

model-comparison

Alexander Keth

2017-05-18

Contents

0.1 User Input

2

1 Whole system biomass

3

2 Biomass timeseries

4

3 Relative biomass timeseries

5

4 Physics

6

NOTE: This vignette is optimised for longer simulation runs. Therefore the output is not as pleasant due to the fact that the dummy setas file have a running time of 5 years.

In order to use this vignette make sure to render `model-preprocess.Rmd` for each simulation first. Save the resulting list of dataframes as shown in `data-raw/data-vignette-model-pr`. Of course, you can also use a personalised version of `model-preprocess.Rmd`. Please make sure to add all resulting dataframes to the list of dataframes at the end of the preprocess vignette and change `model-comparison.Rmd` accordingly.

```
library("atlantistools")
library("ggplot2")
library("gridExtra")
```

```
gen_labels <- list(x = "Time [years]", y = "Biomass [t]")
```

```
# You should be able to build the vignette either by clicking on "Knit PDF" in RStudio or with
# rmarkdown::render("model-comparison.Rmd")
```

0.1 User Input

This section is used to read in the simulations. In order to demonstrate the vignette, dummy simulations are generated. Please change this accordingly.

```
result <- preprocess
```

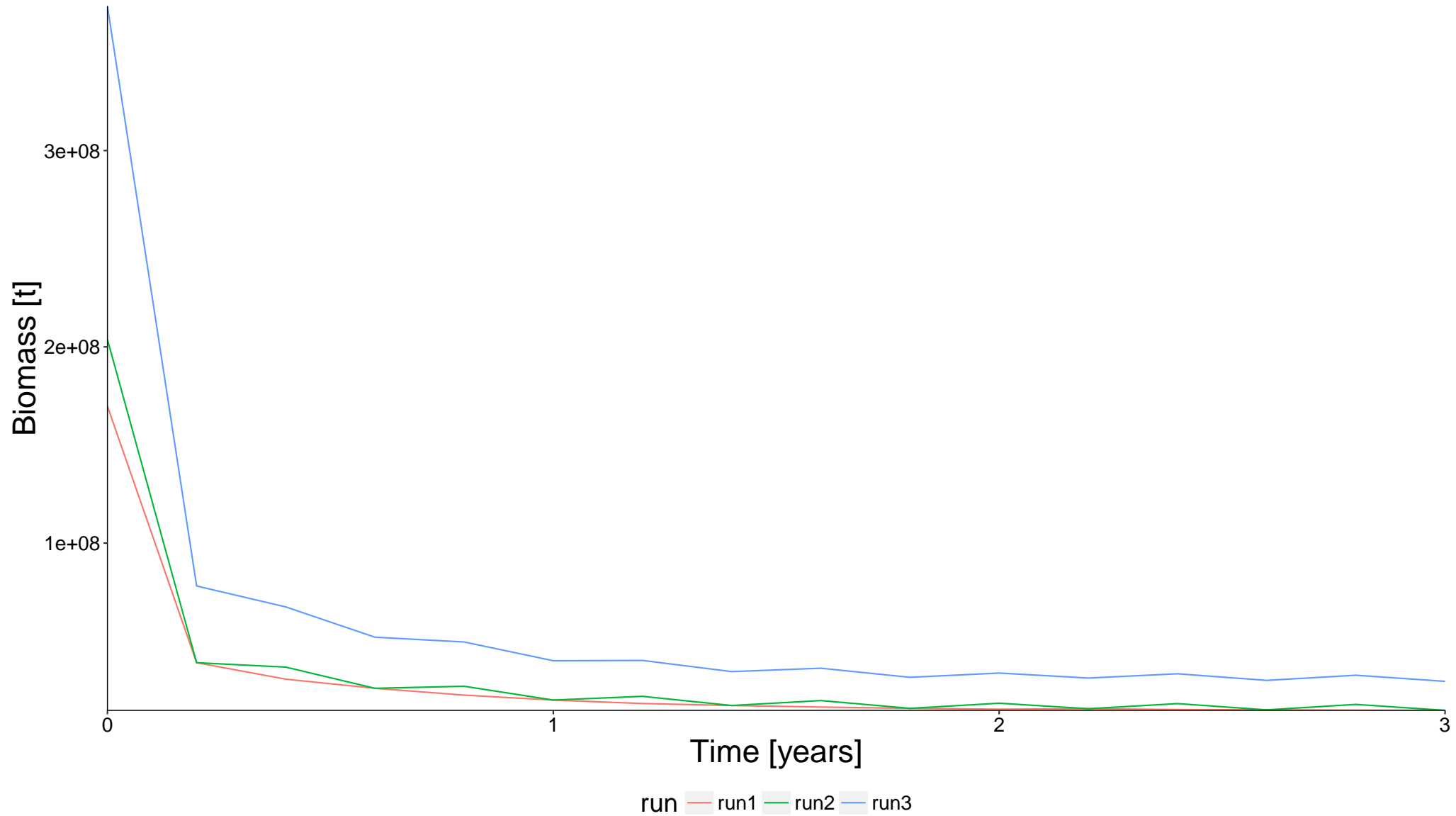
```
dummy_setas <- function(list, mult) {
  for (i in seq_along(list)) {
    if (is.data.frame(list[[i]])) {
      mult <- rep_len(mult, length.out = nrow(list[[i]]))
      list[[i]][, ncol(list[[i]])] <- list[[i]][, ncol(list[[i]])] * mult
    }
  }
  return(list)
}
```

```
store_data <- list(result,
  dummy_setas(result, mult = c(1, 1.2)),
  dummy_setas(result, mult = c(2, 2.2)))
```

```
result <- combine_runs(outs = store_data, runs = c("run1", "run2", "run3"))
```

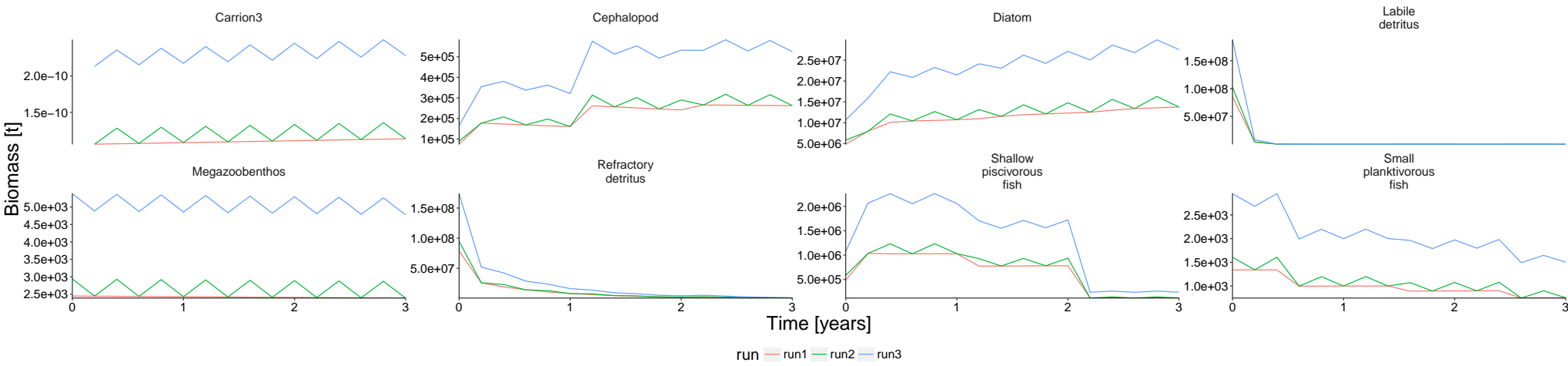
1 Whole system biomass

```
sum_bio <- agg_data(result$biomass, groups = c("time", "run"), fun = sum)
plot <- plot_line(sum_bio, wrap = NULL, col = "run")
update_labels(plot, gen_labels)
```



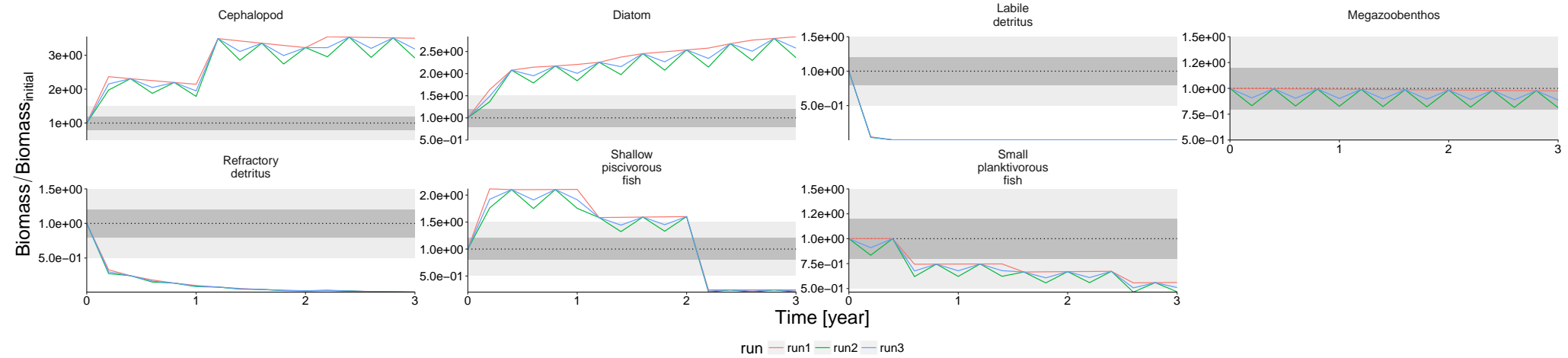
2 Biomass timeseries

```
plot_line(result$biomass, col = "run", ncol = 4) %>% update_labels(gen_labels)
```



3 Relative biomass timeseries

```
df <- convert_relative_initial(result$biomass)
plot <- plot_line(df, col = "run", ncol = 4)
plot <- plot_add_box(plot)
update_labels(plot, labels = list(x = "Time [year]", y = expression(Biomass/Biomass[initial])))
```



4 Physics

```
plot <- plot_line(result$physics, wrap = NULL, col = "run")  
custom_grid(plot, grid_x = "polygon", grid_y = "variable")
```

