A model for changes in length frequencies

1 Data

Read in length data and modify some column names and variable labels for use below.

```
library (gdata)
setwd("../data")
neph.dat <- read.xls("Celtic Warrior Diamond mesh July 2014 Celtic Sea.xls",
    sheet = "Nephrops Lengths", stringsAsFactors = FALSE)
## Show the first 2 rows
head(neph.dat, 2)
             Vessel DATE HAUL COMPARTMENT Mesh.Size SPECIES
## 1 Celtic Warrior 2014-07-19 1 Control 70mm Nephrops   
## 2 Celtic Warrior 2014-07-19 1 Control 70mm Nephrops
## Carapace.Length..mm.. COUNT SUBSRATIO
                        16 1 1
## 1
## 2
                         17
                               11
## Change the carapace length name
names (neph.dat) [names (neph.dat) == "Carapace.Length..mm.."] <- "Carapace.Length"</pre>
## Make the 'HAUL' variable character
neph.dat$HAUL <- paste("H", neph.dat$HAUL, sep = "")</pre>
```

Make one row per length measurement assuming, for example, that a sub-sampling ratio of 0.1 corresponds to 10% of the catch sampled (CHECK).

Read in the haul weights

```
setwd("../data")
weight.dat <- read.xls("Celtic Warrior Diamond mesh July 2014 Celtic Sea.xls",
    sheet = "Weights", stringsAsFactors = FALSE)
## Show the first 2 rows
head(weight.dat, 2)
##
          Date Haul.. Compartment Mesh. Size Species Total. weight..kg.
## 1 2014-07-19 1 TEST1 90mm Bulk 26.28
## 2 2014-07-19
                   1
                            TEST1
                                      90mm Haddock
                                                               0.38
## Sbsample.weight..kg.
## 1
## 2
## create a new 'HAUL' variable for the merge
weight.dat$HAUL <- paste("H", weight.dat$Haul.., sep = "")</pre>
## re-name total weight column
names (weight.dat) [names (weight.dat) == "Total.weight.kg."] <- "Total.Weight"</pre>
```

Merge the bulk weights with the length data

Produce a summary plot of the data by length, haul and catch weight.

```
library(ggplot2)
library(gridExtra)

## quick function for plot
plot.lfreq <- function(data, title.string){
  p <- ggplot(data, aes(x = Carapace.Length, group = HAUL)) +
      geom_density(position = "stack", aes(fill = Total.Weight), colour = 1, lwd = 0.01) +</pre>
```

```
xlim(10, 45) + scale_fill_gradient2(low = "white", high = "blue", limits theme(axis.text.x=element_blank())
return(p)
}

p70mm <- plot.lfreq(neph.70mm, "70mm")
p80mm <- plot.lfreq(neph.80mm, "80mm")
p90mm <- plot.lfreq(neph.90mm, "90mm")
p100mm <- plot.lfreq(neph.100mm, "100mm")
grid.arrange(p70mm, p80mm, p90mm, p100mm, ncol = 1)</pre>
```

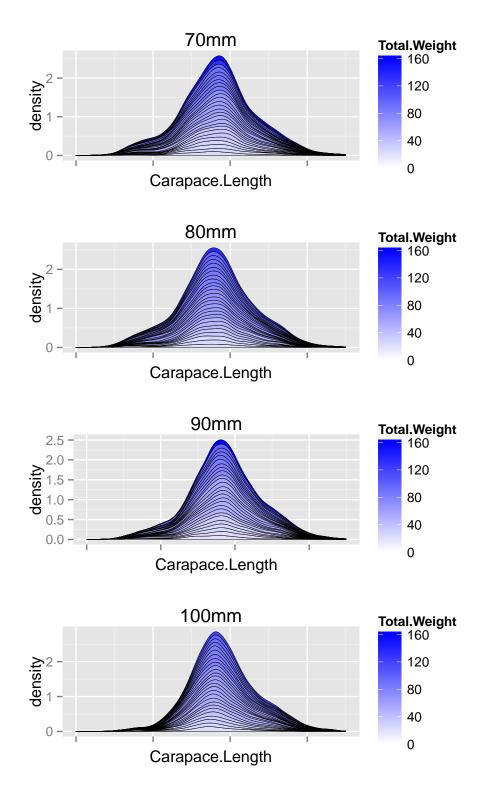


Figure 1: Stacked carapace length densities. Each haul is coloured according to the total bulk weight in that haul.