## Day 1, exercise 1: Decomposing the crude death rate using Kitagawa decomposition

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- $t_1$  is the initial period and  $t_2$  is the final period
- $D_x$  = number of deaths at age x
- $M_x = \text{death rate}$
- $N_x$  = is the mid-year population
- N = total population over ages

Note that  $D_x = M_x * N_x$  and that the difference between two crude death rates can be expressed as

$$\Delta CDR = \sum_{x} M_x(t_2) \frac{N_x(t_2)}{N(t_2)} - \sum_{x} M_x(t_1) \frac{N_x(t_1)}{N(t_1)}.$$
 (1)

The aim is to disentangle the effects of changing rates and the effect of the change in the composition of the population.

The file Kitagawa\_Data.RData contains mortality and exposure data for Israel in 1990 and 2016. The exercise consists in decomposing the changes in CDR following Kitagawa.

Start by loading the data.

```
load('Kitagawa_Data.RData')
```

The exercises in this course will use the package tidyverse, which you should have already met with Tim Riffe.

We need to load this package too (and install it, if you haven't already).

```
#install.packages("tidyverse") #to install the package
library(tidyverse)
```

```
## -- Attaching packages ----- tidyverse 1.3.2 --
## v ggplot2 3.4.0
                  v purrr
                           0.3.5
## v tibble 3.1.8
                   v dplyr
                           1.0.10
## v tidyr
         1.2.1
                   v stringr 1.4.1
## v readr
         2.1.3
                   v forcats 0.5.2
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                masks stats::lag()
```

Period 1 corresponds to 1990 and period 2 to 2016 in this case. Start by selecting deaths and exposures (approximated usually by the midyear population) for each of the periods and store them in vectors.

```
# time period 1
# Select deaths for first period
Dx1 <- Deaths %>%
  filter(Year==1990) %>%
  pull(Total)
# Select population for first period
Nx1 <- Exposures %>%
```

```
filter(Year==1990) %>%
  pull(Total)
# Do the same for period 2
Dx2 <- Deaths%>%
  filter(Year==2016) %>%
  pull(Total)
Nx2 <- Exposures%>%
  filter(Year==2016) %>%
  pull(Total)
```

We can calculate the crude death rates directly from these.

```
# get the crude death rate in time 1
CDR1 <- sum(Dx1)/sum(Nx1)
#crude death expressed by 1000 in time 1
CDR1*1000
## [1] 6.686454
# the same for period 2
CDR2 <- sum(Dx2)/sum(Nx2)
#crude death expressed by 1000 in time 2
CDR2*1000
## [1] 6.882632
#change in CDR
(Dif <- (CDR2 - CDR1)*1000)</pre>
```

## ## [1] 0.1961774

What would we have expected in this period? what do the CDRs suggest? Does it make sense?

Let's decompose and see if effectively mortality went up or if it is a compositional effect.

Remember Kitagawa's formula:

$$\Delta CDR = \underbrace{\sum_{x} \left( \frac{M_x(t_2) + M_x(t_1)}{2} \right) \left( \frac{N_x(t_2)}{N(t_2)} - \frac{N_x(t_1)}{N(t_1)} \right)}_{\text{Changes in x-composition}} + \underbrace{\sum_{x} \left( \frac{N_x(t_2)}{N(t_2)} + \frac{N_x(t_1)}{N(t_1)} \right) \left( M_x(t_2) - M_x(t_1) \right)}_{\text{Changes in rates}}$$

$$(2)$$

We need to calculate the age-specific mortality rates, which are the deaths divided by the exposure for both periods.

```
# time period 1
# Select deaths for first period
# get age-specific mortality rates
Mx1 <- Dx1/Nx1
# replace NA's with zero (just for simplicity)
Mx1 <- Mx1 %>%
    replace_na(replace=0)
# do the same for the second period
```

```
Mx2 <- Dx2/Nx2
Mx2 <- Mx2 %>%
  replace_na(replace=0)
```

Now let's calculate Kitagawa's components.

```
RC <- sum(0.5*(Nx2/sum(Nx2) + Nx1/sum(Nx1))*(Mx2-Mx1))
RC*1000

## [1] -1.898981

CC <- sum(0.5*(Mx2+Mx1)*(Nx2/sum(Nx2)-Nx1/sum(Nx1)))
CC*1000
```

```
## [1] 2.095158
```

What do they tell us about changes in mortality between 1990 and 2016? What was the role of population composition vs death rates?

Let us check that the decomposition results fit with the original difference.

```
RC*1000 + CC*1000

## [1] 0.1961774

Dif

## [1] 0.1961774
```