Summative Assessment PDP

echo: false

Install packages

```
# Install necessary packages
# install.packages("tidyverse")
# install.packages("ggplot2")
# install.packages("HMDHFDplus")
# install.packages('MortalityLaws')
# install.packages('DemoDecomp')
```

Load packages

```
# load necessary packages
library(tidyverse)
```

```
— Attaching core tidyverse packages –
                                                            — tidyverse 2.0.0 —
✓ dplyr
           1.1.4
                     ✓ readr
                                 2.1.5
✓ forcats 1.0.0
                                 1.5.1

✓ stringr

✓ ggplot2 3.5.1

✓ tibble

                                 3.2.1
✓ lubridate 1.9.3

✓ tidyr

                                 1.3.1
           1.0.2
✓ purrr
— Conflicts —
                                                      – tidyverse_conflicts() —
* dplyr::filter() masks stats::filter()
                 masks stats::lag()
* dplyr::lag()
i Use the conflicted package (<http://conflicted.r-lib.org/>) to force all conflicts to
become errors
```

```
library(HMDHFDplus)
library(ggplot2)
library(MortalityLaws)
library(DemoDecomp)
library(readxl)
library(stringr)

# don't display in scientific notation
options(scipen = 999)
```

Question 1 - Mortality

Obtain single year LTs for Latvia and Italy from HMD

```
# Obtain 1 year LTs for Latvia and Italy
```

```
# code in comments for anonymity purposes

# LT_females_Latvia <- readHMDweb(CNTRY = "LVA", item = "fltper_1x1", username = "***", p

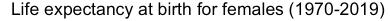
# LT_females_Italy <- readHMDweb(CNTRY = "ITA", item = "fltper_1x1", username = "***", pa</pre>
```

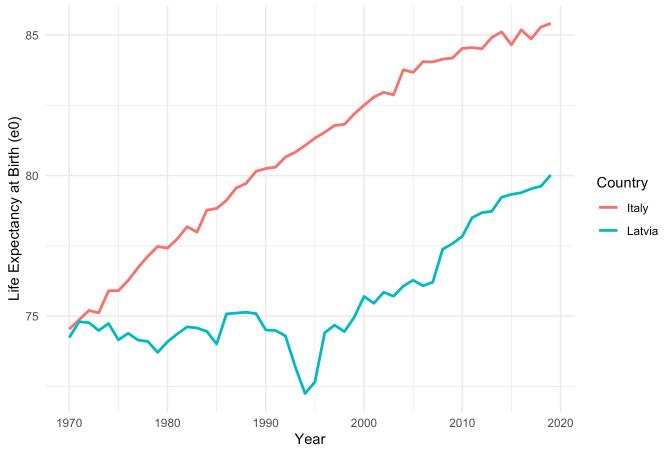
Create data frames that give e0 for both Latvia and Italy from 1970 to 2019. We will also merge these data frames so that we can plot e0 for both countries over time.

```
# Filter data for e0 from 1970 to 2019 - Latvia
e0 females Latvia DF <- LT females Latvia %>%
  filter(Age == 0, Year >= 1970, Year <= 2019) %>%
  select(Year, ex) %>%
  mutate(Country = "Latvia")
# change name of ex column to e0
colnames(e0_females_Latvia_DF) <- c("Year","e0","Country")</pre>
# Filter data for e0 from 1970 to 2019 - Italy
e0_females_Italy_DF <- LT_females_Italy %>%
  filter(Age == 0, Year >= 1970, Year <= 2019) %>%
  select(Year, ex) %>%
  mutate(Country = "Italy")
# change name of ex column to e0
colnames(e0_females_Italy_DF) <- c("Year","e0","Country")</pre>
# Combine both datasets
e0 Latvia and Italy DF <- bind rows(e0 females Latvia DF, e0 females Italy DF)
```

1. Plot e0 for both countries from 1970 - 2019 and describe the results

```
# Plot life expectancy at birth over time
ggplot(e0_Latvia_and_Italy_DF, aes(x = Year, y = e0, color = Country)) +
    geom_line(linewidth = 1) +
    labs(title = "Life expectancy at birth for females (1970-2019)",
        x = "Year",
        y = "Life Expectancy at Birth (e0)",
        color = "Country") +
    theme_minimal()
```





Describe results:

Italy e0 appears to be improving at a constant rate over time. On the other hand, Latvia e0 stagnates somewhat from 1970 - 1990 and then e0 sharply drops in 1990. Around 1995, e0 in Latvia improves back to pre-1990 levels and then improves at a constant rate. It is also worthwhile to note that both Italy and Latvia have similar e0 in 1970, when the first year is specified. However, after 50 years, Latvia e0 in 2019, which is ~80, falls behind Italy by about 5 years, which is ~85.

2. What was the difference in life expectancy at birth between these two countries in 1970, and in 2019?

```
# Latvia
e0_Latvia_1970 <- e0_females_Latvia_DF %>%
filter(Year == 1970) %>%
pull(e0)

e0_Latvia_2019 <- e0_females_Latvia_DF %>%
filter(Year == 2019) %>%
pull(e0)

e0_Latvia_2019 - e0_Latvia_1970 # 5.78 year difference in e0 from 1970 - 2019
```

```
# Italy
e0_Italy_1970 <- e0_females_Italy_DF %>%
    filter(Year == 1970) %>%
    pull(e0)

e0_Italy_2019 <- e0_females_Italy_DF %>%
    filter(Year == 2019) %>%
    pull(e0)

e0_Italy_2019 - e0_Italy_1970 # 10.87 year difference in e0 from 1970 - 2019
```

[1] 10.87

3.

subset the data for both countries in 2019 and for ages from 10 onward

```
# Subset data for year 2019 and ages 10 and on - Latvia
LT_females_Latvia_2019 <- LT_females_Latvia %>%
   filter(Age >= 10, Year == 2019)

# Subset data for year 2019 and ages 10 and on - Italy
LT_females_Italy_2019 <- LT_females_Italy %>%
   filter(Age >= 10, Year == 2019)
```

2). randomly assign NA values to 30 ages

```
# Latvia - Randomly assign NA values to 30 ages
set.seed(123)  # For reproducibility
random_rows <- sample(nrow(LT_females_Latvia_2019), 30)  # Select 30 random row indices

# Assign NA to all columns in the selected rows
LT_females_Latvia_2019[random_rows, ] <- NA

# Italy - Randomly assign NA values to 30 ages
set.seed(123)  # For reproducibility
random_rows <- sample(nrow(LT_females_Italy_2019), 30)  # Select 30 random row indices

# Assign NA to all columns in the selected rows
LT_females_Italy_2019[random_rows, ] <- NA</pre>
```

3). fit a Makeham model for both countries.

Latvia:

```
# first, remove rows with NA values as we did in the first week's HW.
# MortalityLaw() function will not operate properly otherwise.
LT_females_Latvia_2019_clean <- na.omit(LT_females_Latvia_2019)</pre>
# obtain mxs from LT_females_Latvia_2019_clean
mx_values_Latvia <- LT_females_Latvia_2019_clean$mx</pre>
# obtain ages from LT_females_Latvia_2019_clean
ages_Latvia <- LT_females_Latvia_2019_clean$Age</pre>
# create an object that transforms the mx values from LT_females_Latvia_2019_clean into l
log mx values Latvia <- log(mx values Latvia)</pre>
# append this object to the LT females Latvia 2019 clean dataset
LT females Latvia 2019 clean$log mx <- log mx values Latvia
# See available Mortality laws for reference
# availableLaws(law = NULL)
# We can see that the makeham mortality model is typically used to
# estimate adult mortality based on the legend (3 makeham)
# Fit the makeham model, using poisson regression - Latvia
makeham_fitted_Latvia <- MortalityLaw(x = LT_females_Latvia_2019_clean$Age, mx = mx_value
```

Italy:

```
# first, remove rows with NA values
LT_females_Italy_2019_clean <- na.omit(LT_females_Italy_2019)

# obtain mxs from LT_females_Italy_2019_clean
mx_values_Italy <- LT_females_Italy_2019_clean$mx

# obtain ages from LT_females_Italy_2019_clean
ages_Italy <- LT_females_Italy_2019_clean$Age

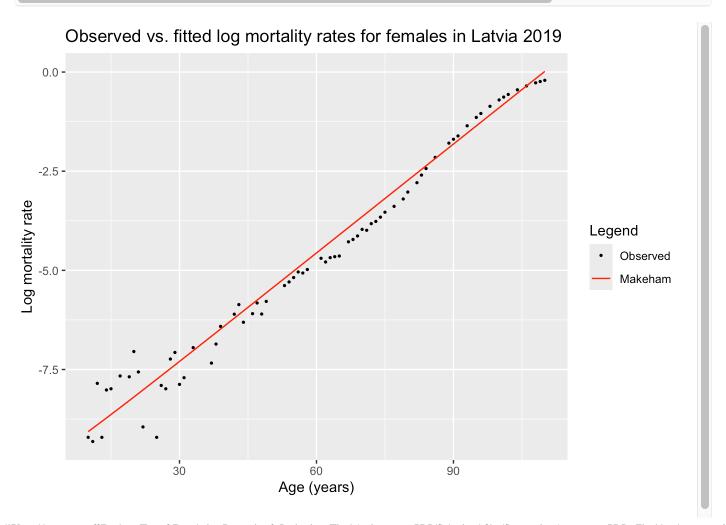
# create an object that transforms the mx values from LT_females_Italy_2019_clean into lo
log_mx_values_Italy <- log(mx_values_Italy)

# append this object to the LT_females_Italy_2019_clean dataset
LT_females_Italy_2019_clean$log_mx <- log_mx_values_Italy
# Fit the makeham model, using poisson regression - Italy
makeham_fitted_Italy <- MortalityLaw(x = LT_females_Italy_2019_clean$Age, mx = mx_values_</pre>
```

4. Plot observed and fitted values, and describe your findings. How good is the fit?

Latvia:

```
# create DF that includes ages, makeham mxs, and log makeham mxs
# will use this DF to plot results
makeham DF Latvia <- data.frame(Age = ages Latvia,</pre>
                                makeham_fitted_Latvia = predict(makeham_fitted_Latvia, x
  mutate(log_makeham_fitted_mx_Latvia = log(makeham_fitted_Latvia))
# Create a plot of the observed log mx rates and the makeham fitted log mx rates
makeham_Latvia <- ggplot() +</pre>
  geom_point(data = LT_females_Latvia_2019_clean, aes(x=Age, y=log_mx, color = "Observed"
  geom\_line(data = makeham\_DF\_Latvia, aes(x = Age, y = log\_makeham\_fitted\_mx\_Latvia, colo
  scale color manual(name = "Legend",
                     values = c("Observed" = "black",
                                 "Makeham" = "red"),
                     limits = c("Observed", "Makeham")) +
  labs(title = "Observed vs. fitted log mortality rates for females in Latvia 2019",
       y = "Log mortality rate",
       x = 'Age (years)') +
  theme(legend.position.inside = c(0.85, 0.25))
makeham_Latvia
```

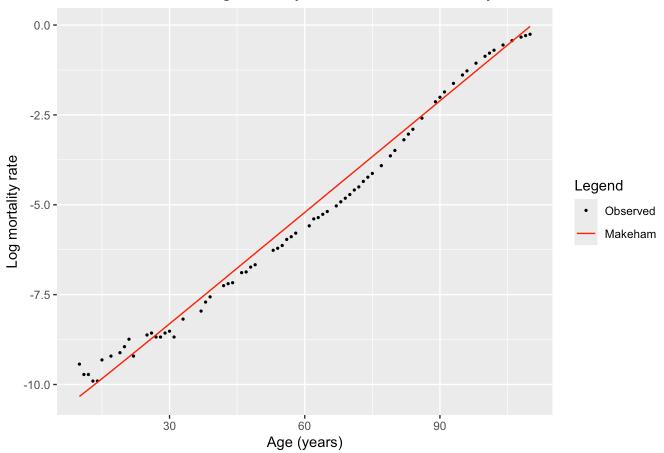


For Latvia, this model fits relatively well for ages 30 and on. However, from ages 60-90, it seems to overestimate the mortality rates, which may be a cause for concern.

Italy:

```
# create DF that includes ages, makeham mxs, and log makeham mxs
makeham DF Italy <- data.frame(Age = ages Italy,</pre>
                               makeham_fitted_Italy = predict(makeham_fitted_Italy, x = a
 mutate(log makeham fitted mx Italy = log(makeham fitted Italy))
# Create a plot of the observed log mx rates and the makeham fitted log mx rates - Italy
makeham Italy <- ggplot() +</pre>
  geom_point(data = LT_females_Italy_2019_clean, aes(x=Age, y=log_mx, color = "Observed")
 geom_line(data = makeham_DF_Italy, aes(x = Age, y = log_makeham_fitted_mx_Italy, color
 scale_color_manual(name = "Legend",
                     values = c("Observed" = "black",
                                "Makeham" = "red"),
                     limits = c("Observed", "Makeham")) +
 labs(title = "Observed vs. fitted log mortality rates for females in Italy 2019",
       y = "Log mortality rate",
       x = 'Age (years)') +
  theme(legend.position.inside = c(0.85, 0.25))
makeham Italy
```

Observed vs. fitted log mortality rates for females in Italy 2019



For Italy, this model fits relatively well, and actually performs better for ages < 30. However, similar to what we saw with Latvia, the model overestimates mortality from ages 45 - 90, which may be a cause for concern.

5. Write down the Makeham model, what does each parameter mean?

 $\mu(x) = A * \exp(Bx) + C$ where,

- $\mu(x)$ is the death rate or force of mortality at age x
- A is a parameter that determines the initial mortality rate
- B is a parameter that determines the rate at which mortality increases with age
- C represents mortality due to factors independent of age (makeham term)
- Note: the Makeham model extends on the Gompertz model, $\mu(x) = A * \exp(Bx)$, by incorporating C, the age-independent mortality component.

6. Say you have data for a country from 1970 to 2019, and you would like to forecast life expectancy to 2040. How would you do it?

To answer this question, I would use the Lee-Carter model (Lecture 9: Forecasting, Slide 4) to forecast life expectancy as we did in the forecasting practical in class on March 14th. The model is specified by

the function: $ln(mx,t) = \alpha x + \beta x \kappa t + \epsilon x, t$ where

- αx is the general shape of the log-mortality at age x
- βx is the rate of mortality improvement at age x
- kt is the general level of mortality at time t
- ex,t is the error with mean 0 and variance σ^2 , e, which reflects the age-specific influences that aren't captured in the model

We first can use the data we have for the country of interest from 1970 to 2019 and model mortality by inputting the data for the relevant parameters. The main parameter of interest in forecasting mortality is κt , as this is the only parameter that contains time in its dimension (error does as well, but we measure this independently). We can identify how κt changes from 1970 to 2019 to see if mortality conditions are improving or worsening over time (Slide 11).

We can then forecast mortality rates for each age group from 2020 to 2040 (20 years) using the κ t values alongside the α x and β x values.

Lastly, through constructing a life table, we can then convert these forecasted mortality rates into forecasted life expectancy values.

Question 2 - Lifespan inequality and decomposition

1. Build a function to calculate the standard deviation from a set of mortality rates.

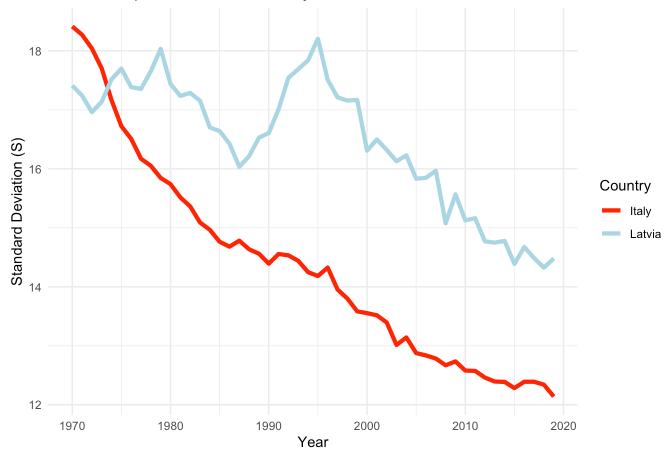
```
sd_function <- function(age, dx, ex, ax, l0) {
  alphax <- age + ax
  eta <- ex[1] +age[1]
  inner <- dx * (alphax - eta)^2
  V <- sum(inner) / l0
  S <- sqrt(V)
  return(S)
}</pre>
```

2. Calculate the standard deviation for both countries from 1970 to 2019. [describe the results, maybe a plot is useful]

```
# Latvia - create a DF that stores e0 and S
results_Latvia <- LT_females_Latvia %>%
  filter(Year >= 1970 & Year <= 2019) %>%
  group_by(Year) %>%
  summarize(
    ex = first(ex),
    ax = list(ax),
```

```
dx = list(dx),
    lx = list(lx),
    S = sd_function(
      age = Age
      dx = dx[[1]],
      ex = ex,
      ax = ax[[1]],
      10 = 100000
    )
  ) %>%
  ungroup() %>%
  select(Year, e0 = ex, S)
# Italy - create a DF that stores e0 and S
results_Italy <- LT_females_Italy %>%
  filter(Year >= 1970 & Year <= 2019) %>%
  group by (Year) %>%
  summarize(
    ex = first(ex),
    ax = list(ax),
    dx = list(dx),
    lx = list(lx),
    S = sd_function(
     age = Age
     dx = dx[[1]],
      ex = ex,
      ax = ax[[1]],
      10 = 100000
    )
  ) %>%
  ungroup() %>%
  select(Year, e0 = ex, S)
# Plot both countries together
# add a country label
results_Italy$Country <- "Italy"</pre>
results Latvia$Country <- "Latvia"
# Merge the two data frames
results_combined <- rbind(results_Italy, results_Latvia)</pre>
# Plot
ggplot(results\_combined, aes(x = Year, y = S, color = Country)) +
  geom\_line(linewidth = 1.5) +
  labs(title = "SD of Lifespan for females in Italy and Latvia 1970-2019",
       x = "Year",
       y = "Standard Deviation (S)") +
  scale_color_manual(values = c("Italy" = "red", "Latvia" = "lightblue")) +
  theme_minimal()
```

SD of Lifespan for females in Italy and Latvia 1970-2019



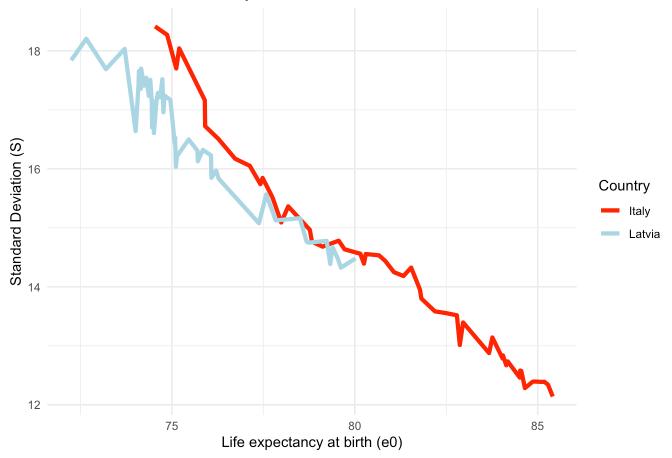
3. Plot life expectancy at birth vs standard deviation, what do you observe?

Simple plot of e0 vs. S

```
# first: a simple plot with e0 on the x-axis and S on the y-axis
# this follows what we have seen in the alternative measures
# of longevity lecture (Slide 27)

ggplot(results_combined, aes(x = e0, y = S, color = Country)) +
geom_line(linewidth = 1.5) +
labs(title = "e0 vs S for females in Italy and Latvia 1970-2019",
        x = "Life expectancy at birth (e0)",
        y = "Standard Deviation (S)") +
scale_color_manual(values = c("Italy" = "red", "Latvia" = "lightblue")) +
theme_minimal()
```

e0 vs S for females in Italy and Latvia 1970-2019



Observations:

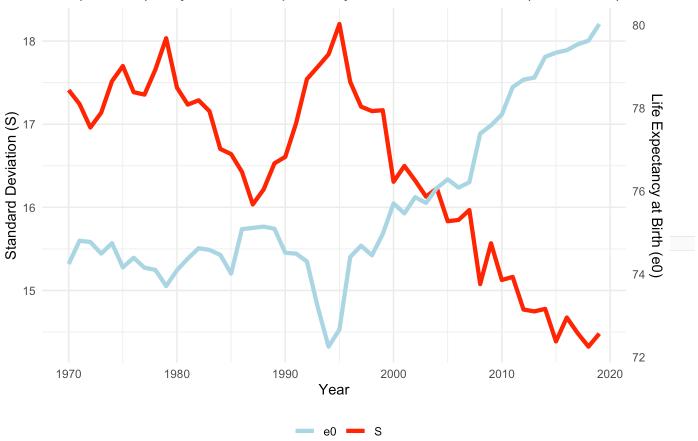
- This plot matches what we saw in the Alternative measures of longevity lecture (Slide 27: Life expectancy and life span inequality (σ)).
- Generally, we see that as life expectancy at birth improves, life span inequality reduces, which is the case in plot above.

Can also be useful to see the relationship between e0 and S by plotting Latvia and Italy individually.

```
# Latvia
# to plot e0 and SD on the y-axis, we need to scale both approrpriately
# scaling factors for e0 to match S's range
min_S_Latvia <- min(results_Latvia$S, na.rm = TRUE)
max_S_Latvia <- max(results_Latvia$S, na.rm = TRUE)
min_e0_Latvia <- min(results_Latvia$e0, na.rm = TRUE)
max_e0_Latvia <- max(results_Latvia$e0, na.rm = TRUE)

# Scale e0 to match S's scale
results_Latvia$e0_scaled <- (results_Latvia$e0 - min_e0_Latvia) / (max_e0_Latvia - min_e0)
# plot e0 vs S
ggplot(results_Latvia, aes(x = Year)) +
geom_line(aes(y = S, color = "S"), linewidth = 1.5) +</pre>
```

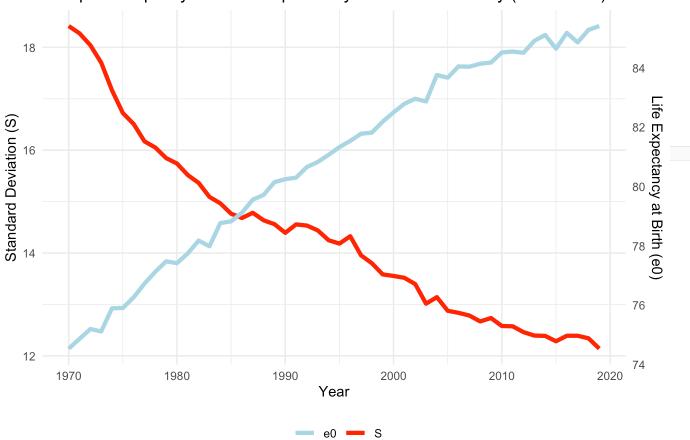
Lifespan Inequality and Life Expectancy for Females in Latvia (1970-2019)



```
# Italy
# scaling factors for e0 to match S's range
min_S_Italy <- min(results_Italy$S, na.rm = TRUE)
max_S_Italy <- max(results_Italy$S, na.rm = TRUE)
min_e0_Italy <- min(results_Italy$e0, na.rm = TRUE)
max_e0_Italy <- max(results_Italy$e0, na.rm = TRUE)

# Scale e0 to match S's scale
results_Italy$e0_scaled <- (results_Italy$e0 - min_e0_Italy) / (max_e0_Italy - min_e0_Italy)</pre>
```

Lifespan Inequality and Life Expectancy for Females in Italy (1970-2019)



4. Using the Horiuchi model for decomposition, decompose the change in lifespan inequality between 2000 and 2019 in both countries by age.

To answer this question, we will need to first define an sd.frommx function in R

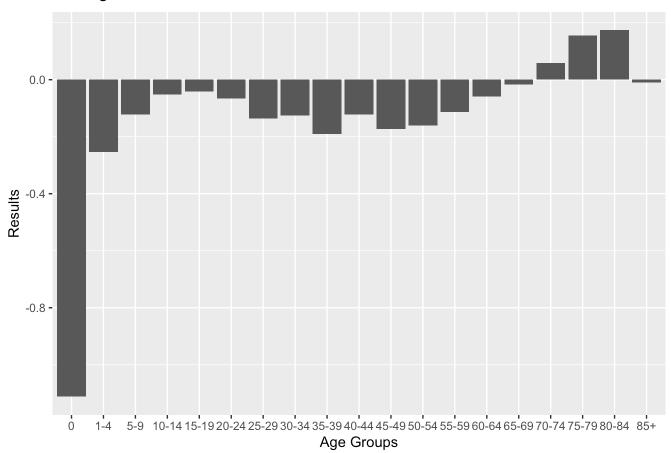
```
# sd.frommx is defined as we saw in the practical on Change in LSV 1996-2023
sd.frommx <- function(nmx = mx, sex=1, age = c(0, 1, seq(5, 85, 5)), nax = NULL){
      <- c(diff(age), 999)
  if (is.null(nax)) {
    nax < -0.5 * n
    if (n[2] == 4) {
      if (sex == 1) {
        if (nmx[1] >= 0.107) {
           nax[1] < -0.33
           nax[2] <- 1.352
        }
        else {
           nax[1] < -0.045 + 2.684 * nmx[1]
           nax[2] <- 1.651 - 2.816 * nmx[1]
        }
      }
      if (sex == 2) {
        if (nmx[1] >= 0.107) {
           nax[1] < -0.35
           nax[2] <- 1.361
        }
        else {
           nax[1] \leftarrow 0.053 + 2.8 * nmx[1]
           nax[2] <- 1.522 - 1.518 * nmx[1]
      }
    }
  }
                <- (n * nmx)/(1 + (n - nax) * nmx)
  nqx
                <- c(nqx[-(length(nqx))], 1)
  nqx
  nqx[nqx > 1] <- 1
  npx <- 1 - nqx
  lx <- cumprod(c(1, npx))</pre>
  ndx <- -diff(lx)
  lxpn \leftarrow lx[-1]
  nLxpn \leftarrow n * lxpn + ndx * nax
  nLx <- c(nLxpn[-length(nLxpn)], lxpn[length(lxpn)-1]/nmx[length(nmx)])</pre>
  Tx <- rev(cumsum(rev(nLx)))</pre>
  lx <- lx[1:length(age)]</pre>
  ex <- Tx/lx
  nax[length(nax)] <- ex[length(ex)]</pre>
  vx \leftarrow sum(ndx*(age+nax-ex[1L])^2)
  sd <- sqrt(vx)</pre>
  return(sd)
}
```

We also need to create an object that gives us labels for age groups

Latvia:

```
# obtain the mortality rates for each age
# Note that the length of these vectors is 111. This is because we
# have mortality rates for each individual age
mx1 Latvia <- LT females Latvia[LT females Latvia$Year == 2000,]$mx
mx2 Latvia <- LT females Latvia[LT females Latvia$Year == 2019,]$mx
# In order for the horiuchi model to work properly, the pars1 and pars2
# parameters need to be of length 19, not 111. We need to transform the mx1_Latvia and mx
# here, we create a vector that will be used to transform our data into
# the typical unabridged LT format
age\_groups \leftarrow c(0, rep(1, 4), rep(seq(2, 18), each = 5), rep(18, 21))
# Compute mean mortality rates for each group
mx1 Latvia 19 <- tapply(mx1 Latvia, age groups, mean)
mx2_Latvia_19 <- tapply(mx2_Latvia, age_groups, mean)</pre>
# Convert to numeric vector
mx1_Latvia_19 <- as.numeric(mx1_Latvia_19)</pre>
mx2_Latvia_19 <- as.numeric(mx2_Latvia_19)</pre>
# age specific contributions to the change in Lifespan variation
Results horiuchi Latvia <- horiuchi(func = sd.frommx,
                                     pars1 = mx1 Latvia 19,
                                     pars2 = mx2 Latvia 19, N = 100)
# Extract just the labels
age labels <- as.character(age names)</pre>
# Convert to an ordered factor, resolves issues had beforehand with plotting
age labels <- factor(age labels, levels = age labels)</pre>
# Plot Results
    qqplot() +
      ggtitle(bquote(~"Change in SD Latvia 2000-2019")) +
      geom_bar(aes(x = age_labels, y = Results_horiuchi_Latvia),
               stat = "identity", position = "stack") +
      xlab("Age Groups") +
      ylab("Results")
```

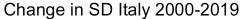
Change in SD Latvia 2000-2019

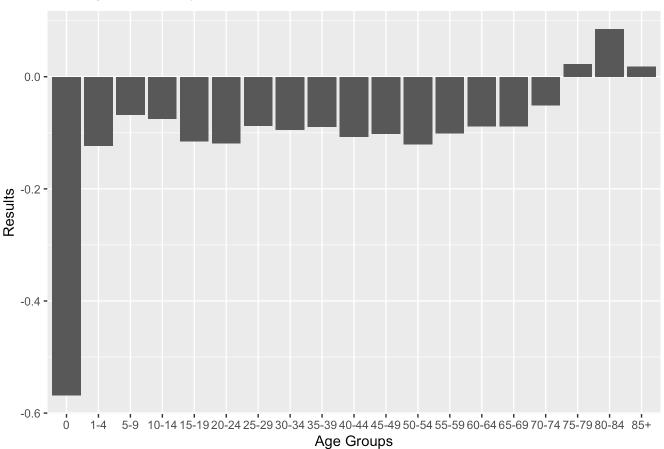


Italy:

```
# Italy
mx1_Italy <- LT_females_Italy[LT_females_Italy$Year == 2000,]$mx</pre>
mx2_Italy <- LT_females_Italy[LT_females_Italy$Year == 2019,]$mx</pre>
# Compute mean mortality rates for each group
mx1_Italy_19 <- tapply(mx1_Italy, age_groups, mean)</pre>
mx2_Italy_19 <- tapply(mx2_Italy, age_groups, mean)</pre>
# Convert to numeric vector
mx1 Italy 19 <- as.numeric(mx1 Italy 19)</pre>
mx2_Italy_19 <- as.numeric(mx2_Italy_19)</pre>
# age specific contributions to the change in Lifespan variation
Results_horiuchi_Italy <- horiuchi(func = sd.frommx,</pre>
                                     pars1 = mx1_Italy_19,
                                     pars2 = mx2_Italy_19, N = 100)
# Plot results
ggplot() +
  ggtitle(bquote(~"Change in SD Italy 2000-2019")) +
  geom_bar(aes(x = age_labels, y = Results_horiuchi_Italy),
           stat = "identity", position = "stack") +
```

xlab("Age Groups") +
ylab("Results")





5. Which age groups contribute to the changes observed? interpret your results and speculate what causes of death may be driving the observed trends.

It appears that the first age group from ages 0 to 1 is contributing the most to changes in SD for both countries from 2000 to 2019. It may be that infant mortality was heavily reduced during this time, thus making the contribution to changes in SD more noticeable for this particular age group.

As far as possible causes of death that may be driving this trend, it could be that less infants are dying from conditions of the perinatal period. Additionally, it may be that this age group is experiencing less deaths due to infectious disease, possibly due to improvements in treatment.

6. How do we interpret lifespan inequality and why do authors think it is important?

Lifespan inequality can be seen as an alternative measure of longevity, particularly alongside life expectancy. While life expectancy can be a useful measure for monitoring improvements in health and development across the globe, it doesn't tell us the whole story. Understanding lifespan inequality,

which is interested in capturing the variation in when people die, is equally important for researchers and authors to consider.

For example, two countries can have an improved life expectancy at birth of 15 years from 1950 to 2020, but they can experience different lifespan variations over this same time period. This is an important factor in measuring inequality because it involves going a step further to unpack whether deaths are more concentrated or dispersed around a specific age. In an ideal state, deaths would be more clustered around a specific age so as to indicate that individuals are dying "equally", for lack of a better term.

Question Three - Projections

Load the data for France

```
# set working directory to access France, WW1.xlsx
setwd("~/Desktop/Term 2/Population Dynamics & Projections/Summative Assessment - work/")
# load France, WW1 excel file
France_WWI <- read_excel("France, WW1.xlsx")</pre>
```

Some data handling:

```
# change column names, will make it simpler to follow code from practical
colnames(France_WWI) <- c("age","nNx_m","nNx_m1920.5","nNx_f", "nNx_f1920.5", "nLx_m","nL

# Read the data as a matrix
France_WWI <- as.matrix(France_WWI)

# create separate matrices for females and males
female <- France_WWI[,c("age","nLx_f","nFx","nNx_f")]
male <- France_WWI[,c("age","nLx_m","nNx_m")]</pre>
```

Important definitions:

```
year <- 1910.5 # mid-year population
ages <- nrow(female) # The number of age groups
radix <- 1 # The radix of a life table
n <- 1 # The length of the age intervals—which is the same for all
SRB <- 1.05 # The Sex Ratio at Birth</pre>
```

Female population, 1910.5:

```
 nLx_f \leftarrow female[,"nLx_f"] \ \# \ creates \ a \ vector \ of \ nLx \ values \ for \ females \ in \ 1910.5 \\ nSx_f \leftarrow c(nLx_f[1]/(n*radix),nLx_f[2:(ages - 1)]/nLx_f[1:(ages - 2)],nLx_f[ages]/sum(nLx nFx \leftarrow female[,"nFx"] \ \# \ creates \ a \ vector \ of \ nFx \ values \ for \ females \ in \ 1910.5
```

```
# Incorporates specific assumptions about fertility.
nGx <- (nFx + c(nFx[2:ages]*nSx_f[2:ages],0))*n/2</pre>
```

Create LESIiE matrix which will be used for population projections:

```
# Define a space for the Leslie matrix
lESliE <- matrix(0,ages,ages)

# Inserts multipliers to project the first age group-depending on the age distribution of
lESliE[1,] <- nGx*1/(1 + SRB)*nSx_f[1]

# Inserts multipliers to project (part of) the open-ended age group
lESliE[ages,ages] <- nSx_f[ages]

# Inserts multipliers to project the remaining age groups
for (x in 1:(ages - 1)) {
    lESliE[x + 1,x] <- nSx_f[x + 1]
}</pre>
```

Create sub-matrices (IESliE_ff, IESliE_mm, IESliE_fm, and IESliE_mf):

```
# Note we use the subscript ff to define a Leslie matrix to project female population usi

LESLIE_ff <- LESLIE

# Defines a space for the Leslie matrix to project male population using only male inform

LESLIE_mm <- matrix(0, ages, ages)

# Defines a space for the Leslie matrix to project male population using only female info

LESLIE_fm <- matrix(0, ages, ages)

# Defines a null space for the Leslie matrix to project female population using only male

LESLIE_mf <- matrix(0, ages, ages)
```

Male population, 1910.5:

```
# Male population
nLx_m <- male[,"nLx_m"]

# Calculation of survivorship ratios is the same as for female population, just using nLx
nSx_m <- c(nLx_m[1]/(n*radix),nLx_m[2:(ages - 1)]/nLx_m[1:(ages - 2)],nLx_m[ages]/sum(nLx

# Inserts multipliers to project the first age group-depending on the age distribution of
lESliE_fm[1,] <- nGx*SRB/(1 + SRB)*nSx_m[1]

# Inserts multipliers to project (part of) the open-ended age group
lESliE_mm[ages,ages] <- nSx_m[ages]</pre>
```

```
# Inserts multipliers to project the remaining age groups
for (x in 1:(ages - 1)) {
   leslie_mm[x + 1,x] <- nsx_m[x + 1]
}</pre>
```

Now, extend LESLiE matrix to 202x202 to account for both sexes:

```
# Concatenates the Leslie matrices for the two sexes
lESliE <- rbind(cbind(lESliE_ff,lESliE_mf),cbind(lESliE_fm,lESliE_mm))</pre>
```

Current vs. projected:

```
current <- rbind(matrix(female[,"nNx_f"],ages),matrix(male[,"nNx_m"],ages))
sum(current) # 39,185,119</pre>
```

[1] 39185119

```
projected <- lESliE%*%current
sum(projected) # 39,261,845</pre>
```

[1] 39261845

Now create a data frame called pop that takes the columns of interest:

```
pop <- France_WWI[,c("age","nNx_m","nNx_f")] #Retains only relevant information for the t
colnames(pop)[colnames(pop) == "nNx_f"] <- str_c("nNx_f", sprintf("%.1f", year)) # Renames
colnames(pop)[colnames(pop) == "nNx_m"] <- str_c("nNx_m", sprintf("%.1f", year))

# don't execute command more than once, otherwise, pop object will change
# for loop iterates every 5 years (from 1910.5-1915.5 and 1915.5-1920.5)

for (t in seq(year, year + 5*n, 5*n)) {
    current <- rbind(matrix(pop[,str_c("nNx_f",sprintf("%.1f",t))],ages),matrix(pop[,str_c(
    projected <- lESliE%*%current #Projects the population n years using the most recent da
    female <- matrix(projected[1:ages,],ages) # Extracts female population
    colnames(female)[1] <- str_c("nNx_f",sprintf("%.1f",t + 5*n)) # Gives a name to the pro
    male <- matrix(projected[(1 + ages):(2*ages),],ages) # Extracts male population
    colnames(male)[1] <- str_c("nNx_m",sprintf("%.1f",t + 5*n)) # Gives a name to the proje
    pop <- cbind(pop,male,female) # Concatenates the table to include the new set of projec
}

pop # lists the current population for males and females in 1910.5,</pre>
```

```
age nNx_m1910.5 nNx_f1910.5 nNx_m1915.5 nNx_f1915.5 nNx_m1920.5
[1,] 0 367716.62 358234.02 368144.49208 356954.12272 368372.32812
[2,] 1 339158.18 334671.47 346740.93085 340321.51302 347144.39578
[3,] 2 327918.23 324483.68 332405.17053 328068.69769 339836.94053
```

```
326062.55 324714.45433 321309.73080 329157.55724
 [4,]
        3
            329206.46
 [5,]
        4
                        328307.88 327138.94564 323917.93444 322675.15110
            331455.83
 [6,]
       5
            332074.81
                        329878.16 329911.08719 326715.38751 325614.32158
 [7,]
        6
            331059.81
                        328866.72 330827.36280 328599.77583 328671.76807
 [8,]
       7
            337225.14
                        334094.89 330040.41541 327743.21761 329808.68395
 [9,]
                        333224.65 336321.55936 333088.16935 329156.08594
            337009.72
       8
                        331817.71 336187.57815 332356.52446 335501.09629
[10,]
       9
            336030.96
[11,]
                        331396.77 335293.22094 331033.14904 335449.49524
      10
            335039.61
                        322115.12 334339.64242 330632.22404 334592.72351
[12,]
      11
            324803.11
[13,]
      12
            330785.48
                        326452.84 324115.76181 321331.53948 333632.11303
[14,]
      13
            333849.14
                        328227.50 330057.74814 325544.36189 323402.70341
                        322815.44 333086.18215 327199.42575 329303.45490
[15,]
      14
            326656.03
[16,]
      15
                        323320.21 325774.37628 321656.60256 332187.17327
            326395.44
                        325556.47 325266.65408 322012.98846 324647.73821
[17,]
      16
            327013.43
                        321406.39 325641.04640 324194.50369 323901.60121
[18,]
      17
            321388.54
[19,]
            308707.11
                        312820.16 319797.64282 319918.89063 324029.09900
      18
[20,]
      19
            303616.91
                        314456.78 306997.78042 311253.70453 318026.90431
[21,]
      20
            311757.44
                        318102.02 301846.32748 312826.36739 305207.48191
                        313602.77 309745.72823 316355.69337 299898.57024
[22,]
      21
            315925.65
                        316305.93 313699.64728 311762.56148 307563.26905
[23,]
      22
            311329.86
[24,]
      23
                        311400.09 309171.16209 314396.53892 311524.51775
            301101.06
                        318786.05 299139.08984 309527.74520 307156.60727
[25,]
      24
            308463.53
[26,]
      25
            310244.61
                        319359.46 306511.94045 316843.75719 297246.49424
[27,]
      26
            303369.22
                        309734.18 308170.03859 317369.58299 304462.32900
[28,]
                        310290.47 301288.51268 307778.62113 306056.40408
      27
            303717.59
                        307428.73 301602.21725 308323.56679 299190.05829
[29,]
      28
            302512.45
                        304847.03 300340.36057 305483.78480 299436.66344
[30,]
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            299051.31
                        304950.67 296906.05619 302927.44204 298185.85972
[31,]
      30
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[32,]
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            292003.32
                        295310.19 296733.72570 302985.89551 294740.10241
[33,]
      32
            295319.75
                        296588.36 289824.06338 293362.41755 294519.16547
[34,]
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                        297976.70 293005.02419 294577.07731 287552.41294
            296552.22
                        308539.09 294118.62758 295934.66861 290600.54107
[35,]
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            305302.02
[36,]
            299747.60
                        301699.41 302770.90618 306394.21235 291680.22995
      35
                        284471.86 297161.98597 299485.28299 300159.21320
[37,]
            285173.08
      36
                        288746.86 282632.87251 282389.61626 294514.98611
[38,]
      37
            288074.44
[39,]
      38
            256053.28
                        258388.06 285481.91591 286645.14915 280089.31978
[40,]
      39
            257118.22
                        260164.71 253569.64245 256445.64680 282712.82970
[41,]
      40
                        272857.94 254452.78137 258209.69196 250940.99046
            264841.32
                        256001.15 262008.69895 270751.27550 251731.27135
[42,]
      41
            249376.34
[43,]
      42
            246522.64
                        251948.79 246584.76359 253950.03666 259075.71299
                        258053.24 243644.53026 249860.20521 243705.92857
[44,]
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            253769.90
[45,]
      44
            253555.46
                        259742.07 250673.75502 255824.38604 240671.92087
[46,]
      45
            247243.77
                        253033.52 250300.89310 257379.70107 247456.17688
                        248094.06 243946.68816 250633.84627 246963.04346
[47,]
      46
            241970.54
                        249729.40 238646.02125 245754.78433 240595.01840
[48,]
      47
            240378.55
[49,]
                        240641.08 236866.15223 247248.40720 235158.93908
      48
            233978.27
                        243661.37 230369.00495 238051.87814 233212.33973
[50,]
      49
            235950.71
                        238370.50 232161.93031 240907.76772 226669.85352
[51,]
      50
            229428.33
[52,]
      51
            218551.83
                        226963.33 225499.98718 235574.67407 228186.78195
                        210380.17 214551.19634 224153.75338 221372.16616
[53,]
      52
            201916.00
[54,]
      53
            198738.00
                        208312.33 198027.67262 207622.15668 210419.55105
```

[55 ,]	54	197078.83	210295.50	194783.65164	205451.03750	194087.45785
[56 ,]	55	188158.00	202582.50	192910.89277	207282.12238	190664.25417
[57 ,]	56	182971.50	192670.67	183946.16558	199484.89605	188592.66693
[58 ,]	57	184477.00	196200.00	178617.54794	189472.22420	179569.02058
[59 ,]	58	185019.17	202564.50	179751.81806	192777.99956	174042.44962
[60,]	59	182784.83	197319.83	179996.60697	198787.04220	174872.24349
[61,]	60	172232.67	187457.67	177485.05418	193180.45487	174777.67460
[62,]	61	160620.50	190740.50	166830.53291	183223.55746	171918.17425
[63,]	62	148549.83	176629.00		186152.69236	
[64,]	63	138892.67			171897.93193	
[65,]	64	144584.00			153932.00399	
[66,]	65	142125.50			161605.98926	
[67,]	66	132664.00			157706.31659	
[68,]	67	127861.50	145854.00		144405.16957	
[69,]	68	122982.83			139648.92285	
[70,]	69	117002.17	137517.83		135379.26829	
[71,]	70	107989.33			130688.13295	
[72 ,]	71	96270.00			120584.97815	
[73 ,]	72	86433.00	105734.83		106785.07394	92834.21672
[74,]	73	80159.17	98107.50	79257.50993	98568.46553	81572.82178
[75 ,]	74	77232.67	95528.50	72881.36869	90738.16705	72061.57203
[76,]	75	67464.17	84599.50	69650.15849	87560.20288	65726.05712
[77,]	76	56859.33	72289.83	60191.86504	76768.45901	62142.21475
[78 ,]	77	49526.83	64150.83	50109.75700	64977.75348	53046.69841
[79 ,]	78	43301.67	56347.00	43117.73510	57003.78388	43625.22754
[80,]	79	40756.17	55811.50	37389.54282	49756.15751	37230.72119
[81,]	80	33737.05	47465.01	34684.31909	48815.92741	31819.25175
[82,]	81	27411.41	39178.28	28208.46032	40954.54740	29000.49763
[83,]	82	22321.73	32931.17	22606.47965	33387.26985	23263.81547
[84,]	83	17987.65	26937.56	18095.44944	27622.17846	18326.28607
[85,]	84	14106.12	21618.41	14307.65692	22188.68981	14393.40228
[86,]	85	10745.87	17230.10	10989.82083	17475.86243	11146.83457
[87,]	86	8004.64	13622.71	8172.49968	13695.67020	8358.03032
[88,]	87	5854.93	10429.38			
[89,]		4229.02	7695.74	4265.80429	8077.41313	
[90,]	89	3024.22	5739.63	3012.41291	5824.75885	3038.61507
[91,]	90	2026.90	4163.03	2133.77049		2125.43987
[92 ,]	91	1301.18	2763.13	1368.17020		1440.30845
[93,]		846.78	1833.49	852.96563		
[94,]		559.29	1295.41	557.07241		
[95 ,]	94	341.67	881.15	355.61688	887.77225	354.20686
[96,]	95	195.67	560.73	206.18191		214.59820
[97 ,]	96	125.27	339.91	118.44758		124.81089
[98,]		70.05	179.69	82.08982		77.61906
[99,]		36.10	112.18	43.53255		51.01468
[100,]	99	21.92	73.05	16.48085		19.87405
[101,]		22.35	64.67	19.22504		15.50591
, -		_f1920.5				_
		_				

nNx_f1920.5

- [1,] 357175.03331
- [2,] 339105.61349
- [3,] 333607.27036

- [4,] 324859.68151
- [5,] 319196.37602
- [6,] 322346.73585
- [7,] 325449.26010
- [0] 227477 4252
- [8,] 327477.18539
- [9,] 326755.63632
- [10,] 332220.39938
- [11,] 331570.68951
- [12,] 330269.44197
- [13,] 329827.92473
- [14,] 320437.31333
- [15,] 324524.69176
- [16,] 326024.85075
- [17,] 320356.10718
- [17,] 320330.10710
- [18,] 320665.84631
- [19,] 322694.10066
- [20,] 318316.88808
- [21,] 309639.89940
- [22,] 311109.00320
- [23,] 314499.33080
- [24,] 309880.59658
- [24,] 505000155050
- [25,] 312506.17747
- [26,] 307641.86119
- [27,] 314869.55512
- [28,] 315365.81671
- [29,] 305827.64030
- [30,] 306372.96042
- [31,] 303560.18727
- [32,] 300975.70306
- [33,] 300987.49654
- [34,] 291373.01125
- [35,] 292558.34350
- [36,] 293877.41338
- [37,] 304145.63088
- [38,] 297293.14576
- [39,] 280334.17808
- [40,] 284490.31537
- [41,] 254518.57581
- [42,] 256216.12274
- [43,] 268581.98230 [44,] 251844.86210
- - -
- [45,] 247702.11602
- [46,] 253497.64867
- [47,] 254938.80981
- [48,] 248270.62299 [49,] 243313.27825
- [50,] 244588.11314
- [51,] 235361.66838
- [52,] 238082.18240
- [53,] 232658.49771
- [54,] 221215.17301

```
[55,] 204770.34412
 [56,] 202507.07742
 [57,] 204112.65848
 [58,] 196173.33011
 [59,] 186167.56551
 [60,] 189183.04211
 [61,] 194616.88789
 [62,] 188817.08160
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 [66,] 148893.06283
 [67,] 155739.56003
 [68,] 151468.77420
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 [71,] 128655.78095
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       91164.50721
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 [98,]
          232.45099
 [99,]
          136.60056
[100,]
           65.44428
[101,]
           68.17779
```

```
# and projects for years 1915.5 and 1920.5
```

1. Calculate the Total Fertility Rate of this population

```
TFR <- sum(nFx)*n
TFR # TFR = 2.61 births per woman</pre>
```

[1] 2.61

2. Calculate the Net Reproduction Rate of this population

```
NRR <- sum(nFx*nLx_f/radix)*(1/(1+SRB)) # Formula on Slide 27 from Cohort Compoent Model NRR # NRR = 0.9700091 , NRR < 1 posing that the population will decline
```

[1] 0.9700091

3. Calculate the projected number of births from 1910 to 1920 by age group of the mother

The following command only gives us the projected number of births by age group of the mother one year in advance:

```
# projected_births <- ((current + lESliE%*%current)/2) * nFx * n</pre>
```

So, we must create a for loop that gives us the number of births by age group per year from 1910.5 - 1920.5

```
# Set up parameters
years = seq(1910.5, 1919.5, by = 1)
# Initialize birth_by_age_group object
births_by_age_group <- matrix(₀, nrow <- length(current), ncol <- length(years)) # Sto
# Loop from 1910.5 to 1920.5 (each step projects to the next year)
# only complete loop once, otherwise results are conflated
for (t in 1:length(years)) {
 # Project population one year forward
 projected population <- lESliE %*% current
 # Compute births by age group for this particular year
 births_by_age_group[, t] <- ((current + projected) / 2) * nFx * n # this is the formul
 # Update population for next iteration
 current <- projected
}
# Here, I've specified that I'm only interested in looking at the age groups of the femal
births_by_age_group <- births_by_age_group[0:101, ]</pre>
```

```
# Let's also change the column names
colnames(births_by_age_group) <- c("1911.5", "1912.5", "1913.5", "1914.5", "1915.5", "191
births_by_age_group</pre>
```

```
1911.5
                                  1912.5
                                                   1913.5
                                                                     1914.5
 [1,]
         0.00000000000
                                             0.0000000000
                                                               0.0000000000
                            0.0000000000
 [2,]
         0.00000000000
                            0.000000000
                                             0.000000000
                                                               0.0000000000
 [3,]
         0.00000000000
                            0.0000000000
                                             0.0000000000
                                                               0.0000000000
[4,]
         0.00000000000
                            0.0000000000
                                             0.0000000000
                                                               0.0000000000
[5,]
         0.00000000000
                            0.0000000000
                                             0.0000000000
                                                               0.0000000000
[6,]
         0.00000000000
                            0.0000000000
                                             0.000000000
                                                               0.0000000000
[7,]
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                            0.0000000000
                                             0.000000000
                                                               0.0000000000
[8,]
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                                                               0.0000000000
[9,]
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                            0.0000000000
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                                                               0.0000000000
[10,]
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                            0.0000000000
                                             0.0000000000
                                                               0.0000000000
[11,]
         0.00000000000
                            0.0000000000
                                             0.000000000
                                                               0.0000000000
[12,]
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                            0.0000252801
                                             0.0000252801
                                                               0.0000252801
[13,]
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                                             0.2966119131
                                                               0.2966119131
                            0.2966119131
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                           18.0281928957
                                            18.0281928957
                                                             18,0281928957
[15,]
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                                           281.4713612519
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[76 ,]	0.0000000000	0.0000000000
[77 ,]	0.0000000000	0.0000000000
[78 ,]	0.000000000	0.0000000000
[79 ,]	0.000000000	0.000000000
[80,]	0.000000000	0.000000000
-		
[81,]	0.0000000000	0.0000000000
[82,]	0.0000000000	0.0000000000
[83,]	0.0000000000	0.0000000000
[84,]	0.0000000000	0.0000000000
[85,]	0.0000000000	0.0000000000
[86,]	0.0000000000	0.0000000000
[87 ,]	0.0000000000	0.0000000000
[88,]	0.0000000000	0.0000000000
[89 ,]	0.0000000000	0.0000000000
[90,]	0.0000000000	0.0000000000
[91,]	0.0000000000	0.0000000000
[92,]	0.0000000000	0.0000000000
[93 ,]	0.0000000000	0.0000000000
[94,]	0.0000000000	0.000000000
/ -		

```
[95.]
           0.0000000000
                             0.0000000000
 [96,]
           0.0000000000
                             0.0000000000
 [97,]
           0.0000000000
                             0.0000000000
 [98,]
           0.0000000000
                             0.0000000000
 [99,]
           0.0000000000
                             0.0000000000
[100,]
           0.0000000000
                             0.0000000000
[101,]
           0.0000000000
                             0.000000000
```

4. Calculate the projected total number of births for each projected year

5. Calculate how many are projected to be male births

```
male_births <- total_births * SRB / (1 + SRB)
male_births

1911.5    1912.5    1913.5    1914.5    1915.5    1916.5    1917.5    1918.5
403015.3    403127.0    403127.0    403127.0    403127.0    403127.0    403127.0
1919.5    1920.5
403127.0    403127.0</pre>
```

6. Using a plot of overlapping population pyramids, compare the projected population of 1920.5 with the population counts of the same year (post-exposure).

Projected Population - 1920.5:

```
# projected population for 1920.5

pop <- data.frame(pop)
projected_pop_1920.5 <- pop

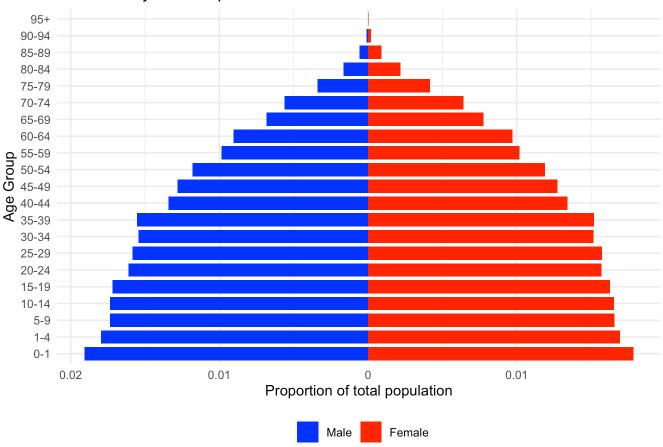
# remove non-relevant column years
projected_pop_1920.5 <- projected_pop_1920.5 %>%
    select(-c(nNx_m1910.5, nNx_f1910.5, nNx_m1915.5, nNx_f1915.5))

# Convert male population to negative values
projected_pop_1920.5 <- projected_pop_1920.5 %>%
    mutate(nNx_m1920.5 = -nNx_m1920.5)
```

Plot a population pyramid for just the projected population in 1920.5:

```
# Transform data frame into age groups and not single year age-intervals
# to make more interpretable on a plot
projected_pop_long <- projected_pop_long %>%
 mutate(Age Group = cut(age,
         breaks = c(0, 1, 5, 10, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80,
         labels = c("0-1", "1-4", "5-9", "10-14", "15-19", "20-24", "25-29", "30-34", "35-
         right = FALSE)) # Ensures left-inclusive intervals
# Plot
ggplot(projected_pop_long, aes(x = Proportion, y = Age_Group, fill = Sex)) +
          geom_col(width = 0.8, position = "identity") + # Use geom_col() instead of geo
          scale_fill_manual(values = c("Male" = "blue", "Female" = "red"), breaks = c("Ma
          scale_x_continuous(labels = abs) + # Keep x-axis values positive
          labs(title = "France Projected Population 1920.5",
               x = "Proportion of total population",
               y = "Age Group",
               fill = "") +
          theme_minimal() +
          theme(legend.position = "bottom")
```

France Projected Population 1920.5



Actual Population - 1920.5:

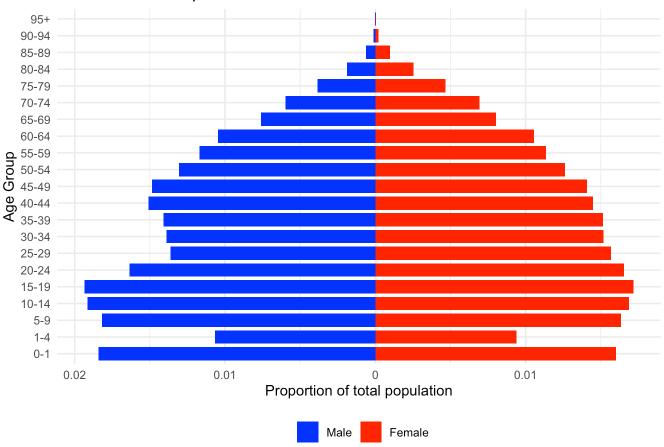
```
# transform France_WWI into a data frame to work with
France WWI <- data.frame(France WWI)</pre>
actual pop 1920.5 <- France WWI
# remove non-relevant column years
actual_pop_1920.5 <- actual_pop_1920.5 %>%
  select(-c(nNx_m, nNx_f, nLx_m, nLx_f, nFx))
# Convert male population to negative values
actual_pop_1920.5 <- actual_pop_1920.5 %>%
  mutate(nNx_m1920.5 = -nNx_m1920.5)
# create an object call actual_pop_long that converts to long format
actual_pop_long <- actual_pop_1920.5 %>%
  pivot_longer(cols = c(nNx_m1920.5, nNx_f1920.5),
               names to = "Sex",
               values_to = "Population")
# Recode Sex labels
actual_pop_long$Sex <- recode(actual_pop_long$Sex,</pre>
                              nNx m1920.5 = "Male",
                              nNx_f1920.5 = "Female")
```

```
# add a column to specify the percentage of pop instead of absolute value
actual_pop_long <- actual_pop_long %>%
    group_by(Sex) %>% # Ensure we calculate proportions separately for each sex
mutate(Proportion = Population / sum(abs(Population))) %>%
ungroup() # Remove grouping after calculation
```

Plot a population pyramid for just the actual population in 1920.5:

```
# Transform data frame into age groups and not single year age-intervals
actual_pop_long <- actual_pop_long %>%
 mutate(Age Group = cut(age,
         breaks = c(0, 1, 5, 10, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80,
         labels = c("0-1", "1-4", "5-9", "10-14", "15-19", "20-24", "25-29", "30-34", "35-
         right = FALSE)) # Ensures left-inclusive intervals
# Plot
ggplot(actual_pop_long, aes(x = Proportion, y = Age_Group, fill = Sex)) +
          geom_col(width = 0.8, position = "identity") + # Use geom_col() instead of geo
          scale fill manual(values = c("Male" = "blue", "Female" = "red"), breaks = c("Ma
          scale_x_continuous(labels = abs) + # Keep x-axis values positive
          labs(title = "France Actual Population 1920.5",
               x = "Proportion of total population",
               y = "Age Group",
               fill = "") +
          theme minimal() +
          theme(legend.position = "bottom")
```

France Actual Population 1920.5

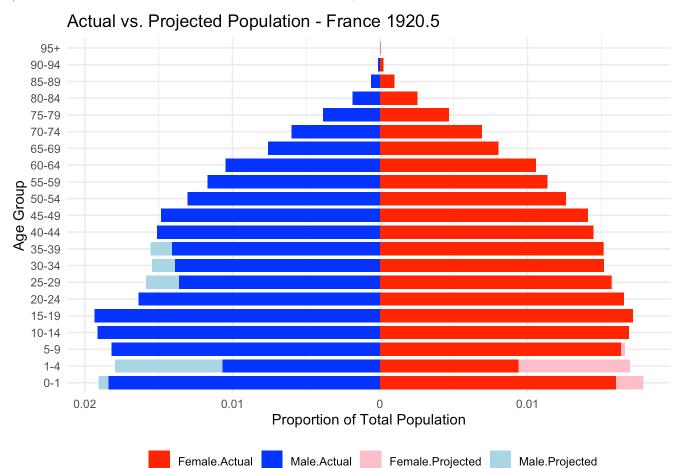


Now, overlap the two population pyramids to identify where differences exist:

```
# combined population pyramids - actual vs. projected
# Merge actual and projected data
    combined_pop_long <- bind_rows(</pre>
      projected_pop_long %>% mutate(Type = "Projected"),
      actual_pop_long %>% mutate(Type = "Actual")
    )
# Create the overlapping population pyramids
ggplot(combined\_pop\_long, aes(x = Proportion, y = Age\_Group, fill = interaction(Sex, Type)
 geom_col(width = 0.8, position = "identity") +
  scale_fill_manual(values = c("Male.Actual" = "blue", "Male.Projected" = "lightblue",
                               "Female.Actual" = "red", "Female.Projected" = "pink")) +
 scale_x_continuous(labels = abs) +
  labs(title = "Actual vs. Projected Population - France 1920.5",
       x = "Proportion of Total Population",
       y = "Age Group",
       fill = "") +
  theme minimal() +
  theme(legend.position = "bottom")
```

Female.Projected

Male.Projected



7. Identify the age groups that were mostly affected during the Great War

Comparing the two population pyramids, it appears that males ages 25 - 40 were severely affected during the Great War. This intuitively makes sense as more men of these age groups were likely fighting during the Great War, which started in 1914 and went until 1919. We can see that this pattern is not noticed with females.

Looking at the plot, it is clear to see that the age group 1-4 was also heavily impacted from the Great War. This makes sense as it is understandable that at this time of crisis, many families are not having as many children.

The difference between the projected population and the actual population for males in this age interval is 630,017. For females, the difference between the projected population and the actual population is 618,130. This is for a total population difference of 1,248,147, which is quite the difference (see R code below for how results were obtained)

```
# Summing the male population in the 1-4 age group - projected
male_pop_age_group_1_4_projected <- projected_pop_long %>%
  filter(Sex == "Male", Age_Group == "1-4") %>%
  summarise(total_population = abs(sum(Population, na.rm = TRUE)))
```

```
# Summing the male population in the 1-4 age group - actual
male_pop_age_group_1_4_actual <- actual_pop_long %>%
filter(Sex == "Male", Age_Group == "1-4") %>%
summarise(total_population = abs(sum(Population, na.rm = TRUE)))

# Difference between the two - males
male_pop_age_group_1_4_projected - male_pop_age_group_1_4_actual
```

```
total_population
1 630017.3
```

```
# Summing the female population in the 1-4 age group - projected
female_pop_age_group_1_4_projected <- projected_pop_long %>%
   filter(Sex == "Female", Age_Group == "1-4") %>%
   summarise(total_population = abs(sum(Population, na.rm = TRUE)))

# Summing the male population in the 1-4 age group - actual
female_pop_age_group_1_4_actual <- actual_pop_long %>%
   filter(Sex == "Female", Age_Group == "1-4") %>%
   summarise(total_population = abs(sum(Population, na.rm = TRUE)))

# Difference between the two - females
female_pop_age_group_1_4_projected - female_pop_age_group_1_4_actual
```

```
total_population
618130
```

```
total_population
1248147
```

References:

Aburto, J.M., Villavicencio, F., Basellini, U., Kjærgaard, S., & Vaupel, J.W. (2020). Dynamics of life expectancy and life span equality. *Proceedings of the National Academy of Sciences of the United States of America*, *117*(10), 5250–5259.

Aburto, J.M. (2025) "Alternative measures of longevity" [Lecture] 2429 Population Dynamics & Projections. London School of Hygiene & Tropical Medicine. 07 March.

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Statement to acknowledge AI use:

I used ChatGPT (OpenAI, 2025) to assist with the debugging of R code as specified on the cover sheet form.