

# DEMG609: Problem Set 3

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```
library(knitr)
library(data.table)
library(ggplot2)
library(formatR)

# Load data for problem set
raw_data <- fread("C:/Users/ngraetz/Documents/repos/demg609/hw3_data_clean.csv")
raw_data[, x := as.numeric(x)]
raw_data[, nNx := gsub(',', '', nNx)]
raw_data[, nNx := as.numeric(nNx)]
raw_data[, nDx := gsub(',', '', nDx)]
raw_data[, nDx := as.numeric(nDx)]

##### calculate_life_table #####
### Define function that calculates complete life table given arguments:
### data = data.table with numeric columns:
### x = beginning of age interval
### nNx = total people reaching age interval
### nDx = total deaths in age interval
### radix = numbers of survivors at age 0 (or births per year in a
### stationary population); defaults to 100000
#####
calculate_life_table <- function(data, radix = 100000) {
# 1. Calculate nmx
data[, nmx := nDx / nNx]

# 2. Calculate nax given Coale and Demeny equations for ages <5 and n/2 for others.
# For 1a0
data[x == 0 & nmx >= .107, nax := 0.35]
data[x == 0 & nmx < .107, nax := 0.053 + (2.8 * nmx)]
# For 4a1
m0 <- data[x == 0, nmx]
data[x == 1 & shift(nmx, 1, type='lag') >= .107, nax := 1.361]
data[x == 1 & shift(nmx, 1, type='lag') < .107, nax := 1.522 - (1.518 * m0)]
# All other age groups
data[x > 1, nax := 5 / 2]
data[x == 85, nax := 1/nmx]

# 3. Calculate nqx
data[x == 0, n := 1]
data[x == 1, n := 4]
data[x > 1, n := 5]
data[, nqx := (n * nmx) / (1 + ((n - nax) * nmx))]
data[x == 85, nqx := 1]

# 4. Calculate npx
data[, npx := 1 - nqx]

# 5. Calculate lx
data[, lx := radix]
for(r in 2:length(data[, lx])) {
```

```

    previous_lx <- data[r-1, lx]
    previous_npx <- data[r-1, npx]
    data[r, lx := previous_lx * previous_npx]
  }

# 6. Calculate ndx
for(r in 1:length(data[, lx])) {
  lx_n <- data[r+1, lx]
  data[r, ndx := lx - lx_n]
  if(r == length(data[, lx])) data[r, ndx := lx]
}

# 7. Calculate nLx
for(r in 1:length(data[, lx])) {
  lx_n <- data[r+1, lx]
  data[r, nLx := (n * lx_n) + (nax * ndx)]
  if(r == length(data[, lx])) data[r, nLx := lx / nm]
}

# 8. Calculate Tx
for(r in 1:length(data[, lx])) {
  data[r:length(data[, lx]), Tx := sum(nLx)]
}

# 9. Calculate ex
data[, ex := Tx / lx]

# Return complete life table
return(data)
}

# Calculate standard life table with data for problem set and default radix
lt <- calculate_life_table(data = raw_data)

```

A.(1)

```

# A.(1)
lt[x == 0, ex]

```

```
## [1] 77.47132
```

A.(2)

```

# A.(2)
lt[x == 35, ex]

```

```
## [1] 44.2476
```

A.(3)

```

# A.(3)
# Given an individual survived until age 35, the expected value

```

```
# of the remaining length of their life in years is 44.2476
```

A.(4)

```
# A.(4)  
prod(lt[x < 25, npx])  
  
## [1] 0.9791639  
lt[x == 25, lx] / lt[1, lx]  
  
## [1] 0.9791639
```

A.(5)

```
# A.(5)  
1 - prod(lt[x >= 25 & x < 50, npx])  
  
## [1] 0.0241032
```

A.(6)

```
# A.(6)  
sum(lt[x >= 15 & x < 65, nLx]) / lt[1, lx]  
  
## [1] 47.9629  
50 - sum(lt[x >= 15 & x < 65, nLx]) / lt[1, lx]  
  
## [1] 2.037096
```

A.(7)

```
# A.(7)  
lt[, nAx := nax * ndx]  
lt[x == 1, nax]  
  
## [1] 1.501346
```

A.(8)

```
# A.(8) = probability of surviving to age 65 * probability of dying between 65-69  
prod(lt[x < 65, npx]) * lt[x == 65, nqx]  
  
## [1] 0.06768248
```

A.(9)

```
# A.(9) probability(30-60) * probability(0-30)  
prod(lt[x < 30, npx]) * prod(lt[x >= 30 & x < 60, npx])
```

```
## [1] 0.916966
  (lt[x == 30, lx] / lt[x == 0, lx]) * (lt[x == 60, lx] / lt[x == 30, lx])

## [1] 0.916966
```

A.(10)

```
# A.(10)
# CBR = CDR
1 / lt[1, ex]
```

```
## [1] 0.012908
# Death rate above 60
1 / lt[x == 60, ex]
```

```
## [1] 0.04769172
# Mean age at death
lt[1, ex]
```

```
## [1] 77.47132
# Given 56,059 births per year in this population, how many people turn 65 each year?
stationary_lt <- calculate_life_table(data = copy(raw_data),
                                     radix = 56059)
stationary_lt[x == 65, lx]
```

```
## [1] 49272.9
```