Method: Life tables for Denmark for the years 1995 and 2012 were constructed from the numbers of deaths and population counts by single year of age, sex and calendar year, using Poisson regression and flexible functions. Life tables for intervening years were constructed by linear interpolation. See article in The Lancet (http://dx.doi.org/10.1016/S0140-6736(14)62038-9) for more details.
Source: Statistics Denmark (Danmarks Statistik)
Registry population (2009): 5,524,430

## Year 1995

Table 1: Summary statistics

| Sex | Life expectancy <br> at birth (years) | Probability (\%) of dying between exact ages |  |  |
| :--- | ---: | ---: | ---: | ---: |
|  | 72.8 | $\mathbf{1 5 - 6 0}$ years | $\mathbf{6 0 - 8 5}$ years | $\mathbf{8 5 - 9 9}$ years |
| Female | 78.1 | 14.7 | 77.1 | 98.5 |

Figure 1: Observed and fitted mortality rates (mx; logarithmic scale) by age and sex


Figure 2: Observed and fitted mortality rates (mx; arithmetic scale) by broad age band and sex


Figure 3: Fitted versus observed mortality rates: residual deviance
We plotted the deviance residuals of the model against age to assess goodness of fit. Deviance residuals are a measure of how closely the modelled values fit the observed data. They should be approximately normally distributed if the model fits the data well. The ideal range is between -2 and +2 (the red lines).


Table 1: Summary statistics

| Sex | Life expectancy <br> at birth (years) | Probability (\%) of dying between exact ages |  |  |
| :--- | ---: | ---: | ---: | ---: |
|  | 77.9 | $\mathbf{1 5 - 6 0}$ years | $\mathbf{6 0 - 8 5}$ years | $\mathbf{8 5 - 9 9}$ years |
| Female | 82.0 | 9.0 | 62.5 | 97.3 |

Figure 1: Observed and fitted mortality rates (mx; logarithmic scale) by age and sex


Figure 2: Observed and fitted mortality rates (mx; arithmetic scale) by broad age band and sex


Figure 3: Fitted versus observed mortality rates: residual deviance
We plotted the deviance residuals of the model against age to assess goodness of fit. Deviance residuals are a measure of how closely the modelled values fit the observed data. They should be approximately normally distributed if the model fits the data well. The ideal range is between -2 and +2 (the red lines).


