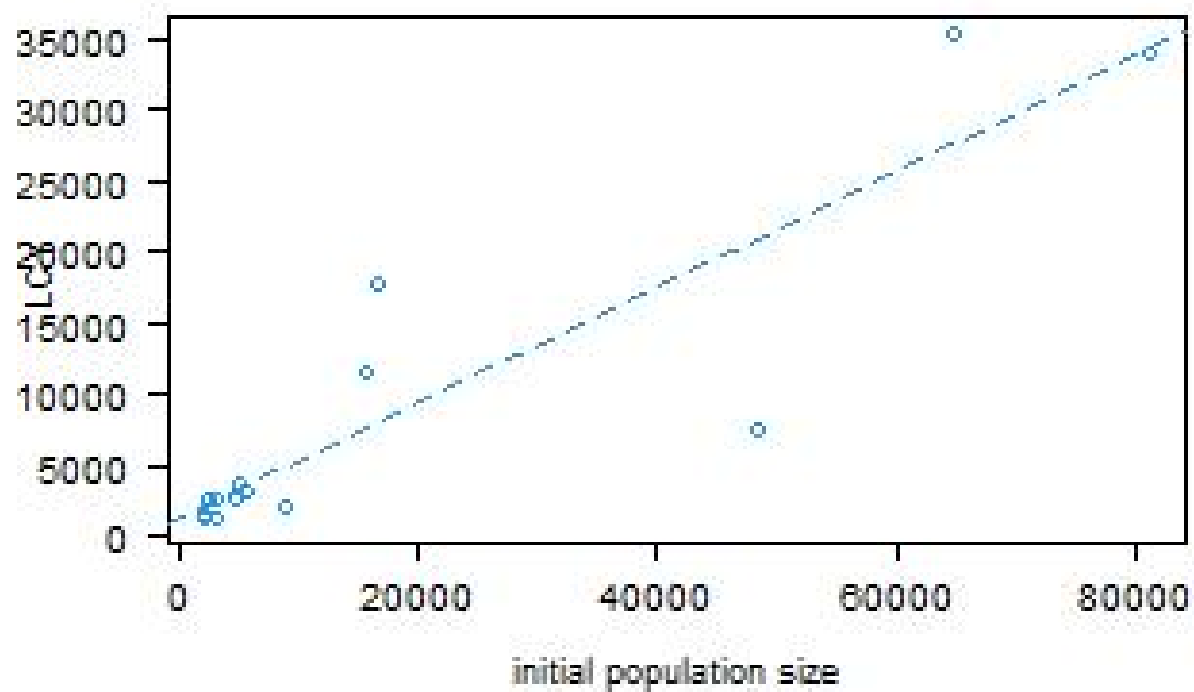


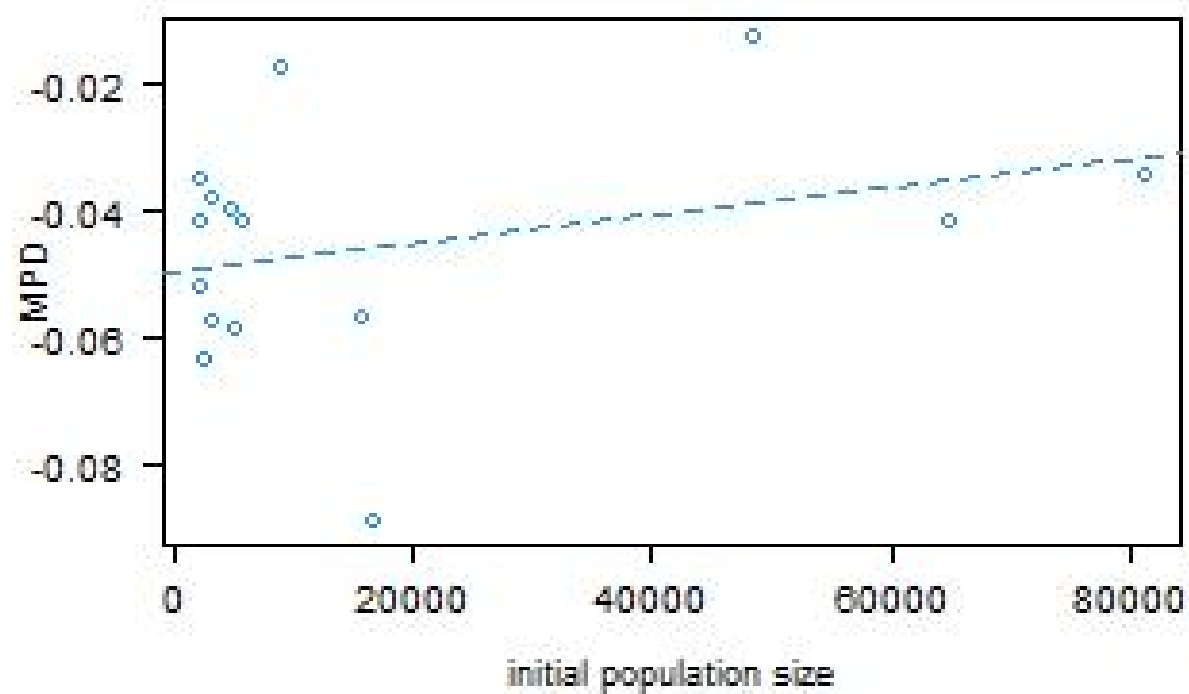


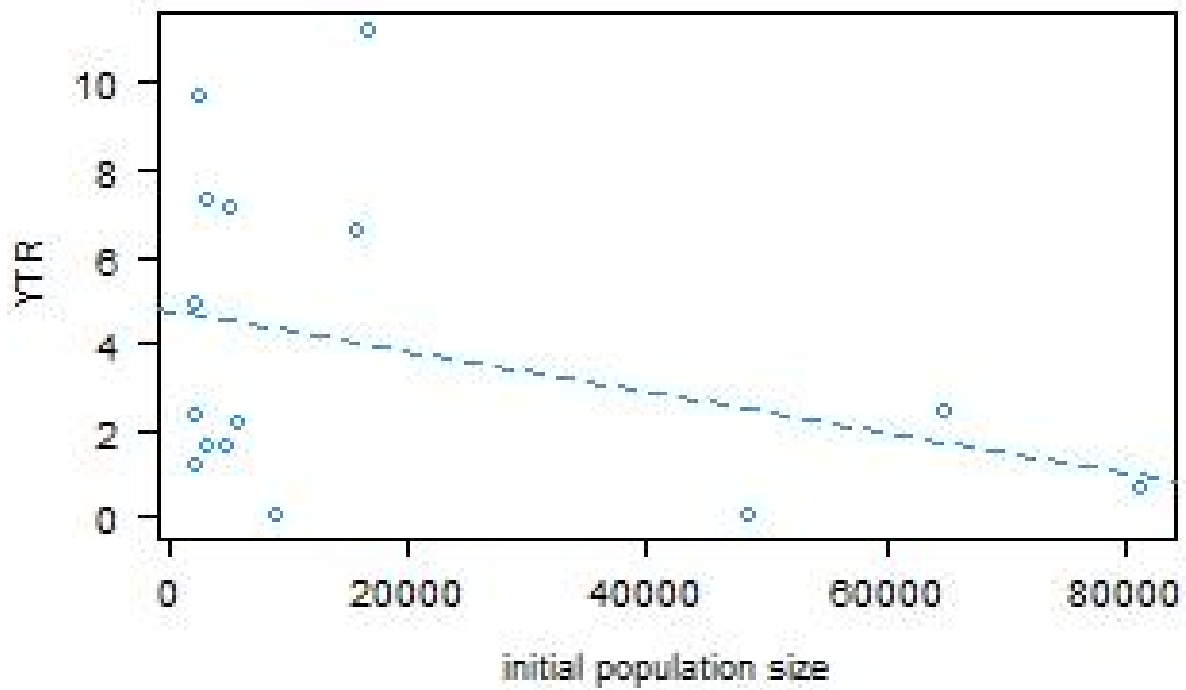


Initial population size

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.

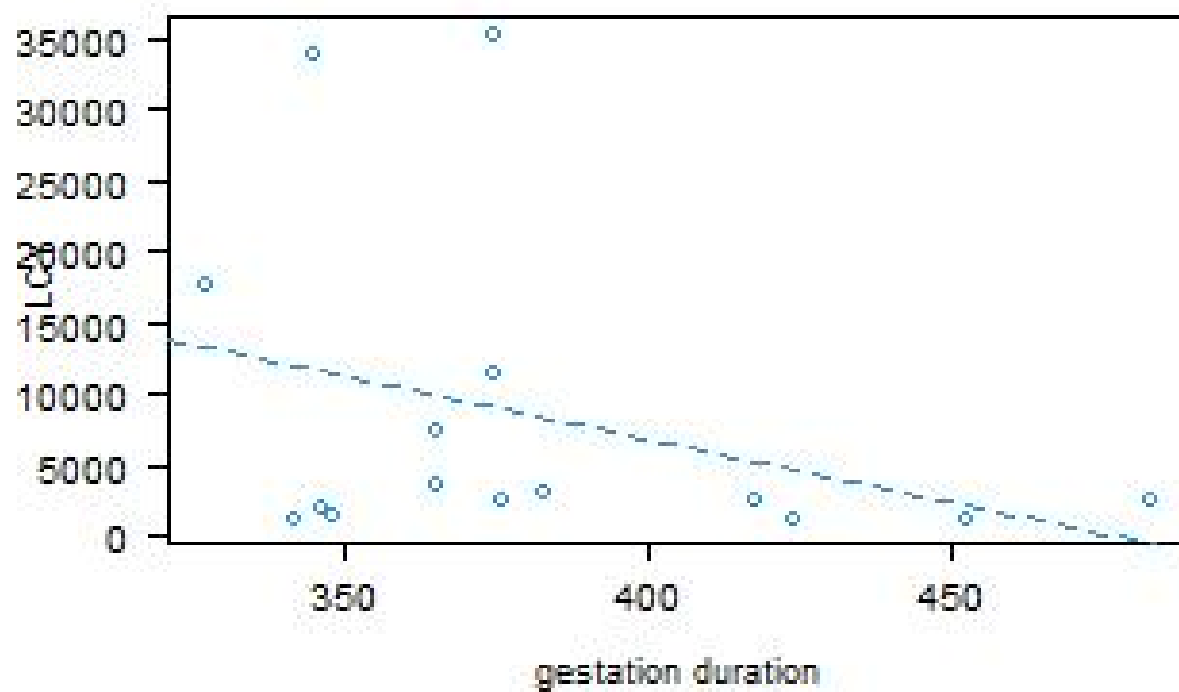


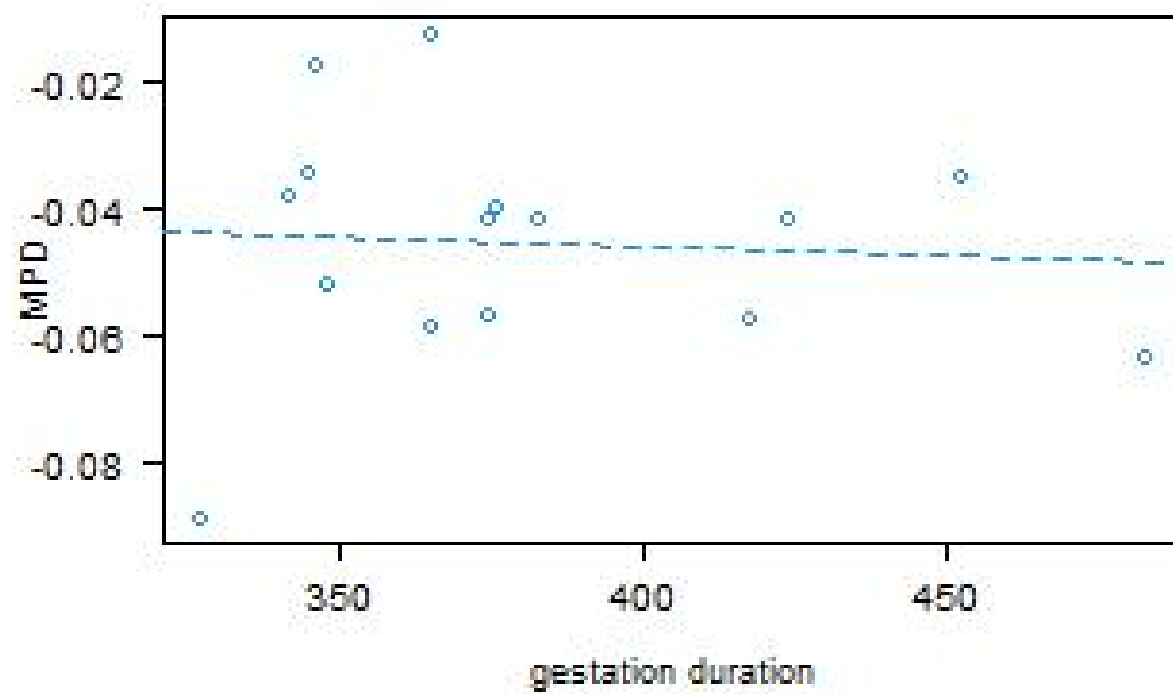


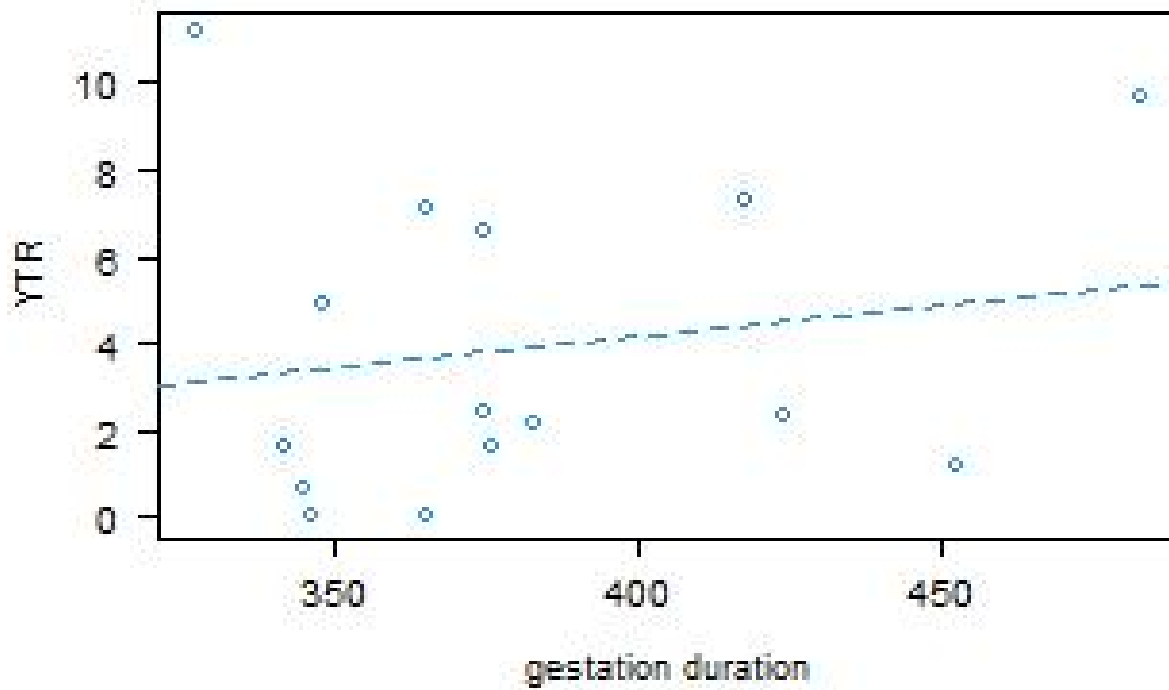


Gestation duration

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

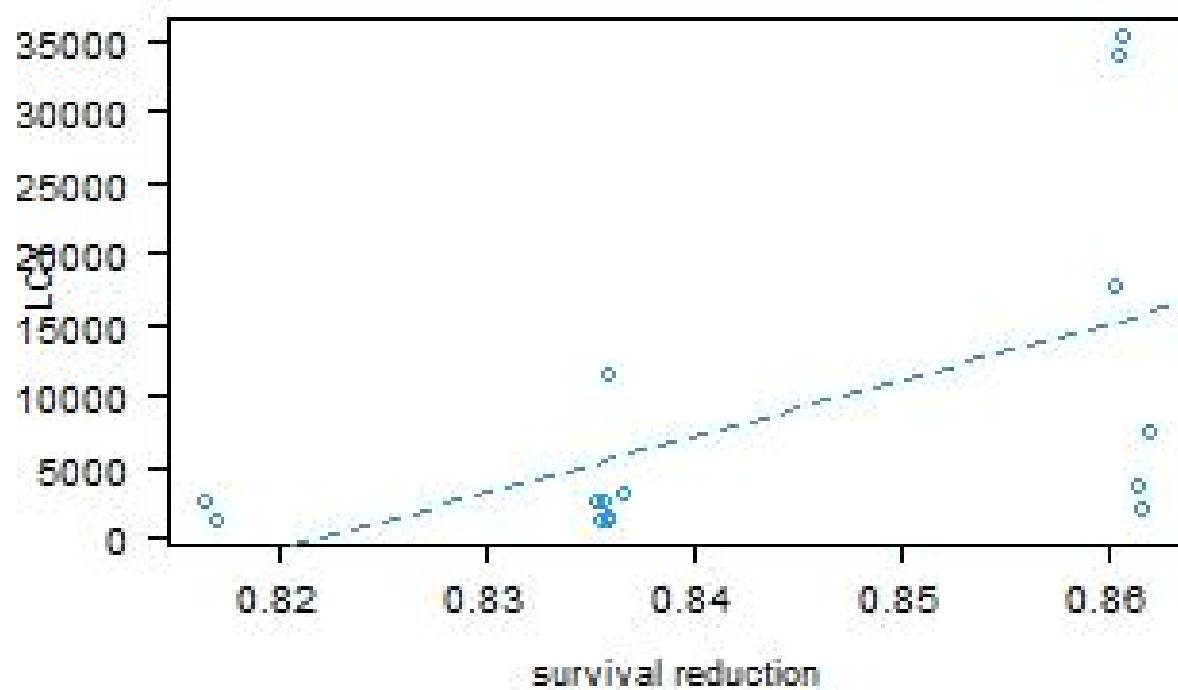


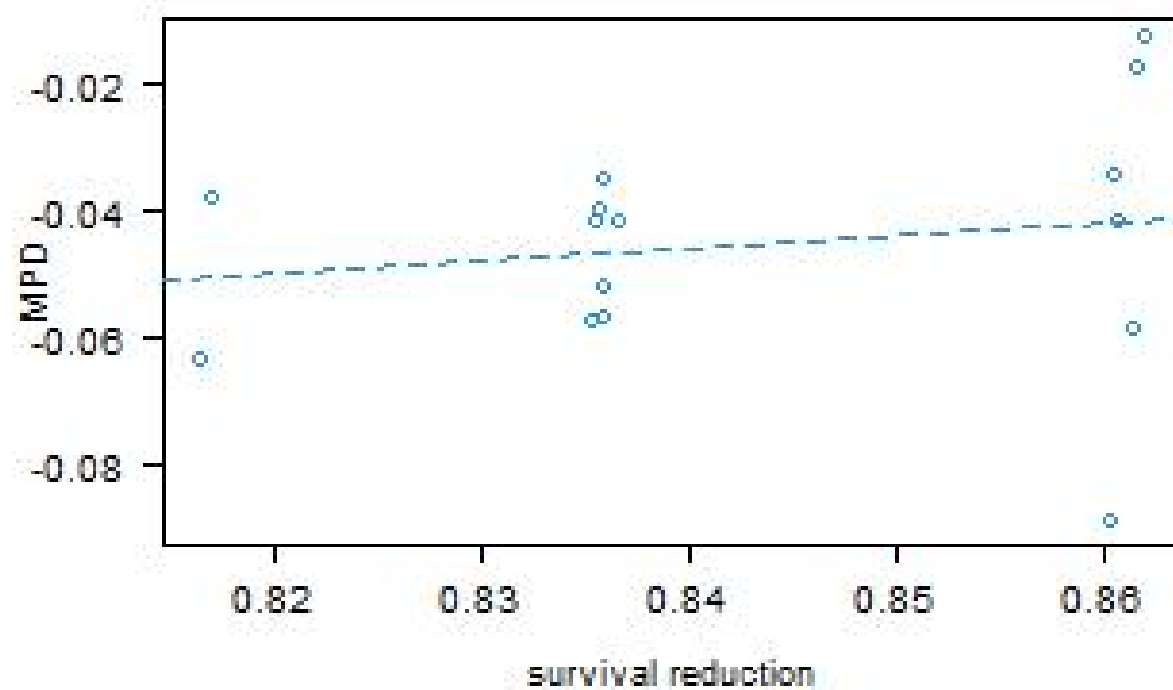


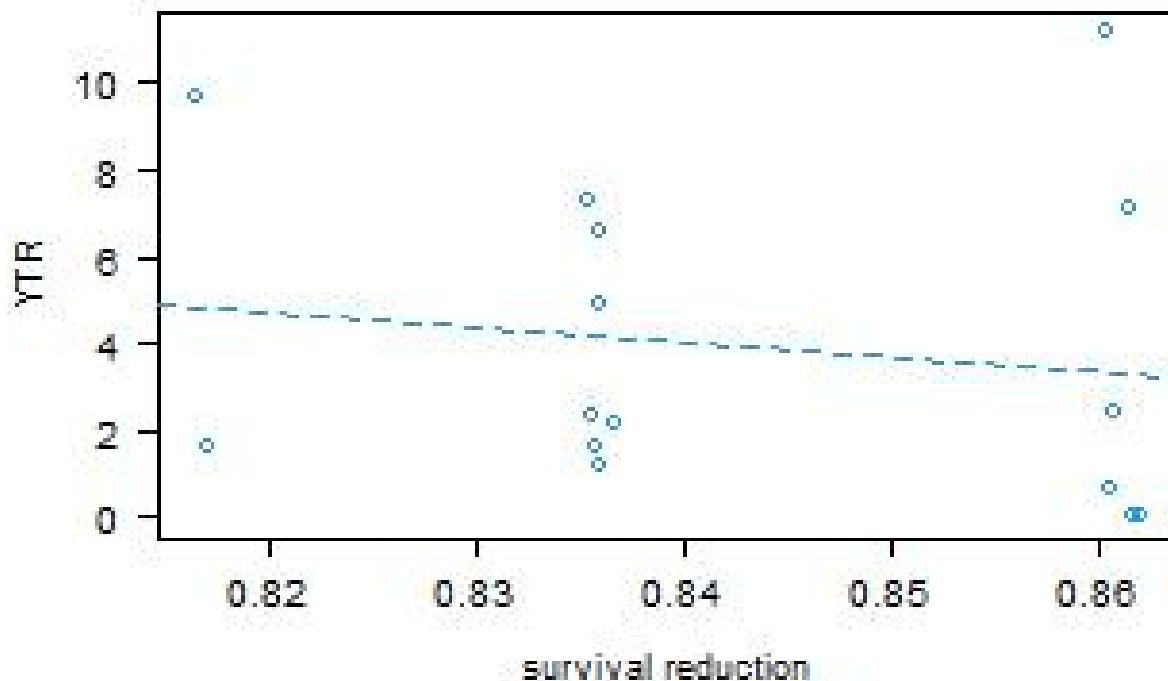


Survival reduction

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.

LCY

As noted above, one can anticipate that LCY would be strongly dependent on the initial population size and the proportion exposed. Therefore, we fit a linear model, treating species as sampling units, to explain LCY as a function of these two quantities. The summary of the corresponding linear model follows. Then we present a plot representing the observed values of LCY and those obtained by predicting from fitting a linear model of LCY as explained by initial population size and the proportion exposed.

```
##
## Call:
## lm(formula = LCY ~ Pexp + Nstart, data = sumres)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -7835.4 -1189.7   855.8  1477.7  6430.2
##
## Coefficients:
##              Estimate Std. Error t value    Pr(>|t|)
## (Intercept) -8585.80860  2457.09226  -3.494    0.004428 **
## Pexp         54656.60562 12278.22337   4.452    0.000791 ***
## Nstart         0.42640    0.03567  11.952 0.0000000505 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

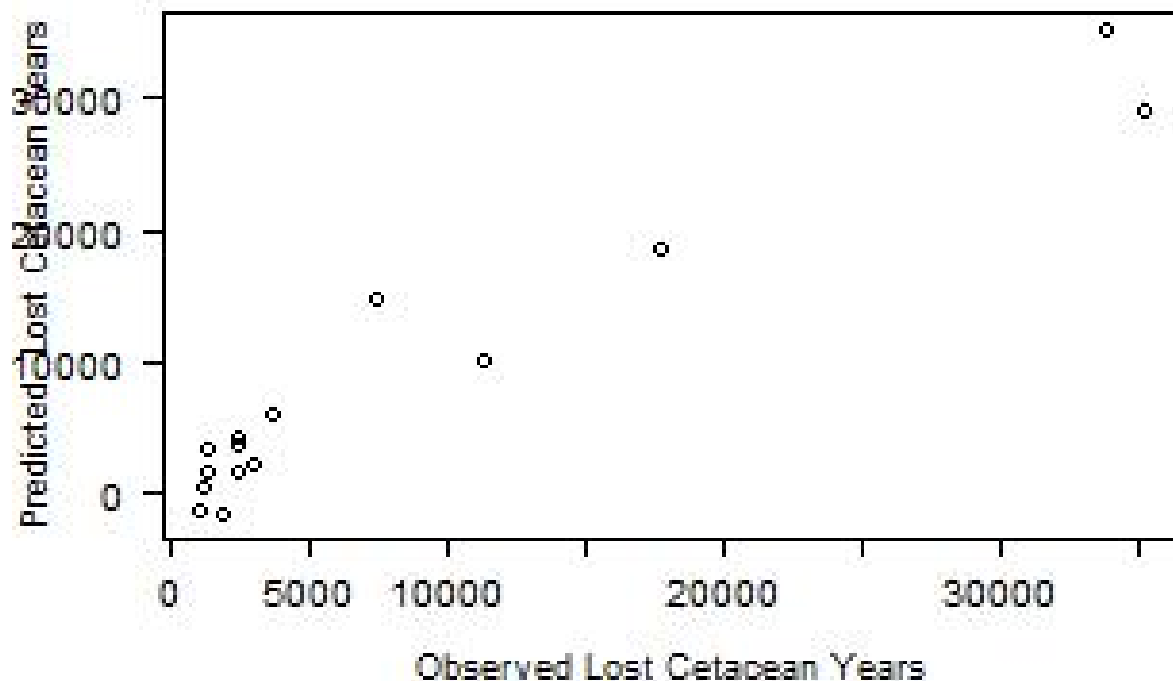
```
##
## Residual standard error: 3380 on 12 degrees of freedom
## Multiple R-squared:  0.9263, Adjusted R-squared:  0.9141
## F-statistic: 75.45 on 2 and 12 DF,  p-value: 0.0000001598
```

The LCY is very well explained by these two variable alone. A bit surprisingly, adding the gestation duration (that scales survival across species) does not even seem to help in explaining the injury

```
##
## Call:
## lm(formula = LCY ~ Pexp + Nstart + inputs$gd, data = sumres)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -7666.8 -1329.1   628.1  1565.4  6130.3
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -13896.68340   9907.08218  -1.403   0.1883
## Pexp         56431.54254  13047.43272   4.325   0.0012 **
## Nstart         0.43441     0.03949  11.000 0.000000283 ***
## inputs$gd      12.73743     22.97221   0.554   0.5904
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 3482 on 11 degrees of freedom
## Multiple R-squared:  0.9283, Adjusted R-squared:  0.9088
## F-statistic:  47.5 on 3 and 11 DF,  p-value: 0.000001384
```

The same seems to happen for the survival reduction, i.e., once population size and proportion exposed are accounted for, the remaining variation in LCY is negligible.

```
##
## Call:
## lm(formula = LCY ~ Pexp + Nstart + inputs$sr, data = sumres)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -7206.9 -1455.1  -213.6  1221.0  6326.1
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  41074.27726  60179.34881   0.683 0.509021
## Pexp         56953.36993  12751.08710   4.467 0.000952 ***
## Nstart         0.45048     0.04645   9.698 0.000001 ***
## inputs$sr    -59856.36279  72473.32346  -0.826 0.426406
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 3426 on 11 degrees of freedom
## Multiple R-squared:  0.9306, Adjusted R-squared:  0.9117
## F-statistic: 49.19 on 3 and 11 DF,  p-value: 0.000001158
```



```
## Analysis of Variance Table
##
## Response: LCY
##      Df      Sum Sq   Mean Sq  F value    Pr(>F)
## Pexp    1   91857453   91857453    8.0395   0.01502 *
## Nstart    1 1632291730 1632291730 142.8602 0.00000005055 ***
## Residuals 12  137109617   11425801
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 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. 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.

YTR

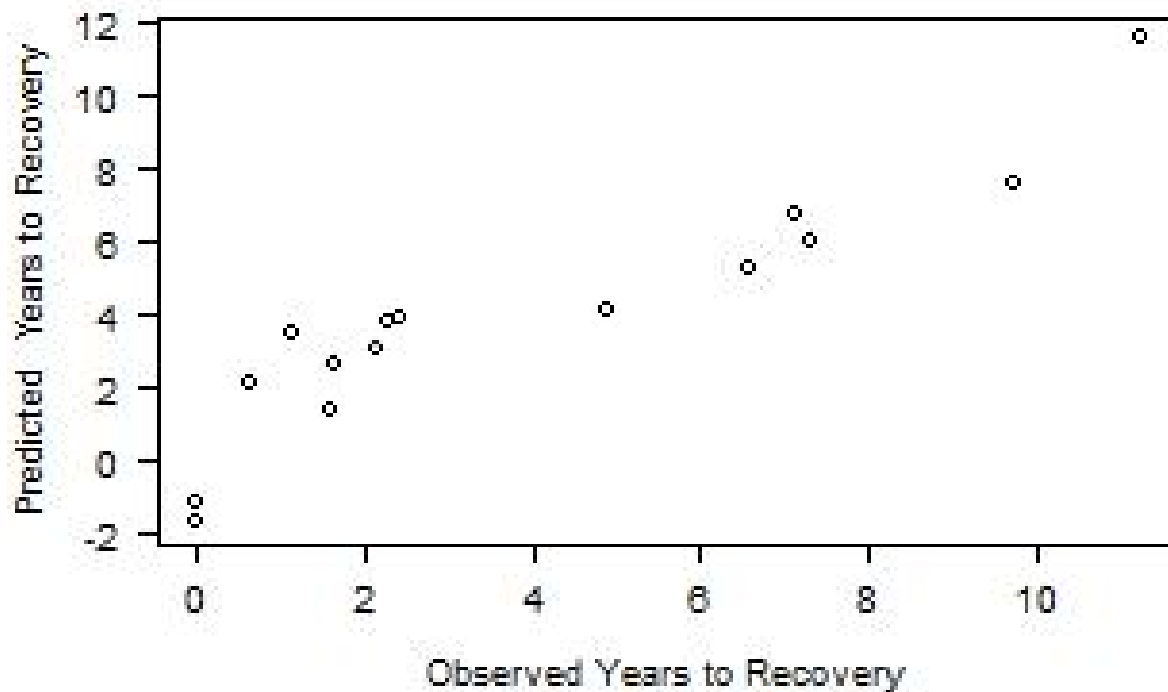
For YTR the following model describes which input parameters help explain the YTR

```
##
## Call:
## lm(formula = YTR ~ Pexp + Nstart + inputs$gd + inputs$sr, data = sumres)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -2.3709 -0.9515 -0.2810  1.0868  2.2222
##
```

```
## Coefficients:
##           Estimate Std. Error t value Pr(>|t|)
## (Intercept) -6.91627839 33.53726432 -0.206 0.8408
## Pexp        44.13367418  6.07960386  7.259 0.0000273 ***
## Nstart      -0.00001197  0.00002185 -0.548 0.5959
## inputs$gd    0.02388869  0.01171380  2.039 0.0687 .
## inputs$sr   -6.75814404 37.56228534 -0.180 0.8608
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 1.61 on 10 degrees of freedom
## Multiple R-squared:  0.8591, Adjusted R-squared:  0.8028
## F-statistic: 15.24 on 4 and 10 DF,  p-value: 0.000294
```

We removing initial population size and survival reduction to obtain a more parsimonious model.

```
##
## Call:
## lm(formula = YTR ~ Pexp + inputs$gd, data = sumres)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -2.364 -1.229  0.256  1.220  2.177
##
## Coefficients:
##           Estimate Std. Error t value Pr(>|t|)
## (Intercept) -14.465792  3.864204 -3.744 0.0028 **
## Pexp        44.996205  5.546928  8.112 0.00000326 ***
## inputs$gd    0.027785  0.009304  2.986 0.0113 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 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
```



```
## Analysis of Variance Table
##
## Response: YTR
##           Df Sum Sq Mean Sq F value    Pr(>F)
## Pexp       1 135.955 135.955   59.199 0.000005592 ***
## inputs$gd   1  20.483  20.483    8.919  0.01135 *
## Residuals 12   27.559    2.297
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 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.

MPD

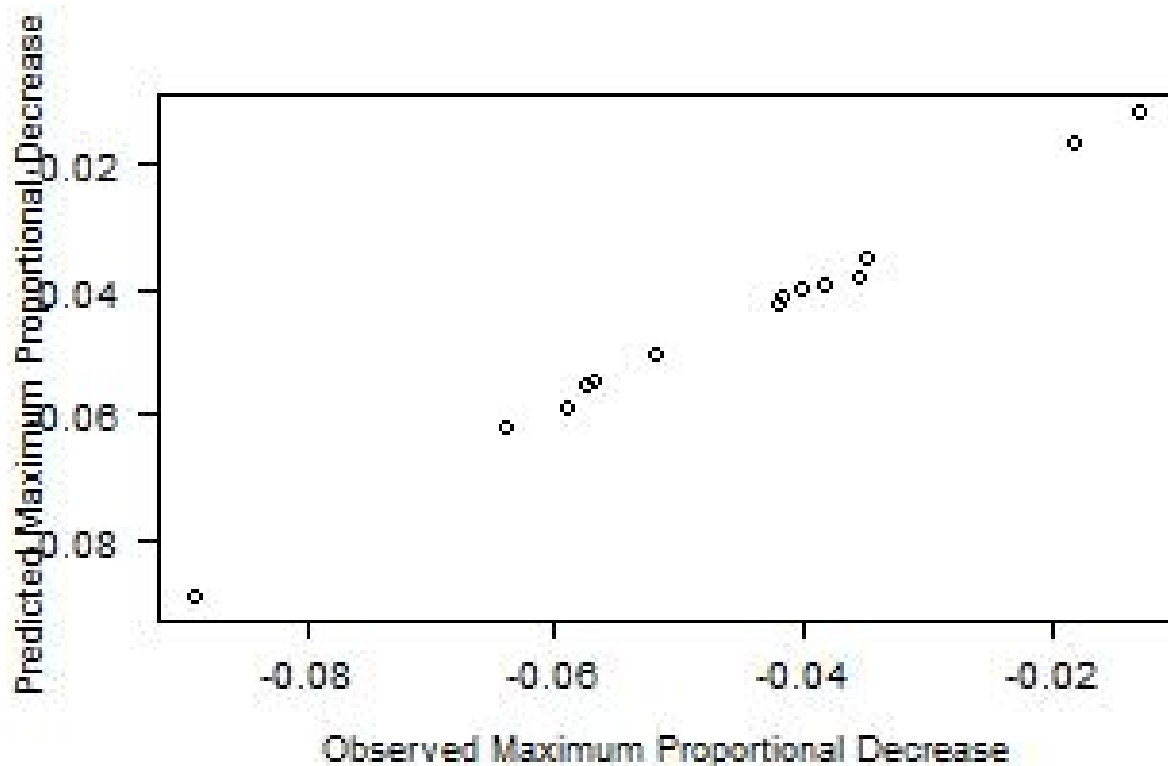
The following model describes which input parameters help explain the MPD

```
##
## Call:
## lm(formula = MPD ~ Pexp + Nstart + inputs$gd + inputs$sr, data = sumres)
##
## Residuals:
##           Min           1Q       Median           3Q          Max
## -0.0017905 -0.0011313  0.0003510  0.0006739  0.0030961
##
## Coefficients:
##              Estimate      Std. Error t value      Pr(>|t|)
```

```
## (Intercept) -0.143158704426  0.034568588676  -4.141          0.002008 **
## Pexp        -0.250521421037  0.006266561376 -39.977 0.000000000000229 ***
## Nstart       0.000000001912  0.000000022525   0.085          0.934020
## inputs$gd    -0.000066021093  0.000012074015  -5.468          0.000274 ***
## inputs$sr     0.196963942281  0.038717385514   5.087          0.000473 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.00166 on 10 degrees of freedom
## Multiple R-squared:  0.9943, Adjusted R-squared:  0.9921
## F-statistic: 438.8 on 4 and 10 DF,  p-value: 0.00000000003484
```

Removing initial population size just to check if the model is more parsimonious, which it happens to be.

```
##
## Call:
## lm(formula = MPD ~ Pexp + inputs$gd + inputs$sr, data = sumres)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -0.0017993 -0.0011580  0.0003586  0.0006988  0.0030760
##
## Coefficients:
##              Estimate Std. Error t value      Pr(>|t|)
## (Intercept) -0.14460731  0.02867493  -5.043    0.000376 ***
## Pexp        -0.25065194  0.00579441 -43.258 0.000000000000123 ***
## inputs$gd    -0.00006607  0.00001150  -5.743    0.000130 ***
## inputs$sr     0.19876972  0.03085665   6.442 0.000047965099459 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.001583 on 11 degrees of freedom
## Multiple R-squared:  0.9943, Adjusted R-squared:  0.9928
## F-statistic: 643.1 on 3 and 11 DF,  p-value: 0.000000000001241
```



```
## Analysis of Variance Table
##
## Response: MPD
##           Df      Sum Sq   Mean Sq    F value    Pr(>F)
## Pexp       1 0.0044337 0.0044337 1769.479 0.0000000000001668 ***
## inputs$gd   1 0.0002966 0.0002966  118.370 0.0000003163698008 ***
## inputs$sr   1 0.0001040 0.0001040   41.496 0.0000479650994587 ***
## Residuals 11 0.0000276 0.0000025
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 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 2.14 %, respectively. About 0.57 % of the variability remains unexplained.

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.Injury.to.GOM.Cetaceans [LINK](#).

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