

# MortalityGaps R Package

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*2018-07-17*

This package contains source code for the Double-Gap model for forecasting life expectancy in human populations.

## Description

Life expectancy is highly correlated over time among countries and between males and females. These associations can be used to improve forecasts. Here we have implemented a method for forecasting female life expectancy based on analysis of the gap between female life expectancy in a country compared with the record level of female life expectancy in the world. Second, to forecast male life expectancy, the gap between male life expectancy and female life expectancy in a country is analysed. We named this method the Double-Gap model. For a detailed description of the method see Pascariu et al. (2017).

## Installation

1. Make sure you have the most recent version of R
2. Run the following code in your R console

```
install.packages("MortalityGaps")
```

## Updating to the latest version of the package

You can track and contribute to the development of `MortalityGaps` on GitHub. To install it:

1. Install the release version of `devtools` from CRAN with `install.packages("devtools")`.
2. Make sure you have a working development environment.
  - **Windows:** Install Rtools.
  - **Mac:** Install Xcode from the Mac App Store.
  - **Linux:** Install a compiler and various development libraries (details vary across different flavors of Linux).
3. Install the development version of `MortalityGaps`.

```
R devtools::install_github("mpascariu/MortalityGaps")
```

## Help

All functions are documented in the standard way, which means that once you load the package using `library(MortalityGaps)` you can just type `?DoubleGap` to see the help file.

## Examples

```
library(MortalityGaps)
```

```
##  
## MortalityGaps: The Double-Gap Life Expectancy Forecasting Model  
## Author      : Marius D. Pascariu  
## Last Update : June 20, 2018
```

### Input data

```
# Collection of life expectancies for female populations  
exF <- MortalityGaps.data$exF  
# Life expectancy for male populations  
exM <- MortalityGaps.data$exM
```

```
head(exF)
```

```
##   country Year Age   ex  
## 1     AUS 1950  0 71.72  
## 2     AUS 1950 65 14.74  
## 3     AUS 1951  0 71.59  
## 4     AUS 1951 65 14.66  
## 5     AUS 1952  0 72.04  
## 6     AUS 1952 65 14.89
```

### Fit DG model at age 0 for Australia using data from 1950 to 2014

```
M0 <- DoubleGap(DF = exF,  
               DM = exM,  
               age = 0,  
               country = "AUS",  
               years = 1950:2014)
```

```
M0
```

```
## Double-Gap Model fit  
##  
## Country      : AUS  
## Age (x)      : 0  
## Years in fit: 1950 - 2014
```

### Summary results

```
summary(M0)
```

```
##  
## Coefficients Double-Gap Model:  
##  
## M1: Best-Practice Life Expectancy Model  
##           Estimate Std. Error t value Pr(>|t|)  
## (Intercept) 73.5182019  0.1190168  617.71 < 2.2e-16 ***
```

```

## year          0.2072107  0.0031353  66.09 < 2.2e-16 ***
##
## M2: Best-Practice Gap Model (ARIMA)
##      ar1
## -0.4255166
##
## M3: Sex-Gap Model
##           Estimate Std. Error  z value  Pr(>|z|)
## (Intercept)  0.1929436  0.0237540   8.1226 4.564e-16 ***
## sex_gap1     0.8315822  0.0210144  39.5720 < 2.2e-16 ***
## sex_gap2     0.1495723  0.0208707   7.1666 7.687e-13 ***
## narrow_level -0.0342501  0.0029929 -11.4439 < 2.2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## tau = 77.28 | A = 87.52275 | L = 2.24 | U = 13.68

```

### Forecast life expectancy in Australia until 2050

```
P0 <- predict(M0, h = 36)
```

### Plot the results

```
plot(P0)
```

