## Demographic Methods - Practical 4 (Life Tables II)

2025-10-20

### The heading of the R script

The R Script starts by clearing all generated data if any. Packages and libraries are not required this time.

rm(list = ls())

#### Reading the data

In this exercise, we are given the number of survivors  $l_x$  from two life tables corresponding to the male and female populations in Athens in 1981. Additionally, we are provided with the life expectancy at age 10  $e_{10}$ , for each population. As shown in the command lines below, the  $l_x$  values and the exact ages x are combined into a vector using the R function "c()".

```
##
           lx_m
                   lx_f
       х
## 1
       0 100000 100000
## 2
          97731
                  98179
       1
## 3
       5
          97203
                  97884
## 4
      10
          96925
                  97752
## 5
      15
          96723
                  97658
## 6
      20
          96364
                  97512
## 7
      25
          95894
                  97313
## 8
      30
          95445
                  97068
## 9
      35
          94958
                  96761
          94343
## 10 40
                  96343
## 11 45
          93436
                  95721
## 12 50
          91883
                  94669
## 13 55
          89212
                  92945
## 14 60
          84660
                  90131
## 15 65
          77483
                  85492
## 16 70
          66830
                  77598
## 17 75
          52309
                  65075
## 18 80
          34968
                  47402
## 19 85
          18173
                 27701
```

#### Data preparation

Using the R function "cbind( $name_1 = vector_1, \ldots, name_n = vector_n$ )", we can create a matrix with n columns—one column of  $l_x$  values per population—and one row per age interval [x, x + n). Similarly, we can create a vector with two elements, one  $e_{10}$  value for each population. Naming the elements of vectors or matrices is optional, but it helps with identification later. There is also a complementary R function, "rbind()", which performs the same operation by rows.

```
lx = cbind(male = lx_m, female = lx_f)
e10 = cbind(male = e10_m, female = e10_f)
```

As is standard in life table calculations, we can define the length of age intervals based on the values of x, make assumptions about the age distribution of deaths within each interval using  $na_x$ , and identify the open-ended age interval using logical operators.

```
n = c(diff(x,1),NA)

nax = c(0.1,0.4,rep(0.5, length(x) - 3),NA)

sEL = !is.na(n)
```

Note that the value of  $na_x$  may also vary by sex, but no adjustments have been made to account for this.

#### Life table formulae

$$_{n}M_{x} = \frac{_{n}D_{x}}{_{n}N_{x}} \tag{1}$$

$${}_{n}q_{x} = \frac{n \cdot_{n} M_{x}}{1 + n \cdot (1 -_{n} a_{x}) \cdot_{n} M_{x}}$$

$$\tag{2}$$

$$_{\infty}q_x = 1.00 \tag{3}$$

$$_{n}p_{x}=1-_{n}q_{x}\tag{4}$$

$$l_{x+n} =_n p_x \cdot l_x \tag{5}$$

$$_{n}d_{x} =_{n} q_{x} \cdot l_{x} \tag{6}$$

$$_{n}d_{x} = l_{x} - l_{x+n} \tag{7}$$

$$_{n}L_{x} = n \cdot (l_{x} -_{n} d_{x}) + n \cdot_{n} a_{x} \cdot_{n} d_{x}$$

$$\tag{8}$$

$$_{\infty}L_{x} = \frac{l_{x}}{_{\infty}M_{x}} \tag{9}$$

$$T_x = \sum_{a=x}^{\infty} {}_n L_a \tag{10}$$

$$e_x = \frac{T_x}{l_x} \tag{11}$$

## Exercise 1: life table revision question (Compulsory)

Calculate the following values and comment on your results. While you may choose to reconstruct the entire life table starting from  $l_x$ , this is not required to solve questions a, b, c, d, and e.

```
a. _1p_0, _4p_1, _5p_{10}, _{10}p_{75}.
Hint: All these quantities depend on l_x.
```

**a.1** 
$$_{1}p_{0} = \frac{l_{1}}{l_{0}}$$

```
lx[x == 1,]/lx[x == 0,]
       male female
## 0.97731 0.98179
a.2
_4p_1 = \frac{l_5}{l_1}
lx[x == 5,]/lx[x == 1,]
       \mathtt{male}
                     female
## 0.9945974 0.9969953
a.3
_5p_{10} = \frac{l_{15}}{l_{10}}
lx[x == 15,]/lx[x == 10,]
##
          male
                     female
## 0.9979159 0.9990384
_{10}p_{75} = \frac{l_{85}}{l_{75}}
lx[x == 85,]/lx[x == 75,]
##
          male
                     female
## 0.3474163 0.4256781
b. _{1}d_{0}, _{4}d_{1}, _{15}d_{50}.
Hint: Decumulate l_x to calculate the number of deaths in a life table.
b.1
_1d_0 = l_0 - l_1
lx[x == 0,] - lx[x == 1,]
##
      male female
##
      2269
                1821
b.2
_4d_1 = l_1 - l_5
lx[x == 1,] - lx[x == 5,]
##
      male female
##
        528
                 295
b.3
_{15}d_{50} = l_{50} - l_{65}
lx[x == 50,] - lx[x == 65,]
      male female
## 14400
               9177
c. _4q_1, _5q_5, _{15}q_{50}.
Hint: These quantities can be calculated directly from l_x.
_{4}q_{1} = \frac{l_{1}-l_{5}}{l_{1}}
```

```
(lx[x == 1,] - lx[x == 5,])/lx[x == 1,]
             male
                           female
## 0.005402585 0.003004716
c.2
_{5}q_{5}=\frac{l_{5}-l_{10}}{l_{5}}
(lx[x == 5,] - lx[x == 10,])/lx[x == 5,]
             male
                           female
## 0.002859994 0.001348535
c.3
_{15}q_{50} = \frac{l_{50} - l_{65}}{l_{50}}
(1x[x == 50,] - 1x[x == 65,])/1x[x == 50,]
            male
                         female
## 0.15672105 0.09693775
d. _1L_0, _4L_1, _5L_5, _5L_{45}.
Hint: Use the general formula for {}_{n}L_{x} in closed age intervals.
_{1}L_{0} = 1 \cdot l_{1} + 1 \cdot_{1} a_{0} \cdot (l_{0} - l_{1})
n[x == 0]*lx[x == 1,] + n[x == 0]*nax[x == 0]*(lx[x == 0,] - lx[x == 1,])
        male female
## 97957.9 98361.1
d.2
_{4}L_{1} = 4 \cdot l_{5} + 4 \cdot {}_{4} a_{1} \cdot (l_{1} - l_{5})
n[x == 1]*lx[x == 5,] + n[x == 1]*nax[x == 1]*(lx[x == 1,] - lx[x == 5,])
##
         male
                   female
## 389656.8 392008.0
d.3
_5L_5 = 5 \cdot l_{10} + 5 \cdot _5 a_5 \cdot (l_5 - l_{10})
n[x == 5]*lx[x == 10,] + n[x == 5]*nax[x == 5]*(lx[x == 5,] - lx[x == 10,])
##
      male female
## 485320 489090
_{5}L_{45} = 5 \cdot l_{50} + 5 \cdot _{5} a_{45} \cdot (l_{45} - l_{50})
n[x == 45]*lx[x == 50,] + n[x == 45]*nax[x == 45]*(lx[x == 45,] - lx[x == 50,])
##
         male
                   female
## 463297.5 475975.0
e. _1M_0, _4M_1, _{10}M_{60}.
Hint: All these quantities depend on _nd_x and _nL_x.
_{1}M_{0} = \frac{_{1}d_{0}}{_{1}L_{0}} = \frac{l_{0}-l_{1}}{_{1}\cdot l_{1}+1\cdot _{1}a_{0}\cdot (l_{0}-l_{1})}
```

```
(1x[x == 0,] - 1x[x == 1,])/(n[x == 0]*1x[x == 1,] + n[x == 0]*nax[x == 0]*(1x[x == 0,] - 1x[x == 1,]))
               male
                              female
## 0.02316301 0.01851342
_{4}M_{1} = \frac{_{4}d_{1}}{_{4}L_{1}} = \frac{l_{1}-l_{5}}{_{4}\cdot l_{5}+4\cdot _{4}a_{1}\cdot (l_{1}-l_{5})}
(1x[x == 1,] - 1x[x == 5,])/(n[x == 1]*1x[x == 5,] + n[x == 1]*nax[x == 1]*(1x[x == 1,] - 1x[x == 5,]))
                  male
                                     female
## 0.0013550386 0.0007525357
e.3
{}_{10}M_{60} = {}_{5} {}_{\overline{L}65+5} {}_{\overline{L}60} \atop 5\overline{L}_{65}+5\overline{L}_{60}} = {}_{(5 \cdot l_{65}+5 \cdot 5 a_{60} \cdot (l_{60}-l_{65})) + (5 \cdot l_{70}+5 \cdot 5 a_{65} \cdot (l_{65}-l_{70}))} \atop (5 \cdot l_{65}+5 \cdot 5 a_{60} \cdot (l_{60}-l_{65})) + (5 \cdot l_{70}+5 \cdot 5 a_{65} \cdot (l_{65}-l_{70}))}
(1x[x==60,] - 1x[x==70,])/(n[x==60]*1x[x==65,] + n[x==60]*nax[x==60]*(1x[x==60,] - 1x[x==65,])
+ n[x=65]*1x[x=70] + n[x=65]*nax[x=65]*(1x[x=65] - 1x[x=70]))
##
                              female
               male
## 0.02327251 0.01480073
f. e_0 and e_1.
```

#### Life table analysis

## 72.31104 75.07068

Hint: Start by calculating  $T_{10}$ , then find  $T_{85}$  using  $_{75}L_{10}$ . Once  $_nL_x$  has been estimated for all ages, including

the open-ended age interval, proceed to compute  $T_x$  and  $e_x$  following standard life table equations.

R function "apply(input, dimension, function)" repeats the same function for each row (dimension = 1) or each column (dimension = 2). Matrix or vector multiplication should use the notation "%\*%".

```
f.1
e0
ex[x == 0,]
## male female
## 71.64988 74.68726
f.2
e1
ex[x == 1,]
## male female
```

- g. Ask Copilot to estimate the number of years that a newborn female is expecting to live between exact ages 20 and 65.
- h. Ask Copilot to estimate the number of years that a 20-year female is expecting to live between exact ages 20 and 65.

# Exercise 2: application of the stationary population model (Compulsory)

A job training program has 150 training positions that are always filled. The program admits 30 new candidates each month. Every month, 10 trainees quit the program while 20 more find a placement. These conditions have prevailed as long as anyone can remember.

- a. What is the monthly rate of leaving the training program?  $CDR = CBR = \frac{l_x}{T_r} = \frac{30}{150} = 200$  exits per 1,000
- b. What is the probability that a trainee will leave the program by finding a placement?  $\frac{placements}{all~exists} = \frac{20}{30} = 0.6666667$
- c. How long on average does a trainee stay in the program?  $e_0=\frac{1}{CDR}=\frac{1}{0.2}=5$  months
- d. Ask Copilot to solve and explain this exercise by including the problem and questions a, b, and c in the prompt.

# Exercise 3: additional life table questions based on the data of Exercise 1 (Optional)

Hint: Bear in mind that both life tables use the same radix of 100,000 individuals, which does not reflect the slight excess of male births. To aggregate the two populations into a single life table, or to compare quantities that depend on population size—such as the number of survivors or person-years lived—it would be necessary to adjust for the Sex Ratio at Birth. You can assume 105 male births per 100 female births.

a. Calculate the Infant Mortality Rate.

```
SRB = 1.05

sum(ndx[x == 0,]*c(1.05,1))/sum(1x[x == 0,]*c(SRB,1))*1000
```

```
## [1] 20.50463
```

b. If you were told that there had been 35,000 males aged 5-9 in Athens in mid-1981, would you be able to say how many there would be aged 10-14 exactly 5 years later? What assumptions would you have to make?

Hint: The command "nLx[x == 5,"male"]" returns the function  $_nL_x$  at x equal to 5 for the male population.  $_5N_{10} = \frac{_5L_{10}}{_5L_5} \cdot _5N_5 = \frac{_5L_{10}}{_5L_5} \cdot _35000$ 

```
nLx[x == 10, "male"]/nLx[x == 5, "male"]*35000
```

```
## male
## 34913.46
```

Assuming a closed population and keeping constant the mortality rates of 1981.

c. What proportion of male population is aged 5-9 in the stationary population represented by the life table in Table 1?

```
\frac{5D_0}{T_0}
nLx[x == 5,"male"]/Tx[x == 0,"male"]
```

```
## male
## 0.06773493
```

d. What would be the sex ratio in the 25-29 age group in the stationary populations represented by the life tables in Table 1?

```
Hint: you may need to make an assumption about the Sex Ratio at Birth. \frac{5L_{25,male}}{5L_{25,female}} \cdot SRB
sprintf("%.5f",nLx[x == 25, "male"]/nLx[x == 25, "female"]*SRB)
```

```
## [1] "1.03357"
```

e. What would be the Crude Birth Rate in the stationary populations represented by the life table of Athens in 1981? and the Crude Death Rate?  $CDR = CBR = \frac{l_0}{T_c}$ 

```
sum(lx[x == 0,]*c(SRB,1))/sum(Tx[x == 0,]*c(SRB,1))*1000
```

```
## [1] 13.67399
```

f. The two questions below pertain to girl twins were born to a woman on her 20th birthday. Her husband was exactly 5 years older than herself. You can assume that the mortality regimes of Athens in 1981 apply to each member of the family. 1. What is the probability that both mother and children are alive when the twins celebrate their 10<sup>th</sup> anniversary, but that the father had died?

```
\begin{split} P(event) &= P(father\,died) \cdot P(mother\,survived) \cdot P(one\,twin\,survived) \cdot P(other\,twin\,survived) \\ P(event) &= \left[1 - \frac{l_{35,male}}{l_{25,male}}\right] \cdot \left[\frac{l_{30,female}}{l_{20,female}}\right] \cdot \left[\frac{l_{10,female}}{l_{0,female}}\right]^2 \\ \text{sprintf("\%.7f",(1 - lx[x == 35,"male"]/lx[x == 25,"male"])} \\ &\quad *(lx[x == 30,"female"]/lx[x == 20,"female"]) \\ &\quad *(lx[x == 10,"female"]/lx[x == 0,"female"])^2) \end{split}
```

```
## [1] "0.0092844"
```

2. What is the probability that at least one child survived but only one parent is alive 10 years after the birth of the twins?

Hint: in order to obtain the probability that two events jointly occur, we have to multiply probabilities (AND = multiplication, assuming event independence); in other to obtain the probability that either one of two events occur we add the two probabilities (OR = addition).

 $P(at \ least \ one \ twin \ survived) = 1 - P(both \ twins \ died)$ 

```
P1 = 1 - (1 - lx[x == 10, "female"]/lx[x == 0, "female"])^2

sprintf("%.10f",P1)
```

## [1] "0.9994946496"

 $P(only\ one\ parent\ survived) = 1 - P(both\ parent\ survived) - P(both\ parent\ died)$ 

```
P2 = 1 - (lx[x == 30,"female"]/lx[x == 20,"female"])*(lx[x == 35,"male"]/lx[x == 25,"male"]) - (1 - lx[x == 30,"female"]/lx[x == 20,"female"])*(1 - lx[x == 35,"male"]/lx[x == 25,"male"]) sprintf("%.10f",P2)
```

## [1] "0.0142251761"

 $P(event) = P(at \ least \ one \ twin \ survived) * P(only \ one \ parent \ survived)$ 

```
sprintf("%.10f",P1*P2)
```

## [1] "0.0142179874"

## Exercise 4: life tables and stationary population theory (Optional)

```
LT
                       = data.frame(x,lx,ex)
print(LT)
##
      х
          lx
                ex
      0 1000 60.0
## 2 10
         950
                NA
## 3 25
         750 50.0
## 4 65
         600 17.5
## 5 85
         300
              5.0
```

Given the stationary population described in the table and the fact that 16.25% of the population is between exact ages 0 and 10, answer the questions:

### Life table analysis

As is standard in life table calculations, we can define the length of age intervals based on the values of x, and identify the open-ended age interval.

```
n = c(diff(x), NA)
sEL = !is.na(n)
```

The number of deaths is calculated as the first difference of the number of survivors. An incomplete set of  $T_x$  is calculated as the product of the life expectancy and the number of survivors. An incomplete set of  ${}_{n}L_{x}$  is calculated as the first difference of  $T_x$ .

 $_{10}L_0$  accounts for 16.25% of  $T_0$ , as incorporated by the following line.

```
nLx[x == 0] = Tx[x == 0]*16.25/100
```

 $_{15}L_{10}$  is still missing. The variable s is created to identify the missing value. The missing value should be equal to the difference between  $T_0$  and the sum of all  $_nL_x$ .

```
s = is.na(nLx)
nLx[s] = Tx[x == 0] - sum(nLx[!s])
```

Once the function  ${}_{n}L_{x}$  has been identified for all ages,  $T_{x}$  and  $e_{x}$  are recalculated to populate their missing values. Then,  ${}_{n}M_{x}$  is calculated as the ratio of the observed number of events  ${}_{n}d_{x}$ , to the exposure to the risk of dying, measured by the number of person-years lived within the age interval  ${}_{n}L_{x}$ .

```
Tx = rev(cumsum(rev(nLx)))
ex = Tx/lx
nMx = ndx/nLx

LT = data.frame(x,n,nMx,lx,ndx,nLx,Tx,ex)
print(LT)
```

```
nMx
                         lx ndx
                                  nLx
                                         Tx
     0 10 0.005128205 1000
                             50
                                 9750 60000 60.00000
## 2 10 15 0.015686275
                        950 200 12750 50250 52.89474
## 3 25 40 0.005555556
                        750 150 27000 37500 50.00000
## 4 65 20 0.033333333
                        600
                            300
                                 9000 10500 17.50000
## 5 85 NA 0.20000000
                        300 300
                                 1500
                                      1500 5.00000
```

a. What is the death rate in the age interval [65, 85)  $_{20}M_{65}$ ?

```
nMx[x == 65]
```

## [1] 0.03333333

b. What is the value of  $e_{10}$ ?

```
ex[x == 10]
```

## [1] 52.89474

- c. Ask Copilot to find the value of  $e_{10}$ , analysing the data as an expert demographer.
- d. Ask Copilot to calculate the life expectancy at birth for a theoretical population with a constant force of mortality equal to 0.02.
- e. Ask Copilot to calculate  $e_{10}$  for the same theoretical population.