

Chapter 1 Introduction

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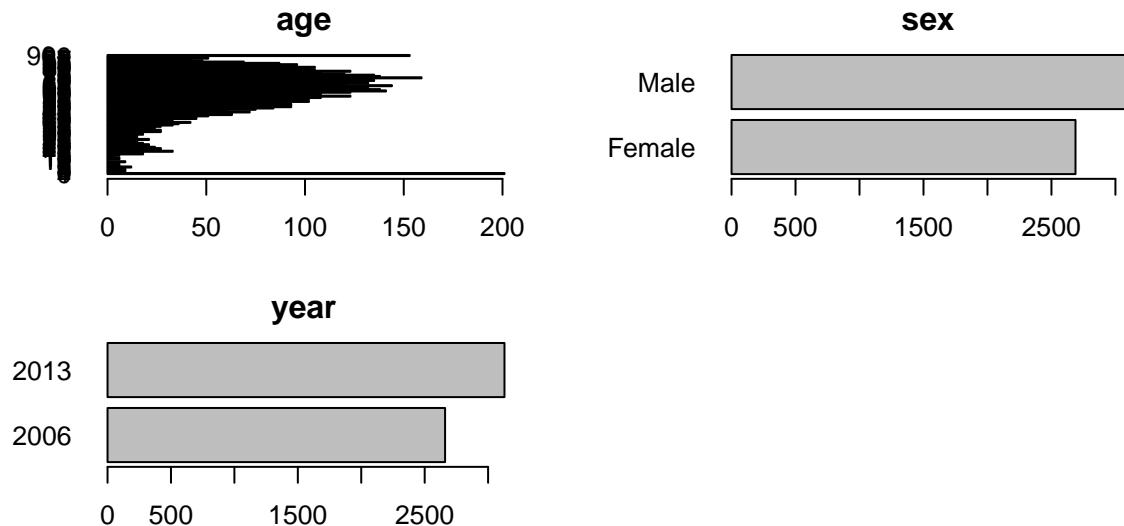
Setup

```
library(bdefdata)
library(demest)
library(demlife)
library(dplyr)
library(latticeExtra)
```

Raw data

```
deaths <- bdefdata::maori_deaths %>%
  Counts(dimtypes = c(year = "state")) %>%
  subarray(year %in% c(2006, 2013)) %>%
  collapseIntervals(dimension = "age", old = c(90:99, "100+"))
```

```
plot(deaths)
```



```
popn <- bdefdata::maori_popn %>%
  Counts(dimtypes = c(year = "state")) %>%
  subarray(year %in% c(2006, 2013))
```

```
plot(popn)
```



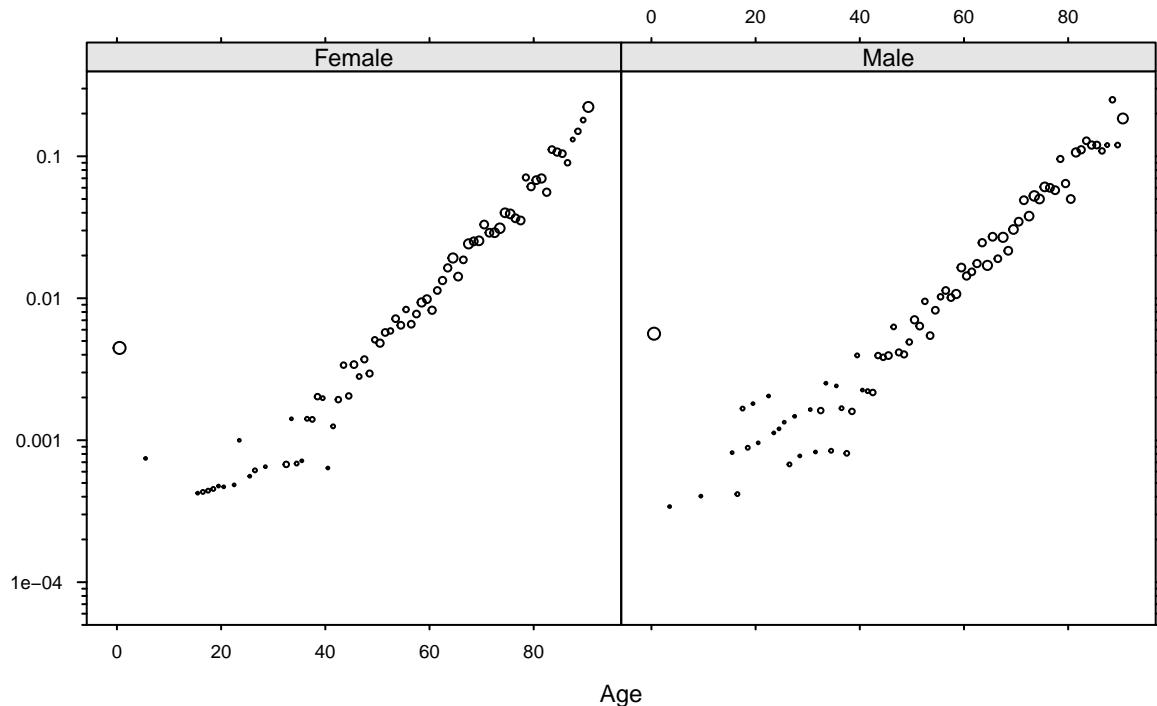
Direct estimates

```
direct <- (deaths / popn) %>%
  as.data.frame(direction = "long", midpoints = "age")

cex <- 0.1 * sqrt(as.numeric(deaths))

p_direct <- xyplot(value ~ age | sex,
                     data = direct,
                     subset = year == "2013",
                     as.table = TRUE,
                     col = "black",
                     ylab = "",
                     xlab = "Age",
                     ylim = c(0.00005, NA),
                     scales = list(x = list(tck = 0.3),
                                   y = list(log = TRUE, tck = 0.6)),
                     yscale.components = yscale.components.log10ticks,
                     par.settings = list(fontsize = list(text = 9),
                                         strip.background = list(col = "grey90")),
                     cex = cex)

p_direct
```



Fit Model

```

model <- Model(y ~ Poisson(mean ~ age * sex + year),
                 age ~ DLM(damp = NULL,
                            covariates = Covariates(infant = TRUE)),
                 age:sex ~ DLM(trend = NULL,
                                damp = NULL),
                 jump = 0.065)

datasets <- list(deaths = deaths)

data_models <- list(Model(deaths ~ Round3()))

filename <- tempfile()

set.seed(0)

## warning - this takes a long time
estimateCounts(model,
                y = deaths,
                exposure = popn,
                dataModels = data_models,
                datasets = datasets,
                filename = filename,
                nBurnin = 400000,
                nSim = 400000,
                nChain = 4,
                nThin = 500)

options(width = 150)
fetchSummary(filename)

```

```

## -----
## model:
## y ~ Poisson(mean ~ age * sex + year),
## age ~ DLM(damp = NULL, covariates = Covariates(infant = TRUE)),
## age:sex ~ DLM(trend = NULL, damp = NULL),
## 0.065
## dimensions: age, sex, year
## -----
## y:
## dimensions: age, sex, year
## n cells: 364
## -----
## Data models:
## *deaths*
## deaths ~ Round3()
## dimensions: age, sex, year
## -----
## Datasets:
## *deaths*
## Object of class "Counts"
## dimensions: age, sex, year
## n cells: 364, n missing: 0, integers: TRUE, n zeros: 33, median: 12
## -----
## MCMC statistics:
## nBurnin: 400000, nSim: 400000, nChain: 4, nThin: 500, nIteration: 3200
## 
## Metropolis-Hastings updates:
##          jump acceptance autocorr
## model.likelihood.rate 0.065      0.364    0.374
## 
## parameters:
##                               Rhat    2.5%   50% 97.5% length
## model.likelihood.rate     1.05 0.000185 0.00391 0.162    364
## model.prior.mean          1.03 -8.59   -5.54 -1.82    364
## model.prior.sd            1.02 0.00172  0.0305 0.0631     1
## model.hyper.age.scaleLevel 1.01 0.00228  0.0357 0.0978     1
## model.hyper.age.scaleTrend 1.05 0.00171  0.0107 0.0622     1
## model.hyper.age.coef       1.04  2.56    3.26  3.9      1
## model.hyper.age.scaleError 1.00 0.000835  0.0155 0.0562     1
## model.hyper.age:sex.scaleLevel 1.04 0.0114  0.0414 0.0828     1
## model.hyper.age:sex.scaleError 1.01 0.00481  0.0285 0.065      1
## y                           1.00    0     12    43    364
## -----

```

Plot

```

rates <- fetch(filename,
               where = c("model", "likelihood", "rate"))

summary(rates)

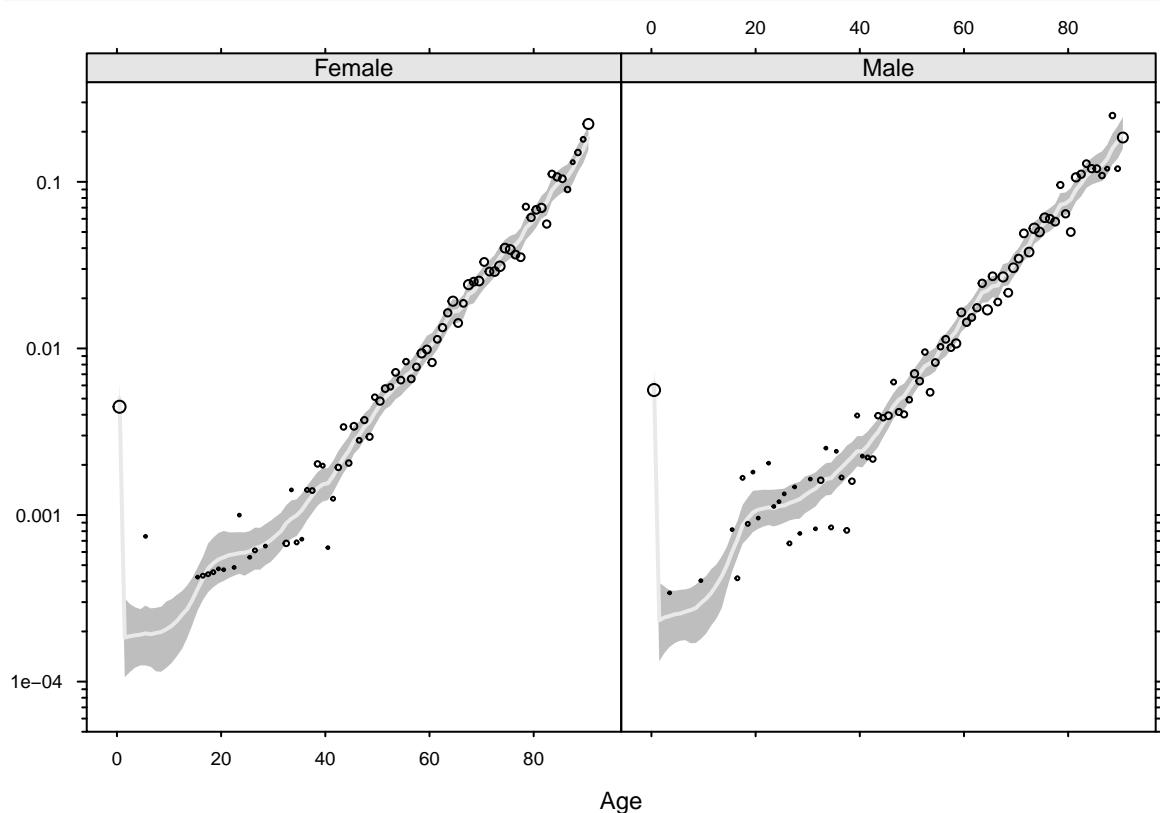
## 
## name:      age        sex        year        iteration

```

```

## length: 91      2      2      3200
## dimtype: age    sex    state   iteration
## dimscale: Intervals Sexes Categories Iterations
## first: 0      Female 2006      1
## last: 90+     Male   2013     3200
##
##      Min.   1st Qu.   Median   Mean   3rd Qu.   Max.
## 0.0000699 0.0008381 0.0038899 0.0252276 0.0271123 0.3452140
p_modelled <- p_direct +
  as.layer(dplot(~ age | sex,
                 data = rates,
                 subarray = year == "2013",
                 midpoints = "age",
                 prob = c(0.025, 0.5, 0.975),
                 lwd = 2,
                 col = "grey",
                 scales = list(y = list(log = TRUE))),
            under = TRUE)
p_modelled

```



Life expectancy

```

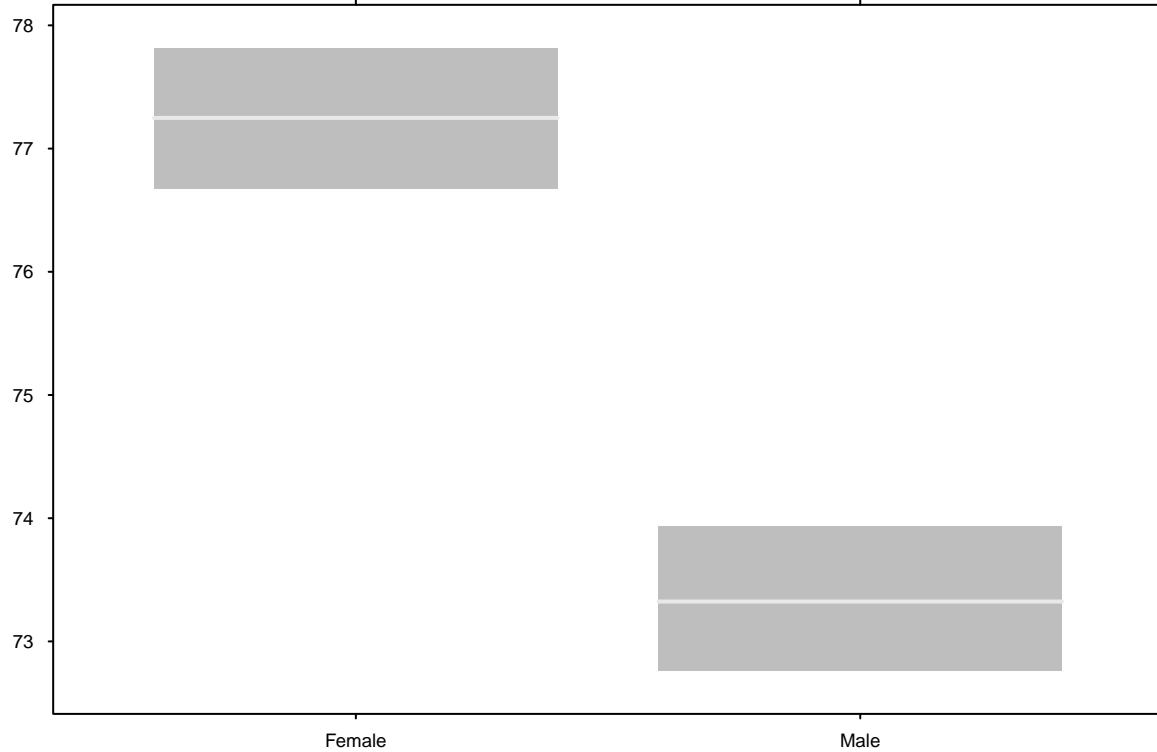
life_exp <- rates %>%
  LifeTable() %>%
  lifeExpectancy()

```

```

p_life_exp <- dplot(~ sex,
                      data = life_exp,
                      subarray = year == "2013",
                      as.table = TRUE,
                      col = "grey",
                      lwd = 2,
                      prob = c(0.025, 0.5, 0.975),
                      ylab = "",
                      xlab = "",
                      scales = list(tck = 0.3),
                      par.settings = list(fontsize = list(text = 9),
                                          strip.background = list(col = "grey90")))
p_life_exp

```



Unconfidentialized deaths

```

y <- fetch(filename, where = "y")
summary(y)

##
##  name:      age       sex     year      iteration
##  length:    91        2       2       3200
##  dimtype:   age       sex     state    iteration
##  dimscales: Intervals Sexes   Categories Iterations
##  first:     0        Female  2006      1
##  last:     90+       Male   2013      3200
##
##  Min. 1st Qu. Median     Mean 3rd Qu.   Max.

```

```
##      0.00    4.00   12.00   15.92   26.00   71.00
prob <- y %>%
  subarray(sex == "Male" & age == "20" & year == "2013") %>%
  table() %>%
  prop.table() %>%
  round(3)
prob

## .
##     4      5      6      7      8
## 0.093 0.215 0.362 0.230 0.100
```