



Following demographic and economic assumptions will be hold:

- $x$ , the retirement age will be set equal to 65 regardless the cohort.
- $m$ , the number of fractional payments per year, will be equal to 12.
- $\ddot{a}_x^{(m)}$  to be the actuarial present value of a yearly annuity of 1 monetary unit. The annuity will be evaluated assuming an interest rate of 4% and an inflation rate of 2%.

The projection has been performed using a mechanical approach, since the purpose of this paper lies in showing the procedure instead of providing sensible results.

Most of this paper is based on the examples provided in [Charpentier \(2012\)](#) and [Charpentier and Dutang \(2013\)](#) online manual.

## 2. Fitting Lee Carter model

Lee Carter original model, [Lee and Carter \(1992\)](#), focuses, as main forecasting methodologies, on the central mortality rates  $m(x, t)$  for age  $x$  in year  $t$  defined as the ratio between the number of deaths  $D(x, t)$ , recorded during the calendar year  $t$  for people aged  $x$ , and the exposure to risk  $E(x, t)$  obtained as the average number of people living during the calendar year  $t$ .

Starting by this sample notation, Lee and Carter (1992) proposed to describe the logarithm of central mortality rates as a linear combination of parameters as expressed by Equation (1):

$$\ln m_{x,t} = a_x + b_x k_t + \varepsilon_{x,t} \quad (1)$$

where  $a_x$  describes the general shape of mortality according to different ages and it represents the logarithm of the geometric mean of empirical mortality rates, averaged over historical years.  $e^{a_x}$  mesure indeed the general shape across age of the mortality schedule.

Furthermore,  $k_t$  reproduces the underlying time trend, while a term  $b_x$  is considered in order to take into account the different effect of time  $t$  at each age.  $b_x$  is assumed to be invariant over time and it explains how rates decline rapidly or slowly in response to change in  $k_t$ .

Finally,  $\varepsilon_{x,t}$  are independent and identical distributed random variables  $N(0, \sigma^2)$  taking into account the age and time specific trends not fully captured by the model.

In the original version, parameters have been estimated by a two-stage process where Singular Value Decomposition (SVD) of the matrix of centered age profiles  $\ln(m_{x,t}) - \hat{a}_x$ , allows a first estimation of parameters  $b_x$  and  $k_t$ .

In order to assure a unique solution for the system of equation of the model, Lee and Carter proposed the following constraints:  $\sum_t k_t = 0$  and  $\sum_x b_x = 1$ .

A second step, based on a refitting of  $\hat{k}_t$  on the number of deaths, is usually suggested in order to assure a better convergence bewteen estimated and observed deaths. The aim is to find the  $\hat{k}_t$  such that  $D(x, t) = E(x, t) \exp(\hat{a}_x + \hat{b}_x \cdot \hat{k}_t)$ .

Alternative frameworks have been proposed over the years in order to improve some drawbacks of original Lee-Carter model (in particular see [? , ? , ? , ? , ? , ? , ?](#))

The one - year survival probability at age  $x$  during calendar year  $t$  is expressed by Equation (2). Equation (2) assumes constant force of mortality to hold between  $[x, x + t)$  and that  $\mu_x \sim m_x$ , that is the force of mortality to be approximated by the central rate of mortality<sup>1</sup>:

$$p_{x,t} = \exp(-\mu_{x,t}) \sim \exp(-m_{x,t}). \quad (2)$$

A longitudinal life table for the cohort of born in calendar year YYYY can be created selecting all  $p_{x,t}$  for which  $t - x = YYYY$ .

We will perform such exercise on Italy HMD data and by applying the original version of Lee-Carter.

```
R> library(demography)
R> library(forecast)
R> library(lifecontingencies)
```

Following code import data from the Human mortality Database and it creates a `demogdata` object from HMD data structure. The `hmd.mx` function downloads all available annual data by single years of age, but for the application we will use the already saved data.

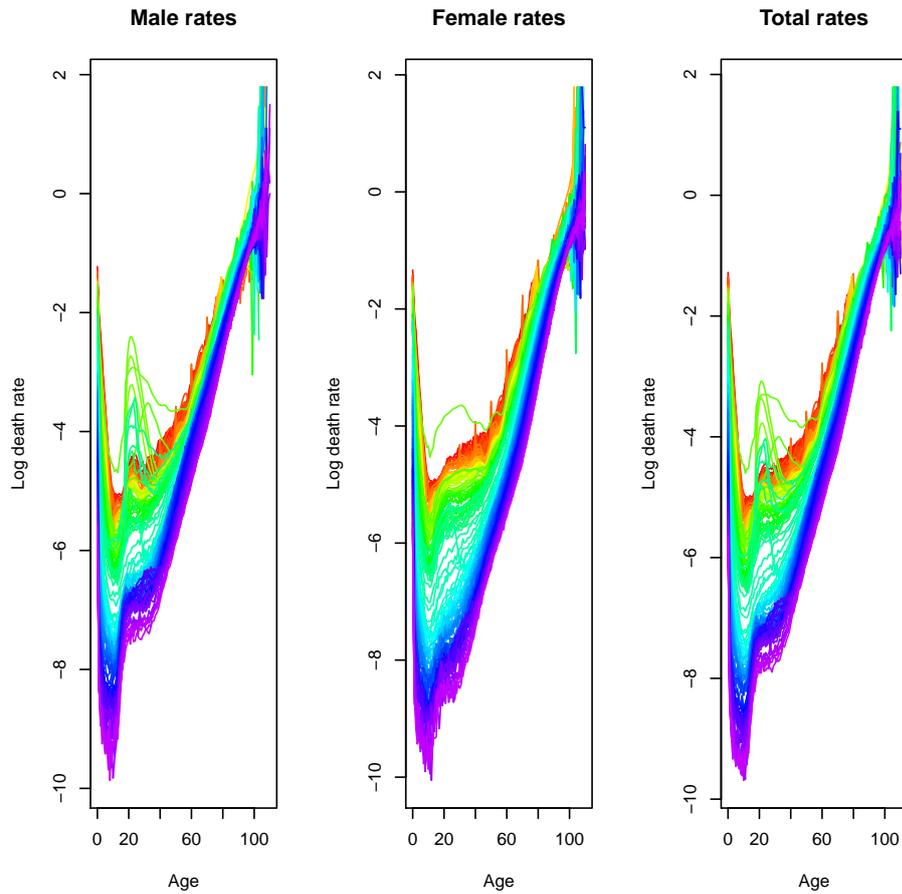
```
R> #italyDemo<-hmd.mx(country="ITA", username="username@email.domain",
R> #password="password", label="Italy")
R> load(file="mortalityDatasets.RData")
```

Plot method is available on `demogdata`. Following figures report for the Italian Population the pattern of logarithm of death rates according to age and time. Several behaviour are shown respectively for male, female and total population.

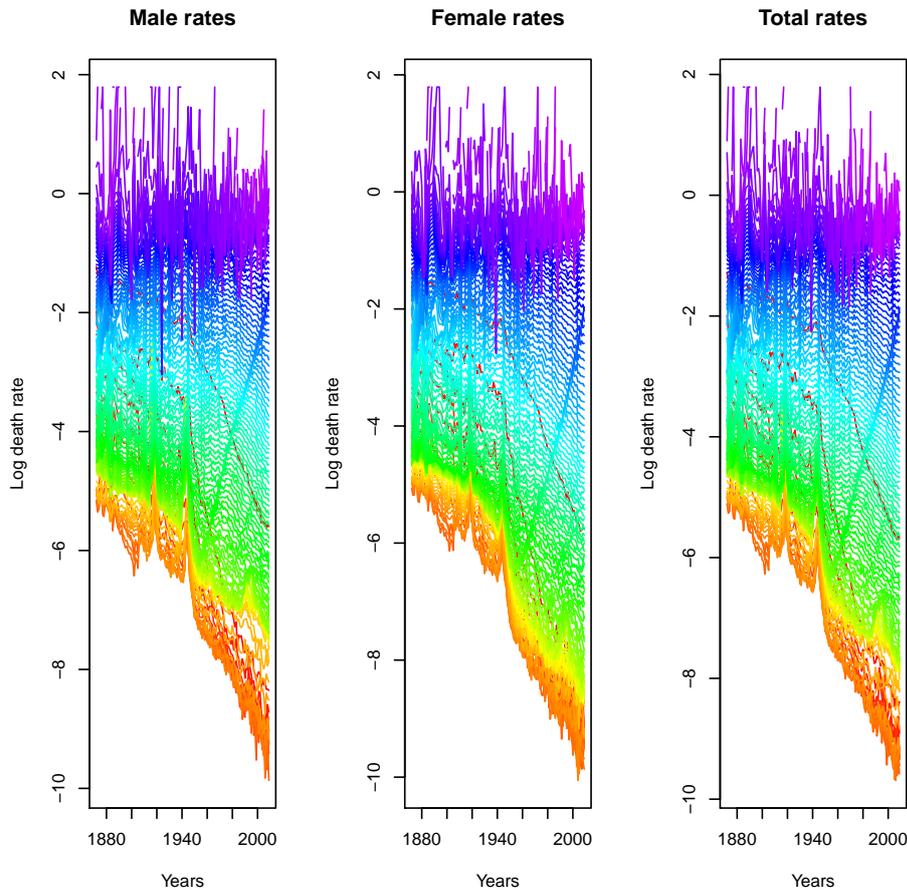
```
R> par(mfrow=c(1,3))
R> plot(italyDemo,series="male",datatype="rate", main="Male rates")
R> plot(italyDemo,series="female",datatype="rate", main="Female rates")
R> plot(italyDemo,"total",datatype="rate", main="Total rates")
```

---

<sup>1</sup>If  $p_{x,t}$  were assumed linear between the two consecutive integer ages, we could write  $m_x = \frac{q_x}{1-\frac{1}{2}q_x}$ .



```
R> par(mfrow=c(1,3))
R> plot(italyDemo,series="male",datatype="rate",
+       plot.type="time", main="Male rates",xlab="Years")
R> plot(italyDemo,series="female",datatype="rate",
+       plot.type="time", main="Female rates",xlab="Years")
R> plot(italyDemo,series="total",datatype="rate",
+       plot.type="time", main="Total rates",xlab="Years")
```



Italian data confirms that mortality is falling at all ages with a different behaviour according to different ages

To fit Lee - Carter model (without going through logarithms) `lca` function can be used. Lee-Carter is here applied separately between male, female and total population and by considering a maximum age equal to 100.

```
R> italyLcaM<-lca(italyDemo,series="male",max.age=100)
R> italyLcaF<-lca(italyDemo,series="female",max.age=100)
R> italyLcaT<-lca(italyDemo,series="total",max.age=100)
```

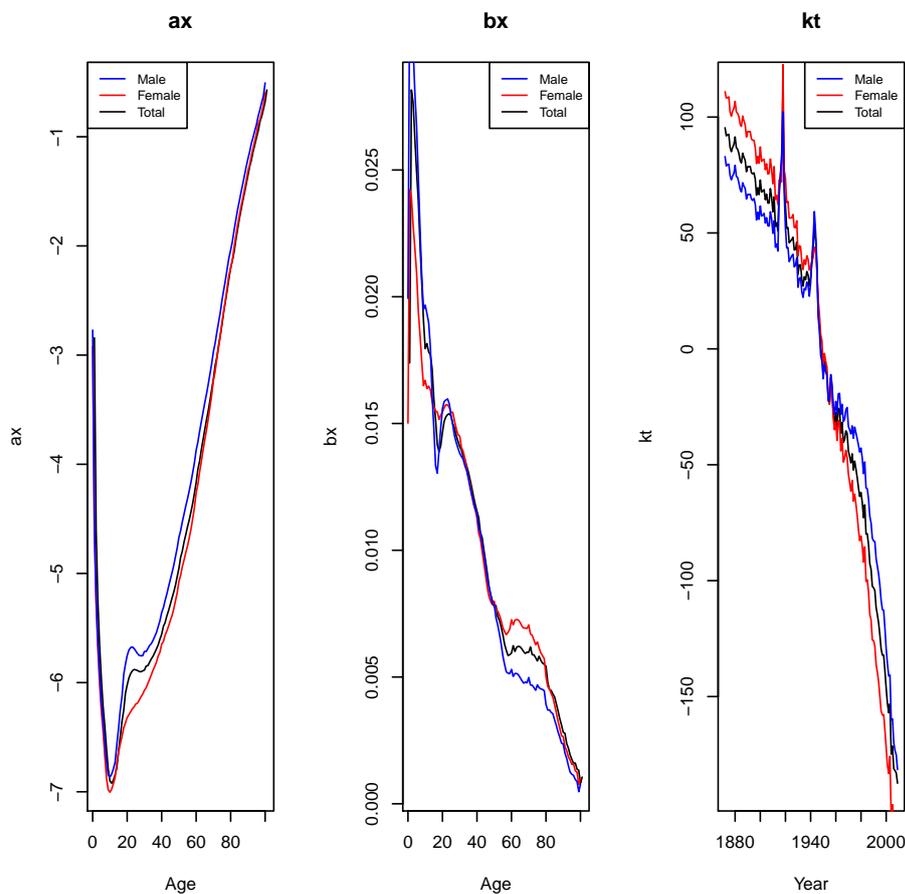
`lca` returned object allows us to inspect  $a_x$ ,  $b_x$  and  $k_t$ . Figures represent the values of the estimated parameters.

```
R> par(mfrow=c(1,3))
R> plot(italyLcaT$ax, main="ax", xlab="Age",ylab="ax",type="l")
R> lines(x=italyLcaF$age, y=italyLcaF$ax, main="ax", col="red")
R> lines(x=italyLcaM$age, y=italyLcaM$ax, main="ax", col="blue")
R> legend("topleft" , c("Male","Female","Total"),
+ cex=0.8,col=c("blue","red","black"),lty=1);
R> plot(italyLcaT$bx, main="bx", xlab="Age",ylab="bx",type="l")
```

```

R> lines(x=italyLcaF$age, y=italyLcaF$bx, main="bx", col="red")
R> lines(x=italyLcaM$age, y=italyLcaM$bx, main="bx", col="blue")
R> legend("topright" , c("Male","Female","Total"),
+ cex=0.8,col=c("blue","red","black"),lty=1);
R> plot(italyLcaT$kt, main="kt", xlab="Year",ylab="kt",type="l")
R> lines(x=italyLcaF$year, y=italyLcaF$kt, main="kt", col="red")
R> lines(x=italyLcaM$year, y=italyLcaM$kt, main="kt", col="blue")
R> legend("topright" , c("Male","Female","Total"),
+ cex=0.8,col=c("blue","red","black"),lty=1);

```



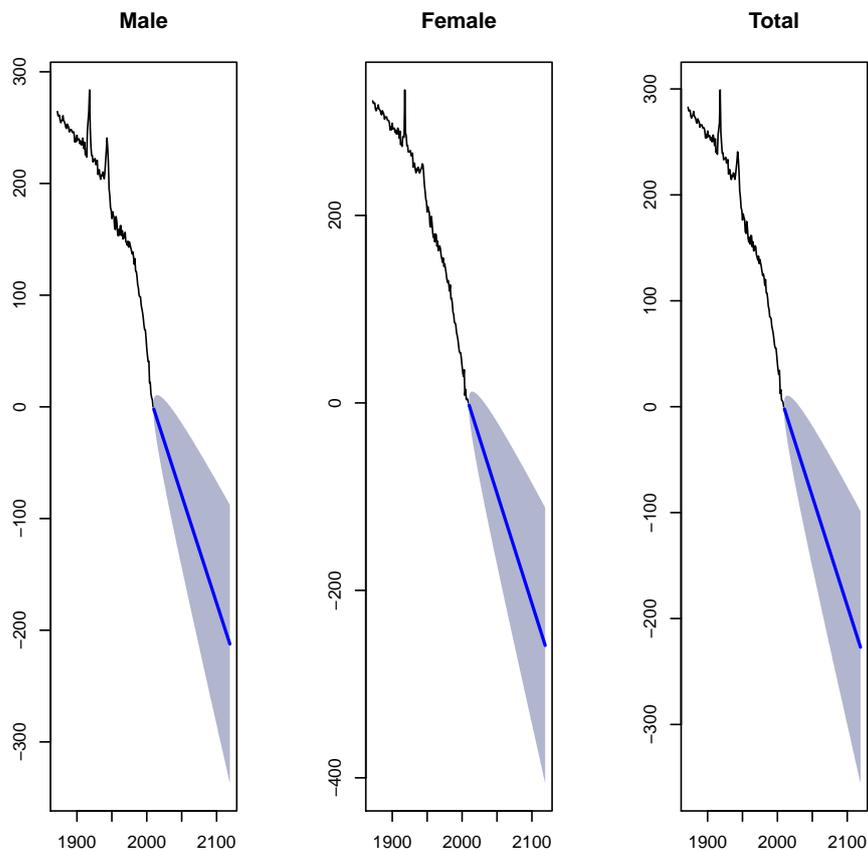
A similar behaviour of parameters is observed according to different data-sets. As expected the average mortality grows when age increases (see  $\hat{a}_x$  pattern). Furthermore it is clearly visible the young mortality hump for males in the age-range (20,30) due to accidental deaths.  $\hat{b}_x$  shows instead a greater value for younger ages and a greatest improvement for females in the age range (60-80). Finally, as expected,  $\hat{k}_t$  has a decreasing trend with the increment of time.

We can therefore use **forecast** package to project the future  $k_t$ s (up to 110). Projection is based on ARIMA extrapolation.

```
R> fM<-forecast(italyLcaM,h=110)
R> fF<-forecast(italyLcaF,h=110)
R> fT<-forecast(italyLcaT,h=110)
```

The predicted values of  $k_t$  rescaled to zero in the last observed year (2009) are here reported.

```
R> par(mfrow=c(1,3))
R> plot(fM$kt.f,main="Male")
R> plot(fF$kt.f,main="Female",)
R> plot(fT$kt.f,main="Total")
```

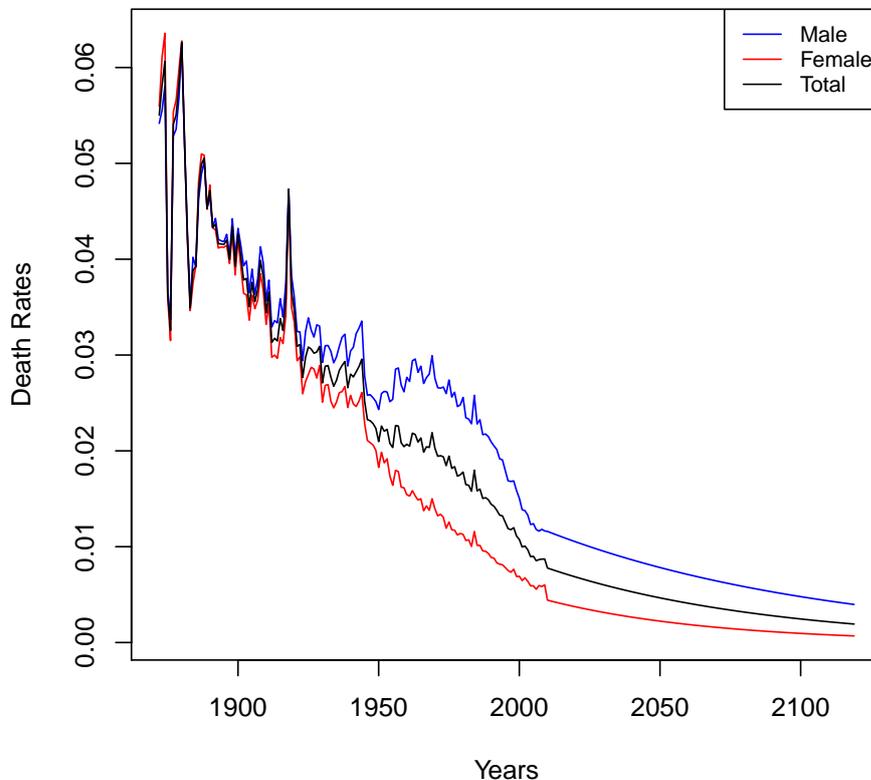


Finally, it's easy to derive the full pattern of rates. Past and forecasted rates are here binded in the same matrix.

```
R> ratesM<-cbind(italyDemo$rate$male[1:100,],fM$rate$male[1:100,])
R> ratesF<-cbind(italyDemo$rate$female[1:100,],fF$rate$female[1:100,])
R> ratesT<-cbind(italyDemo$rate$total[1:100,],fT$rate$total[1:100,])
```

We report here the pattern of past and forecasted rates according to different population for people aged 65. The expected improvement is clearly visible in the Figure.

```
R> par(mfrow=c(1,1))
R> plot(seq(min(italyDemo$year),max(italyDemo$year)+110),ratesF[65,],
+       col="red",xlab="Years",ylab="Death Rates",type="l")
R> lines(seq(min(italyDemo$year),max(italyDemo$year)+110),ratesM[65,],
+        col="blue",xlab="Years",ylab="Death Rates")
R> lines(seq(min(italyDemo$year),max(italyDemo$year)+110),ratesT[65,],
+        col="black",xlab="Years",ylab="Death Rates")
R> legend("topright" , c("Male","Female","Total"),
+        cex=0.8,col=c("blue","red","black"),lty=1);
```



We have applied here the original version of Lee-Carter in order to obtain a forecast of mortality rates. Alternative estimates can be derived by using `lca` function through the **demography** package (as Lee-Miller (?), Booth-Maindonald-Smith (?) and Hyndman-Ullah (?) methods). Finally Lifemetrics package allows to fit ?, ? and ? models.

### 3. Perform actuarial projections

Our aim is to create a function to project life table depending by year of birth, using results from Lee - Carter model. In particular, for ages  $0, 1, \dots, \tau$  on which Lee-Carter model has been fit Equation 3 apply, while for extreme ages,  $\tau+1, \dots, \omega$  on which no data were provided,

it has been assumed that on year probability decreases evenly in 20 steps.

$$\begin{aligned}\ln \hat{\mu}_{x,t} &= \hat{a}_x + \hat{b}_x \hat{k}_t \\ \hat{p}_{x,t} &= \exp(-\hat{\mu}_{x,t})\end{aligned}\quad (3)$$

```
R> createActuarialTable<-function(yearOfBirth,rate){
+
+   mxcoh <- rate[1:nrow(rate),(yearOfBirth-min(italyDemo$year)+1):ncol(rate)]
+   cohort.mx <- diag(mxcoh)
+   cohort.px=exp(-cohort.mx)
+   #get projected Px
+   fittedPx=cohort.px #add px to table
+     px4Completion=seq(from=cohort.px[length(fittedPx)], to=0, length=20)
+     totalPx=c(fittedPx,px4Completion[2:length(px4Completion)])
+     #create life table
+     irate=1.04/1.02-1
+
+     cohortLt=probs2lifetable(probs=totalPx, radix=100000,type="px",
+ name=paste("Cohort",yearOfBirth))
+     cohortAct=new("actuarialtable",x=cohortLt@x, lx=cohortLt@lx,
+ interest=irate, name=cohortLt@name)
+     return(cohortAct)
+   }
R>
R>
```

We can therefore calculate the APV of  $\ddot{a}_{65}^{(12)}$  for the selected cohorts. Values have been derived separately between males and females and by using directly the total population.

```
R> getAnnuityAPV<-function(yearOfBirth,rate) {
+   actuarialTable<-createActuarialTable(yearOfBirth,rate)
+   out=axn(actuarialTable,x=65,m=12)
+   return(out)
+ }
R> rate<-ratesM
R> for(i in seq(1920,2000,by=10)) {
+   cat("For cohort ",i, "of males the e0 is",
+ round(exn(createActuarialTable(i,rate)),2),
+ " and the APV is :",round(getAnnuityAPV(i,rate),2),"\\n")
+ }
R>
```

```
For cohort 1920 of males the e0 is 48.55 and the APV is : 3.92
For cohort 1930 of males the e0 is 60.22 and the APV is : 5.03
For cohort 1940 of males the e0 is 64.36 and the APV is : 5.63
For cohort 1950 of males the e0 is 71.63 and the APV is : 5.92
For cohort 1960 of males the e0 is 74.78 and the APV is : 6.22
```

```

For cohort 1970 of males the e0 is 77.77 and the APV is : 6.52
For cohort 1980 of males the e0 is 80.46 and the APV is : 6.81
For cohort 1990 of males the e0 is 82.24 and the APV is : 7.1
For cohort 2000 of males the e0 is 83.53 and the APV is : 7.38

```

```

R> rate<-ratesF
R> for(i in seq(1920,2000,by=10)) {
+   cat("For cohort ",i, "of females the e0 at birth is",
+   round(exn(createActuarialTable(i,rate)),2),
+   " and the APV is :",round(getAnnuityAPV(i,rate),2),"\\n")
+ }

```

```

For cohort 1920 of females the e0 at birth is 57.38 and the APV is : 6.21
For cohort 1930 of females the e0 at birth is 66.99 and the APV is : 7.23
For cohort 1940 of females the e0 at birth is 71 and the APV is : 7.8
For cohort 1950 of females the e0 at birth is 78.22 and the APV is : 8.24
For cohort 1960 of females the e0 at birth is 81.76 and the APV is : 8.63
For cohort 1970 of females the e0 at birth is 84.54 and the APV is : 8.99
For cohort 1980 of females the e0 at birth is 86.85 and the APV is : 9.33
For cohort 1990 of females the e0 at birth is 88.18 and the APV is : 9.66
For cohort 2000 of females the e0 at birth is 89.27 and the APV is : 9.96

```

```

R> rate<-ratesT
R> for(i in seq(1920,2000,by=10)) {
+   cat("For cohort ",i, "of total population the e0 is",
+   round(exn(createActuarialTable(i,rate)),2),
+   " and the APV is :",round(getAnnuityAPV(i,rate),2),"\\n")
+ }

```

```

For cohort 1920 of total population the e0 is 52.85 and the APV is : 5.17
For cohort 1930 of total population the e0 is 63.6 and the APV is : 6.23
For cohort 1940 of total population the e0 is 67.73 and the APV is : 6.85
For cohort 1950 of total population the e0 is 75.05 and the APV is : 7.24
For cohort 1960 of total population the e0 is 78.42 and the APV is : 7.61
For cohort 1970 of total population the e0 is 81.36 and the APV is : 7.96
For cohort 1980 of total population the e0 is 83.93 and the APV is : 8.31
For cohort 1990 of total population the e0 is 85.55 and the APV is : 8.64
For cohort 2000 of total population the e0 is 86.81 and the APV is : 8.96

```

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## References

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