

Chapter 19 Population in China

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2018-09-10

Setup

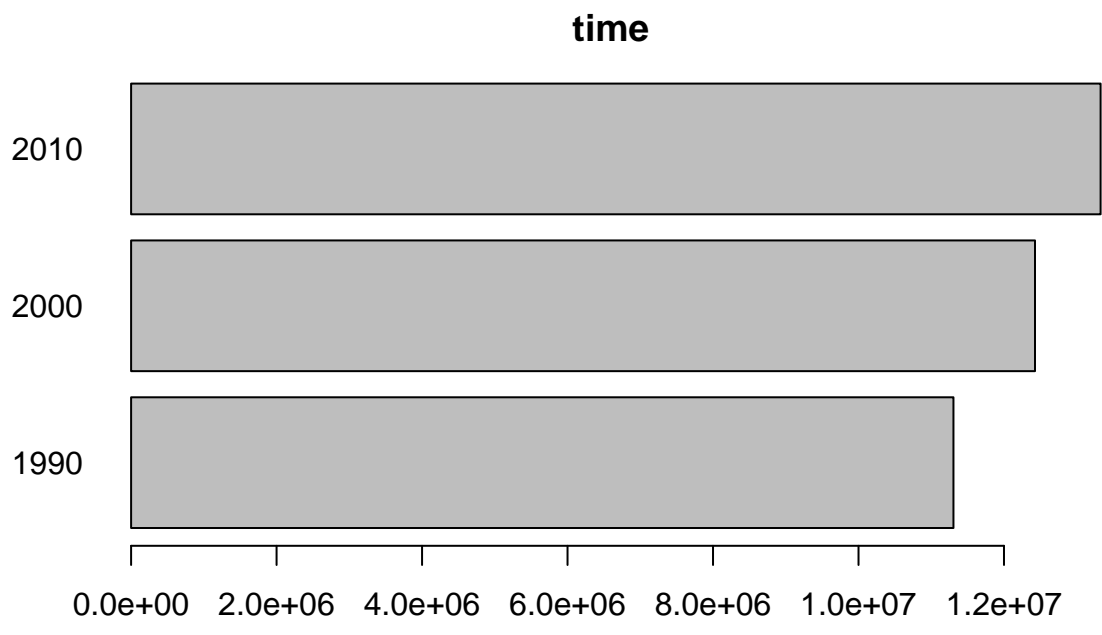
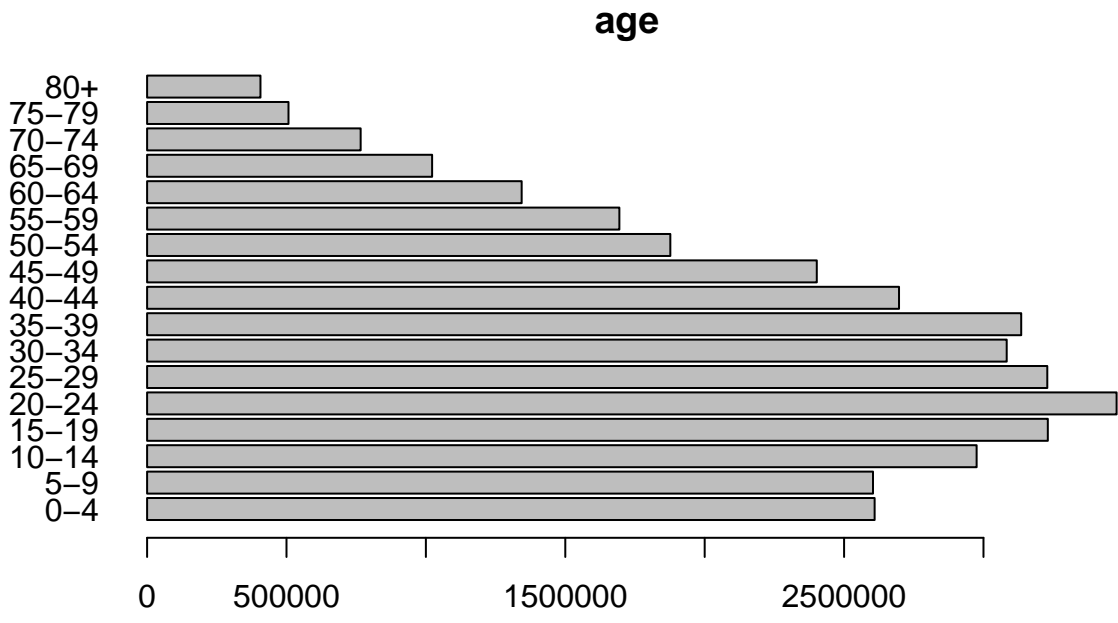
```
library(methods)
library(bdefdata)
library(demest)
library(demlife)
library(dplyr)
library(tidyverse)
library(latticeExtra)
```

Get data

```
census <- bdefdata::china_population_census %>%
  Counts()
summary(census)
```

```
##
## name:      age      time
## length:   17       3
## dimtype:   age      time
## dimscale: Intervals Points
## first:    0-4      1990
## last:     80+     2010
##
##      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
## 76764 414086 787532 726654 1020226 1274125
```

```
plot(census)
```

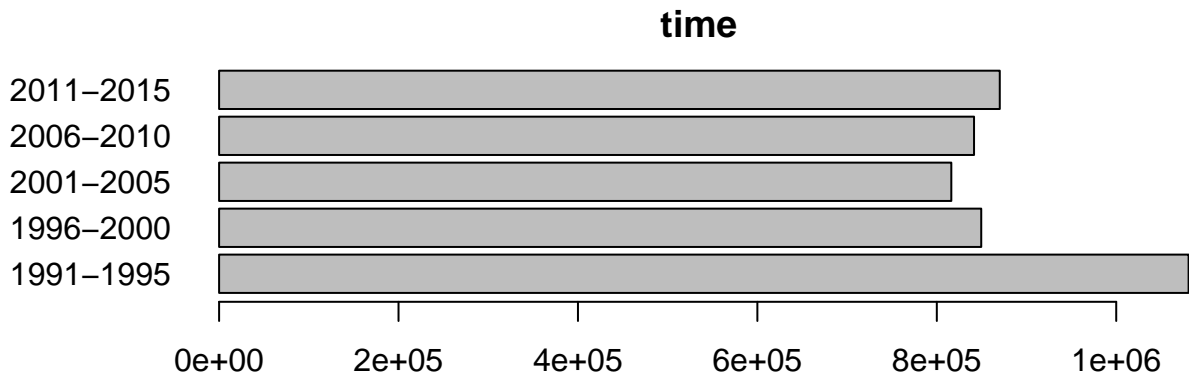
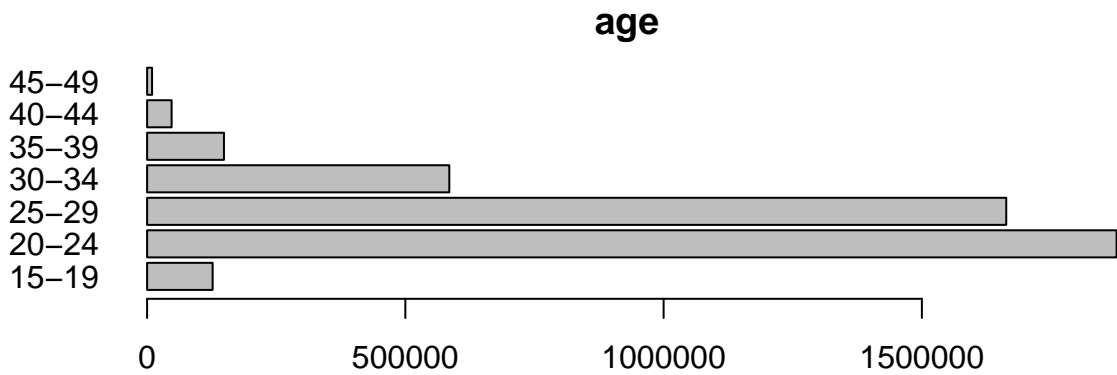


```
births_un <- bdefdata::china_births_un %>%
  Counts()
summary(births_un)
```

```
##
## name:      age      time
## length:   7        5
## dimtype:  age      time
## dimscale: Intervals Intervals
## first:    15-19    1991-1995
## last:     45-49    2011-2015
```

```
##
##   Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##  1330  13835   39630  127373  292470  487940
```

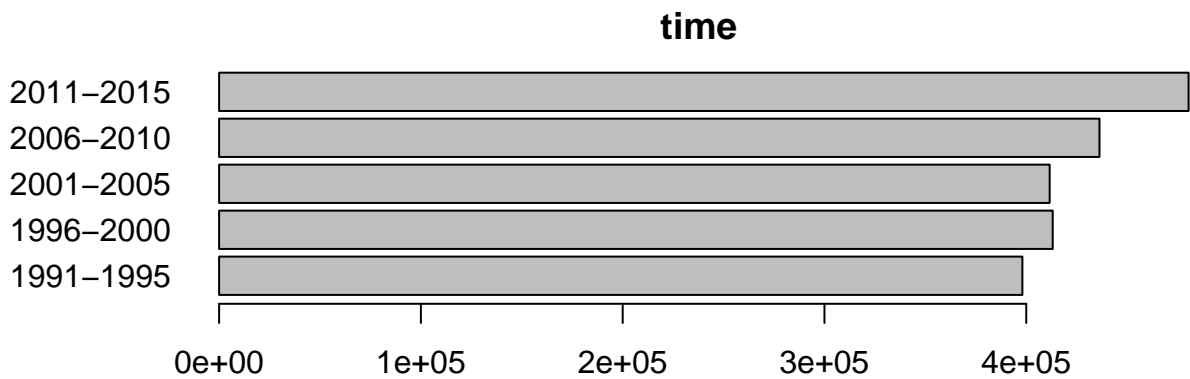
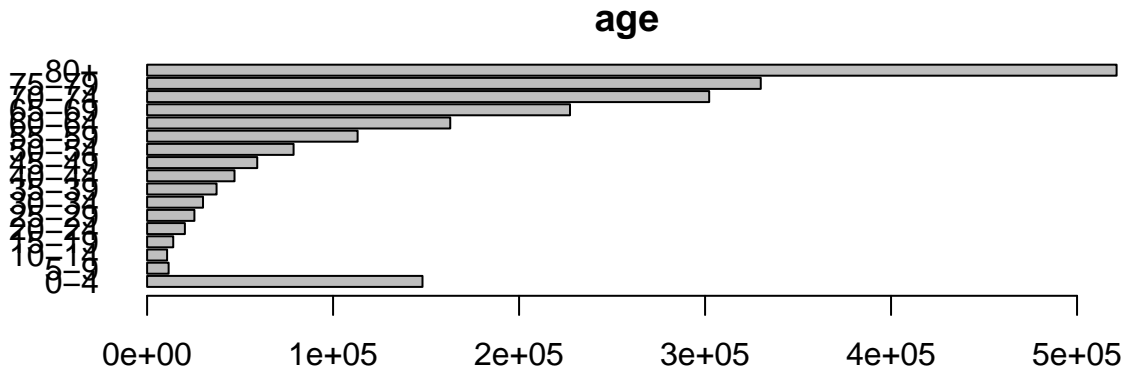
```
plot(births_un)
```



```
deaths_un <- bdefdata::china_deaths_un %>%
  Counts()
summary(deaths_un)
```

```
##
## name:      age      time
## length:   17      5
## dimtype:  age      time
## dimscales: Intervals Intervals
## first:    0-4      1991-1995
## last:     80+     2011-2015
##
##   Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
##   1000  4740   11990  25171  40510  141700
```

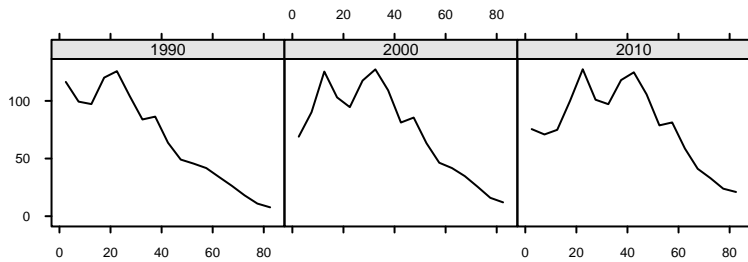
```
plot(deaths_un)
```



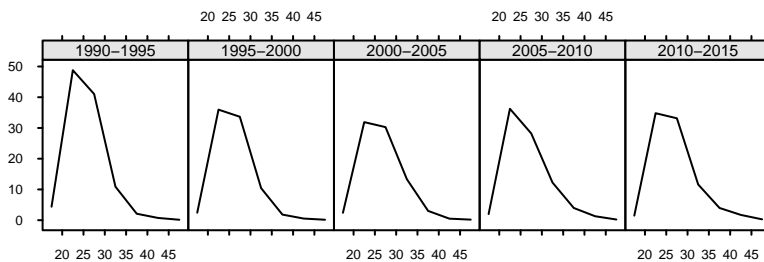
Plot data

```
#Plot census counts of population (millions), by age.
denom_china <- 100
census_df <- census %>%
  as.data.frame(direction = "long", midpoints = "age") %>%
  mutate(count = 1e-6 * denom_china * count)

xyplot(count ~ age | factor(time),
  data = census_df,
  col = "black",
  type = "l",
  layout = c(NA, 1),
  xlab = "",
  ylab = "",
  prepanel = function(y) list(ylim = c(0, max(y, na.rm = TRUE))),
  scales = list(tck = 0.4),
  as.table = TRUE,
  par.settings = list(fontsize = list(text = 6),
    strip.background = list(col = "grey90")))
```



```
#Plot estimates of births (millions), by age of mother.
births_df <- births_un %>%
  as.data.frame(direction = "long", midpoints = "age") %>%
  mutate(count = 1e-6 * denom_china * count) %>%
  mutate(time = gsub("1\\-", "0-", time),
         time = gsub("6\\-", "5-", time))
xyplot(count ~ age | time,
       data = births_df,
       col = "black",
       xlab = "",
       ylab = "",
       type = "l",
       layout = c(NA, 1),
       scales = list(tck = 0.3),
       as.table = TRUE,
       par.settings = list(fontsize = list(text = 6),
                          strip.background = list(col = "grey90")))
```



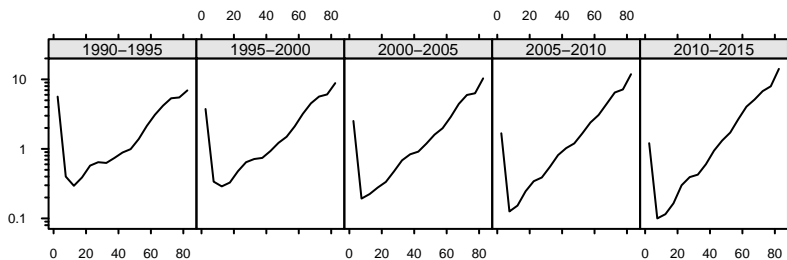
```
#Plot estimates of deaths (millions), by age. The data are shown on a log scale.
deaths_df <- deaths_un %>%
  as.data.frame(direction = "long", midpoints = "age") %>%
  mutate(count = 1e-6 * denom_china * count) %>%
  mutate(time = gsub("1\\-", "0-", time),
         time = gsub("6\\-", "5-", time))

lty <- c("solid", "dotted")
xyplot(count ~ age | time,
       data = deaths_df,
       col = "black",
       type = "l",
       lty = lty,
       xlab = "",
```

```

ylab = "",
lwd = 1,
midpoints = "age",
layout = c(NA, 1),
scales = list(x = list(tck = 0.3),
              y = list(log = TRUE, tck = 0.45)),
yscale.components = yscale.components.log10ticks,
as.table = TRUE,
par.settings = list(fontsize = list(text = 6),
                    strip.background = list(col = "grey90")))

```



```

#Plot changes in cohort size (millions), as implied by the births and census data.
census_df <- census %>%
  as.data.frame(direction = "long", midpoints = "age") %>%
  mutate(cohort = time - age) %>%
  mutate(count = 1e-6 * 100 * count)

births_df <- births_un %>%
  collapseDimension(dimension="age") %>%
  as.data.frame(direction = "long") %>%
  mutate(count = 1e-6 * 100 * count) %>%
  mutate(time = c(1992.5, 1997.5, 2002.5, 2007.5, 2012.5)) %>%
  mutate(cohort = time)

cohorts <- rbind(census_df[census_df$cohort>=1932.5,2:4],births_df[births_df$cohort<=2007.5,]) %>%
  mutate(cohort.name = sprintf("%s-%s",round(cohort-1.5),round(cohort+2.5))) %>%
  arrange(cohort.name,time)

cohorts.births <- cohorts
cohorts.births$count[cohorts.births$time != cohorts.births$cohort] <- NA

p <- xyplot(count ~ time | factor(cohort.name),
            data = cohorts,
            subset = cohort>=1952.5,
            type = "b",
            col = "black",
            xlab = "",
            ylab = "",
            xlim = c(1987, 2013),
            ylim = c(0,NA),
            as.table = TRUE,
            layout = c(NA,2),
            scales = list(tck = 0.3),

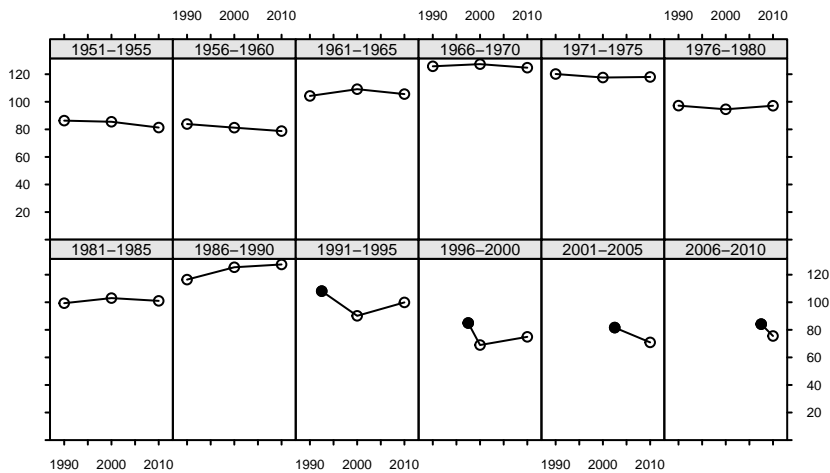
```

```

par.settings = list(strip.background = list(col = "grey90"),
                    fontsize = list(points=6, text=6))
p <- p + as.layer(xyplot(count ~ time | factor(cohort.name),
                      data = cohorts.births,
                      subset = cohort >= 1952.5,
                      type = "p",
                      pch = 19,
                      col = "black"))

plot(p)

```



Fit model

```

filename <- "china_model.est"
if(!file.exists(filename)) {

  #Initial account
  population <- dbind(extrapolate(subarray(census, time == "1990", drop = FALSE),
                                labels = "1995"),
                    extrapolate(subarray(census, time == "2000", drop = FALSE),
                                labels = "2005"),
                    extrapolate(subarray(census, time == "2010", drop = FALSE),
                                labels = c("2015", "2020")),
                    along = "time")

  births <- extrapolate(births_un, labels = "2016-2020")
  deaths <- extrapolate(deaths_un, labels = "2016-2020")

  account <- Movements(population = population,
                      births = births,
                      exits = list(deaths = deaths)) %>%
    makeConsistent()

  #Get datasets.

```

```

datasets <- list(census = census,
                births_un = births_un,
                deaths_un = deaths_un)

#Set up data models.
mean <- census / census
sd <- 0.005 * as(census, "Values")

census_dmodel <- Model(census ~ NormalFixed(mean = mean, sd = sd),
                      series = "population")

births_un_dmodel <- Model(births_un ~ Poisson(mean ~ 1),
                          `(Intercept)` ~ ExchFixed(sd = 0.1),
                          series = "births",
                          priorSD = HalfT(scale = 0.1),
                          jump = 0.01)

deaths_un_dmodel <- Model(deaths_un ~ Poisson(mean ~ 1),
                          `(Intercept)` ~ ExchFixed(sd = 0.1),
                          series = "deaths",
                          priorSD = HalfT(scale = 0.1),
                          jump = 0.03)

dataModels <- list(census_dmodel,
                  births_un_dmodel,
                  deaths_un_dmodel)

#Set up system models
population <- Model(population ~ Poisson(mean ~ age * time,
                                          useExpose = FALSE),
                   age ~ DLM(damp = NULL),
                   time ~ DLM(damp = NULL),
                   age:time ~ DLM(damp = NULL),
                   jump = 0.005)

births <- Model(births ~ Poisson(mean ~ age * time),
               age ~ DLM(damp = NULL),
               time ~ DLM(level = Level(scale = HalfT(scale = 0.1)),
                           trend = NULL,
                           damp = NULL),
               age:time ~ DLM(level = Level(scale = HalfT(scale = 0.05)),
                               trend = NULL,
                               damp = NULL),
               upper = 0.1,
               jump = 0.015)

deaths <- Model(deaths ~ Poisson(mean ~ age * time),
               age ~ DLM(damp = NULL),
               time ~ DLM(level = Level(scale = HalfT(scale = 0.1)),
                           trend = Trend(scale = HalfT(scale = 0.1)),
                           damp = NULL),
               age:time ~ DLM(level = Level(scale = HalfT(scale = 0.05)),
                               trend = NULL,

```



```

                                damp = NULL),
                                upper = 1,
                                jump = 0.03)

systemModels <- list(population,
                     births,
                     deaths)

#Estimate account.
n_burnin <- 700000
n_sim <- 800000
n_chain <- 4
n_thin <- 400

set.seed(0)

estimateAccount(account = account,
                datasets = datasets,
                dataModels = dataModels,
                systemModels = systemModels,
                filename = filename,
                nBurnin = n_burnin,
                nSim = n_sim,
                nChain = n_chain,
                nThin = n_thin)

options(width = 150)
}
fetchSummary(filename)

## -----
## Account:
## *population*
## dimensions: age, time
## n cells: 119
## *births*
## dimensions: age, time, triangle
## n cells: 84
## *deaths*
## dimensions: age, time, triangle
## n cells: 204
## -----
## System models:
## *population*
## population ~ Poisson(mean ~ age * time, useExpose = FALSE),
## age ~ DLM(damp = NULL),
## time ~ DLM(damp = NULL),
## age:time ~ DLM(damp = NULL),
## 0.005
## dimensions: age, time
## *births*
## births ~ Poisson(mean ~ age * time),
## age ~ DLM(damp = NULL),
## time ~ DLM(level = Level(scale = HalfT(scale = 0.1)), trend = NULL,

```

```

##      damp = NULL),
## age:time ~ DLM(level = Level(scale = HalfT(scale = 0.05)), trend = NULL,
##      damp = NULL),
## 0.1,
## 0.015
## dimensions: age, time, triangle
## *deaths*
## deaths ~ Poisson(mean ~ age * time),
## age ~ DLM(damp = NULL),
## time ~ DLM(level = Level(scale = HalfT(scale = 0.1)), trend = Trend(scale = HalfT(scale = 0.1)),
##      damp = NULL),
## age:time ~ DLM(level = Level(scale = HalfT(scale = 0.05)), trend = NULL,
##      damp = NULL),
## 1,
## 0.03
## dimensions: age, time, triangle
## -----
## Datasets:
## *census*
## Object of class "Counts"
## dimensions: age, time
## n cells: 51, n missing: 0, integers: TRUE, n zeros: 0, median: 787532
## *births_un*
## Object of class "Counts"
## dimensions: age, time
## n cells: 35, n missing: 0, integers: TRUE, n zeros: 0, median: 39630
## *deaths_un*
## Object of class "Counts"
## dimensions: age, time
## n cells: 85, n missing: 0, integers: TRUE, n zeros: 0, median: 11990
## -----
## Data models:
## *census*
## census ~ NormalFixed(mean = mean, sd = sd),
## "population"
## dimensions: age, time
## *births_un*
## births_un ~ Poisson(mean ~ 1),
## `(Intercept)` ~ ExchFixed(sd = 0.1),
## HalfT(scale = 0.1),
## 0.01,
## "births"
## dimensions: age, time
## *deaths_un*
## deaths_un ~ Poisson(mean ~ 1),
## `(Intercept)` ~ ExchFixed(sd = 0.1),
## HalfT(scale = 0.1),
## 0.03,
## "deaths"
## dimensions: age, time
## -----
## MCMC statistics:
## nBurnin: 700000, nSim: 800000, nChain: 4, nThin: 400, nIteration: 8000
##

```

```

## Metropolis-Hastings updates:
##
##           jump acceptance autocorr
## systemModels.population.likelihood.count 0.005    0.274    0.347
## systemModels.births.likelihood.rate      0.015    0.308    0.864
## systemModels.deaths.likelihood.rate      0.030    0.308    0.582
## dataModels.births_un.likelihood.rate     0.010    0.320    0.681
## dataModels.deaths_un.likelihood.rate     0.030    0.253    0.623
##
## parameters:
##
##           Rhat      2.5%      50%
## account.population . 1.10    113260  778270
## account.births     . 1.24      610    14929
## account.deaths     . 1.18      484    4920
## systemModels.population.likelihood.count . 1.16    113685  778068
## systemModels.population.prior.mean      1.08     11.7    13.6
## systemModels.population.hyper.age.scaleLevel 1.00    0.0551  0.451
## systemModels.population.hyper.age.scaleTrend 1.01    0.0175  0.246
## systemModels.population.hyper.age.scaleError 1.00    0.0428  0.49
## systemModels.population.hyper.time.scaleLevel 1.00    0.00159 0.0209
## systemModels.population.hyper.time.scaleTrend 1.00    0.00619 0.0226
## systemModels.population.hyper.time.scaleError 1.00    0.000816 0.0134
## systemModels.population.hyper.age:time.scaleLevel 1.00    0.0461  0.107
## systemModels.population.hyper.age:time.scaleTrend 1.00    0.00695 0.0264
## systemModels.population.hyper.age:time.scaleError 1.01    0.0279  0.0835
## systemModels.births.likelihood.rate      . 1.14    0.000334 0.00547
## systemModels.births.prior.mean          . 1.14     -7.98   -5.25
## systemModels.births.prior.sd            . 1.17     0.0443  0.16
## systemModels.births.hyper.age.scaleLevel 1.01     0.103  0.731
## systemModels.births.hyper.age.scaleTrend 1.01     0.0594  0.503
## systemModels.births.hyper.age.scaleError 1.02     0.00805 0.576
## systemModels.births.hyper.time.scaleLevel 1.00     0.00642 0.0696
## systemModels.births.hyper.time.scaleError 1.01     0.0568  0.163
## systemModels.births.hyper.age:time.scaleLevel 1.04     0.00603 0.109
## systemModels.births.hyper.age:time.scaleError 1.00     0.0029  0.0357
## systemModels.deaths.likelihood.rate      . 1.19    0.000269 0.00231
## systemModels.deaths.prior.mean          . 1.19     -8.21   -6.08
## systemModels.deaths.prior.sd            1.06     0.0074  0.0612
## systemModels.deaths.hyper.age.scaleLevel 1.08     0.065  0.937
## systemModels.deaths.hyper.age.scaleTrend 1.05     0.0489  0.411
## systemModels.deaths.hyper.age.scaleError . 1.19     0.0982  0.706
## systemModels.deaths.hyper.time.scaleLevel 1.00     0.00532 0.049
## systemModels.deaths.hyper.time.scaleTrend 1.00     0.00854 0.0511
## systemModels.deaths.hyper.time.scaleError 1.02     0.00124 0.0288
## systemModels.deaths.hyper.age:time.scaleLevel 1.00     0.00442 0.0202
## systemModels.deaths.hyper.age:time.scaleError 1.00     0.00383 0.0155
## dataModels.births_un.likelihood.rate     1.09     0.983  1.14
## dataModels.births_un.prior.mean          1.01     0.104  0.127
## dataModels.births_un.prior.sd            1.05     0.0245  0.0546
## dataModels.deaths_un.likelihood.rate     1.03     0.954  1.17
## dataModels.deaths_un.prior.mean          1.01     0.104  0.166
## dataModels.deaths_un.prior.sd            1.02     0.0985  0.123
##
##           97.5% length
## account.population 1256804    119
## account.births     201329     84

```

```

## account.deaths                    54006    204
## systemModels.population.likelihood.count 1257200 119
## systemModels.population.prior.mean      14    119
## systemModels.population.hyper.age.scaleLevel 1.59    1
## systemModels.population.hyper.age.scaleTrend 1.32    1
## systemModels.population.hyper.age.scaleError 1.32    1
## systemModels.population.hyper.time.scaleLevel 0.186    1
## systemModels.population.hyper.time.scaleTrend 0.131    1
## systemModels.population.hyper.time.scaleError 0.168    1
## systemModels.population.hyper.age:time.scaleLevel 0.169    1
## systemModels.population.hyper.age:time.scaleTrend 0.0623    1
## systemModels.population.hyper.age:time.scaleError 0.505    1
## systemModels.births.likelihood.rate    0.0725    84
## systemModels.births.prior.mean        -2.71    84
## systemModels.births.prior.sd          0.357    1
## systemModels.births.hyper.age.scaleLevel 1.94    1
## systemModels.births.hyper.age.scaleTrend 1.74    1
## systemModels.births.hyper.age.scaleError 1.8    1
## systemModels.births.hyper.time.scaleLevel 0.198    1
## systemModels.births.hyper.time.scaleError 0.35    1
## systemModels.births.hyper.age:time.scaleLevel 0.208    1
## systemModels.births.hyper.age:time.scaleError 0.152    1
## systemModels.deaths.likelihood.rate    0.137    204
## systemModels.deaths.prior.mean        -2.02    204
## systemModels.deaths.prior.sd          0.136    1
## systemModels.deaths.hyper.age.scaleLevel 2.01    1
## systemModels.deaths.hyper.age.scaleTrend 1.23    1
## systemModels.deaths.hyper.age.scaleError 1.84    1
## systemModels.deaths.hyper.time.scaleLevel 0.15    1
## systemModels.deaths.hyper.time.scaleTrend 0.187    1
## systemModels.deaths.hyper.time.scaleError 0.251    1
## systemModels.deaths.hyper.age:time.scaleLevel 0.0619    1
## systemModels.deaths.hyper.age:time.scaleError 0.0582    1
## dataModels.births_un.likelihood.rate    1.35    35
## dataModels.births_un.prior.mean        0.183    35
## dataModels.births_un.prior.sd          0.156    1
## dataModels.deaths_un.likelihood.rate    1.62    85
## dataModels.deaths_un.prior.mean        0.225    85
## dataModels.deaths_un.prior.sd          0.162    1
## -----

```

Results

```

#Population counts
population <- fetch(filename,
                    where = c("account", "population"))
population <- 1e-6 * denom_china * population

census_data <- fetch(filename,
                    where = c("datasets", "census"))
census_data <- 1e-6 * denom_china * census_data

p <- dplot(~ age | factor(time),

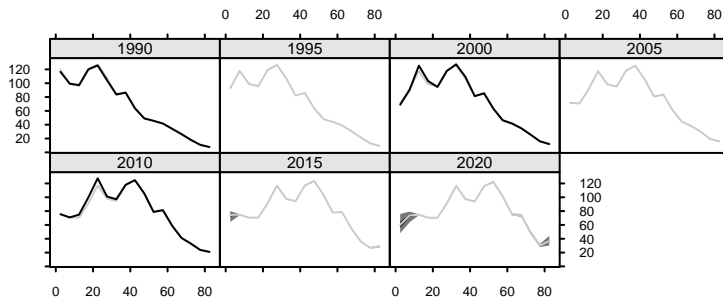
```

```

data = population,
col = "grey20",
xlab = "",
ylab = "",
midpoints = "age",
as.table = TRUE,
scales = list(tck = 0.3),
par.settings = list(fontsize = list(text = 6),
                    strip.background = list(col = "grey90")),
overlay = list(values = census_data,
               col = "black")

```

```
plot(p)
```



```

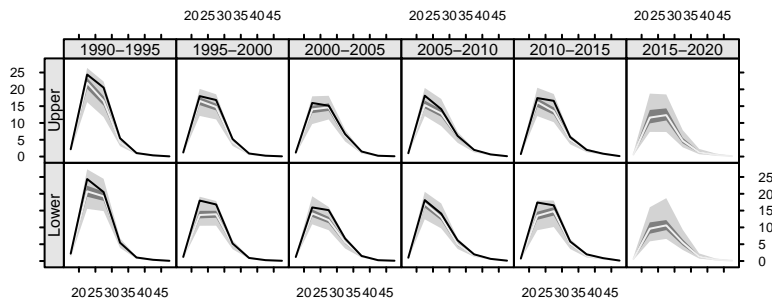
#Birth counts
period_labels <- paste(seq(1990, 2015, 5), seq(1995, 2020, 5), sep = "--")

births <- 1e-6 * denom_china * fetch(filename,
                                     where = c("account", "births")) %>%
  as("array")
dimnames(births)[[3]] <- c("Lower", "Upper")
births <- Counts(births)
births <- births[,2:1,]
births_data <- 0.5 * 1e-6 * denom_china * fetch(filename,
                                                where = c("datasets", "births"))

p <- dplot(~ age | factor(time) * triangle,
           data = births,
           weights = weights,
           col = "grey",
           xlab = "",
           ylab = "",
           midpoints = "age",
           as.table = TRUE,
           scales = list(tck = 0.3),
           strip = strip.custom(factor.levels = period_labels),
           par.settings = list(fontsize = list(text = 6),
                               strip.background = list(col = "grey90")),
           overlay = list(values = births_data,
                          col = "black",
                          type = "l"))

p <- useOuterStrips(p)
p$condlevels[[1]] <- period_labels
plot(p)

```



```

#Death counts
period_labels <- paste(seq(1990, 2015, 5), seq(1995, 2020, 5), sep = "--")

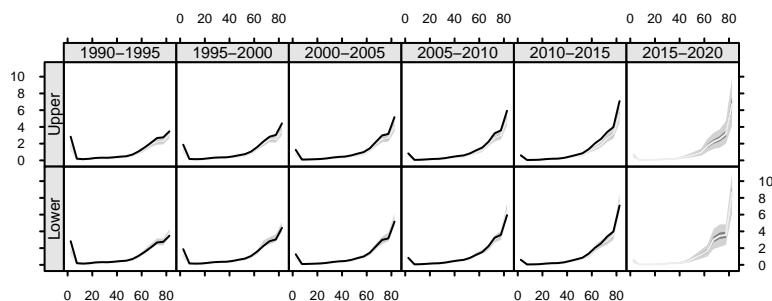
deaths <- (1e-6 * denom_china * fetch(filename,
                                     where = c("account", "deaths"))) %>%
  as("array")
dimnames(deaths)[[3]] <- c("Lower", "Upper")
deaths <- deaths[,2:1,]
deaths <- Counts(deaths)

deaths_data <- (0.5 * 1e-6 * denom_china * fetch(filename,
                                                  where = c("datasets", "deaths_un")))

p <- dplot(~ age | factor(time) * triangle,
           data = deaths,
           col = "grey",
           xlab = "",
           ylab = "",
           midpoints = "age",
           as.table = TRUE,
           scales = list(tck = 0.3),
           par.settings = list(fontsize = list(text = 6),
                               strip.background = list(col = "grey90")),
           overlay = list(values = deaths_data,
                          col = "black"))

p <- useOuterStrips(p)
p$condlevels[[1]] <- period_labels
plot(p)

```



```

#Change in cohort size
period_labels <- paste(seq(1950, 2005, 5), seq(1955, 2010, 5), sep = "--")

population <- fetch(filename,
                    where = c("account", "population"))
births <- fetch(filename,
                where = c("account", "births")) %>%
  collapseDimension(dimension = c("age", "triangle"))

population.pmean <- collapseIterations(population, FUN = mean) %>%
  as.data.frame(direction = "long", midpoints = "age") %>%
  mutate(cohort = time - age) %>%
  mutate(count = 1e-6 * denom_china * count)

births.pmean <- collapseIterations(births, FUN = mean) %>%
  as.data.frame(direction = "long") %>%
  mutate(time = c(1992.5, 1997.5, 2002.5, 2007.5, 2012.5, 2017.5)) %>%
  mutate(cohort = time) %>%
  mutate(count = 1e-6 * denom_china * count)

cohorts.pmean <- population.pmean %>%
  filter(cohort >= 1932.5 & cohort <= 2007.5) %>%
  filter(time %in% c(1990, 2000, 2010)) %>%
  bind_rows(filter(births.pmean, cohort <= 2007.5)) %>%
  select(-age) %>%
  mutate(cohort.name = sprintf("%s-%s",round(cohort-1.5),round(cohort+2.5))) %>%
  arrange(cohort.name, time)

cohorts.pmean.births <- cohorts.pmean %>%
  mutate(count = ifelse(time == cohort, count, NA))

census_df <- census %>%
  as.data.frame(direction = "long", midpoints = "age") %>%
  mutate(cohort = time - age) %>%
  mutate(count = 1e-6 * denom_china * count)

births_df <- births_un %>%
  collapseDimension(dimension = "age") %>%
  as.data.frame(direction = "long") %>%
  mutate(count = 1e-6 * denom_china * count) %>%
  mutate(time = c(1992.5, 1997.5, 2002.5, 2007.5, 2012.5)) %>%
  mutate(cohort = time)

cohorts <- census_df %>%
  filter(cohort >= 1932.5) %>%
  select(time, cohort, count) %>%
  bind_rows(filter(births_df, cohort <= 2007.5)) %>%
  mutate(cohort.name = sprintf("%s-%s",round(cohort-1.5),round(cohort+2.5))) %>%
  arrange(cohort.name,time)
cohorts.births <- cohorts %>%
  mutate(count = ifelse(time == cohort, count, NA))

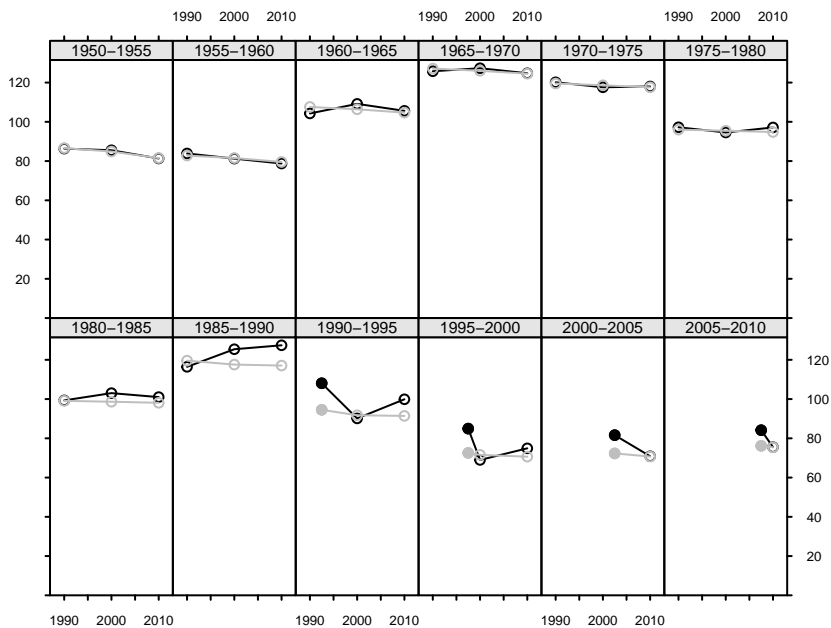
p <- xyplot(count ~ time | cohort.name,

```

```

data = cohorts,
subset = cohort >= 1952.5,
type = "b",
col = "black",
ylim = c(0, NA),
xlab = "",
ylab = "",
xlim = c(1987, 2013),
as.table = TRUE,
layout = c(NA,2),
scales = list(tck = 0.3),
par.settings = list(strip.background = list(col = "grey90"),
                    fontsize = list(points=6, text=6))
p <- p + as.layer(xyplot(count ~ time | cohort.name,
                        data = cohorts.births,
                        subset = cohort>=1952.5,
                        type = "p",
                        pch = 19,
                        col = "black"))
p <- p + as.layer(xyplot(count ~ time | cohort.name,
                        data = cohorts.pmean,
                        subset = cohort>=1952.5,
                        type = "b",
                        col = "grey"))
p <- p + as.layer(xyplot(count ~ time | cohort.name,
                        data = cohorts.pmean.births,
                        subset = cohort>=1952.5,
                        type = "p",
                        pch = 19,
                        col = "grey"))
p$condlevels[[1]] <- period_labels
plot(p)

```

#Death rates

```

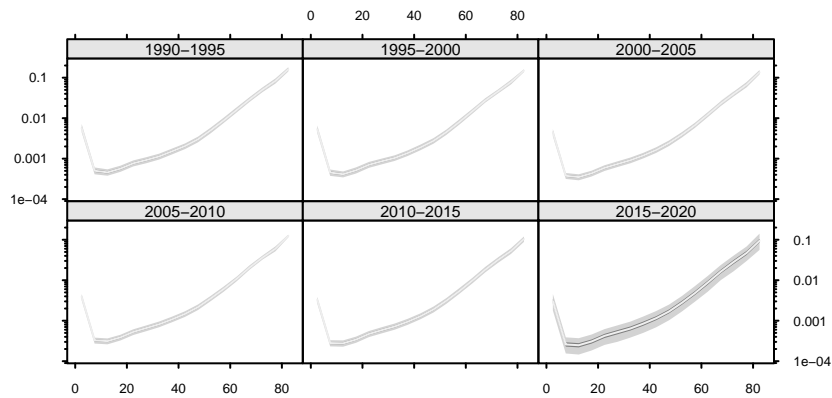
period_labels <- paste(seq(1990, 2015, 5), seq(1995, 2020, 5), sep = "--")

rates <- fetch(filename,
               where = c("systemModels", "deaths", "likelihood", "rate"))
weights <- fetch(filename,
                 where = c("account", "population")) %>%
  exposure(triangles = TRUE)

p <- dplot(~ age | factor(time),
           data = rates,
           weights = weights,
           col = "grey",
           xlab = "",
           ylab = "",
           midpoints = "age",
           as.table = TRUE,
           scales = list(tck = 0.3,
                         y = list(log = TRUE)),
           yscale.components = yscale.components.log10ticks,
           par.settings = list(fontsize = list(text = 6),
                               strip.background = list(col = "grey90")))
p$condlevels[[1]] <- period_labels

plot(p)

```



```

#Life expectancy
period_labels <- paste(seq(1990, 2015, 5), seq(1995, 2020, 5), sep = "-")

rates <- fetch(filename,
               where = c("systemModels", "deaths", "likelihood", "rate"))
weights <- fetch(filename,
                 where = c("account", "population")) %>%
  exposure(triangles = TRUE)

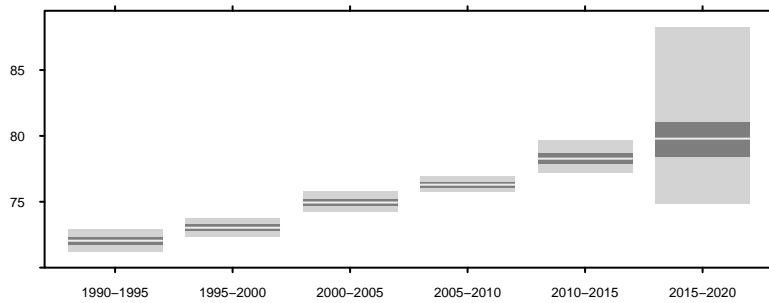
lifeexp <- rates %>%
  collapseDimension(dimension="triangle", weights=weights) %>%
  LifeTable() %>%
  lifeExpectancy()

lifeexp <- as(lifeexp, "array")
dimnames(lifeexp)[[1]] <- period_labels
lifeexp <- Counts(lifeexp,
                 dimtypes = c(time = "state"))

p <- dplot(~ time,
           data = lifeexp,
           col = "grey",
           xlab = "",
           ylab = "",
           as.table = TRUE,
           scales = list(tck = 0.3),
           par.settings = list(fontsize = list(text = 6),
                               strip.background = list(col = "grey90")))

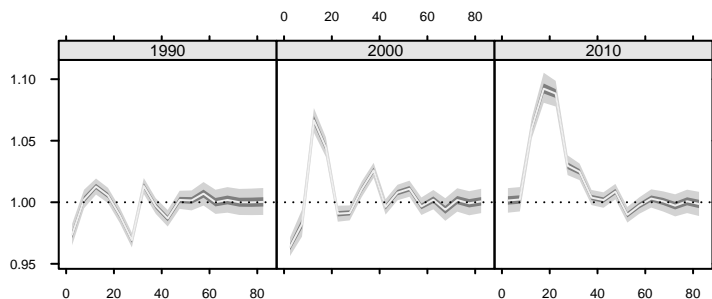
plot(p)

```



```
#Coverage ratio for census counts
coverage <- fetchCoverage(filename, "census")

p <- dplot(~ age | factor(time),
  data = coverage,
  col = "grey",
  xlab = "",
  ylab = "",
  midpoint = "age",
  as.table = TRUE,
  scales = list(x = list(tck = 0.3),
    y = list(tck = 0.5)),
  par.settings = list(fontsize = list(text = 6),
    strip.background = list(col = "grey90")))
p <- p + latticeExtra::layer(panel.abline(h=1,lty="dotted",col="black"))
plot(p)
```



```
#Coverage ratio for UN births estimates
coverage <- fetchCoverage(filename, "births_un") %>%
  subarray(time %in% c("1991-1995", "2001-2005", "2011-2015"))

p <- dplot(~ age | factor(time),
  data = coverage,
  col = "grey",
  xlab = "",
  ylab = "",
  midpoint = "age",
  layout = c(NA,1),
  as.table = TRUE,
  scales = list(x = list(tck = 0.3),
```

```

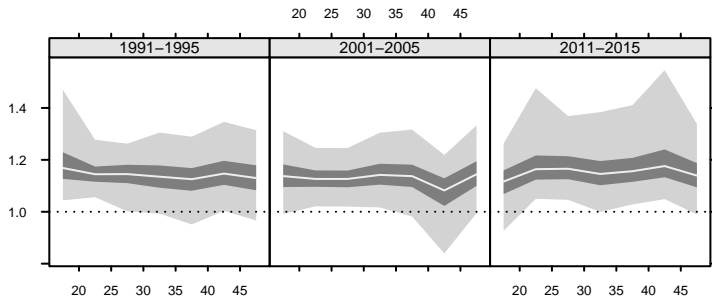
y = list(tck = 0.5)),
par.settings = list(fontsize = list(text = 6),
strip.background = list(col = "grey90"))

```

```

p <- p + latticeExtra::layer(panel.abline(h = 1, lty = "dotted", col = "black"))
plot(p)

```



```

#Coverage ratio for UN deaths estimates
coverage <- fetchCoverage(filename, "deaths_un") %>%
  subarray(time %in% c("1991-1995", "2001-2005", "2011-2015"))

```

```

p <- dplot(~ age | factor(time),
data = coverage,
col = "grey",
xlab = "",
ylab = "",
midpoint = "age",
layout = c(NA,1),
as.table = TRUE,
scales = list(x = list(tck = 0.3),
y = list(tck = 0.5)),
par.settings = list(fontsize = list(text = 6),
strip.background = list(col = "grey90"))

```

```

p <- p + latticeExtra::layer(panel.abline(h = 1, lty = "dotted", col = "black"))
plot(p)

```

