# Example 2.5

Disease mapping: from foundations to multidimensional modeling

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This document reproduces the analysis made at Example 2.5 of the book: "Disease mapping: from foundations to multidimensional modeling" by Martinez-Beneito M.A. and Botella-Rocamora P., published by CRC press in 2019. You can watch the analysis made with full detail at this pdf document, or even execute it if you want with the material available at https://github.com/MigueBeneito/DMBook. Anyway, this pdf file should be enough for following most of the details of the analysis made for this example.

The statistical analysis below has been run in R, by additionally using the library Rmarkdown, so be sure that you have this software installed if you want to reproduce by yourself the content of this document. In that case we advise you to download first the annex material at https://github.com/MigueBeneito/DMBook, open with Rstudio the corresponding .Rproj file that you will find at the folder corresponding to this example and compile the corresponding .Rmd document. This will allow you to reproduce the whole statistical analysis below.

The inference carried out in this example has been undertaken with WinBUGS, that will be introduced in the next chapter. The details for fully understanding this code are given at Chapter 3, nevertheless you may find interesting to take it a look at this code now, or later when you read Chapter 3.

## Libraries and data loading

#### Data preparation

```
# data preparation
year = 1991:2011
year.centered = year - mean(year)

rates = 1e+05 * 0/Pop
```

## MCMC sampling with WinBUGS (centered covariate)

```
RegLin2 = function() {
   for (i in 1:n) {
       rate[i] ~ dnorm(media[i], prec)
       media[i] <- beta1 + beta2 * year[i]</pre>
   }
   prec <- pow(sigma, -2)</pre>
   sigma ~ dunif(0, 1000)
   beta1 ~ dt(0, 1e-06, 2)
   beta2 ~ dt(0, 1e-06, 2)
}
# WinBUGS call for making inference on the model above
# Data
data = list(year = year.centered, rate = rates, n = 21)
# Initial values
inits = function() {
   list(beta1 = rnorm(1, 0, 10), beta2 = rnorm(1, 0, 10), sigma = runif(1,
       0, 10))
}
# Parameters to save
parameters = c("beta1", "beta2", "sigma")
# WinBUGS call
RegLin2WB = pbugs(data = data, inits = inits, param = parameters, model = RegLin2,
   bugs.seed = 1, DIC = F, n.iter = 5500, n.burnin = 500, n.chains = 3,
   n.thin = 1)
# Posterior summaries
round(RegLin2WB$summary, 3)
                      2.5%
                                25%
                                       50%
                                              75% 97.5% Rhat n.eff
           mean
                   sd
## beta1 7.760 0.158 7.446 7.658 7.759 7.864 8.069 1.001 15000
## beta2 -0.124 0.026 -0.176 -0.141 -0.124 -0.107 -0.073 1.001 15000
## sigma 0.717 0.127 0.516 0.627 0.700 0.788 1.010 1.001 5300
```

## MCMC sampling with WinBUGS (uncentered covariate)

```
# Data
data = list(year = year, rate = rates, n = 21)

# Initial values
inits = function() {
    list(beta1 = rnorm(1, 500, 20), beta2 = rnorm(1, -0.1, 0.1), sigma = runif(1, 0, 4))
}

# WinBUGS call
```

```
RegLin2WB.uncentered = pbugs(data = data, inits = inits, param = parameters,
    model = RegLin2, bugs.seed = 1, DIC = F, n.iter = 5500, n.burnin = 500,
   n.chains = 3, n.thin = 1)
# Posterior summaries
round(RegLin2WB.uncentered$summary, 3)
                                                             97.5%
##
                      sd
                             2.5%
                                       25%
                                               50%
                                                       75%
                                                                     Rhat
           mean
## beta1 107.783 277.716 -251.700 -214.100 100.400 444.300 462.700 46.294
## beta2 -0.050
                 0.139
                          -0.227
                                    -0.218 -0.046
                                                     0.111
                                                             0.130 46.301
## sigma
         1.191
                  0.450
                            0.668
                                     0.843
                                             1.008
                                                     1.534
                                                             2.209 3.400
        n.eff
## beta1
            3
## beta2
            3
## sigma
```

## Long MCMC sampling with WinBUGS (uncentered covariate)

```
# WinBUGS call
RegLin2WB.uncentered2 = pbugs(data = data, inits = inits, param = parameters,
   model = RegLin2, bugs.seed = 1, DIC = F, n.iter = 2e+06, n.burnin = 2e+05,
   n.chains = 3, n.thin = 360)
# Posterior summaries
round(RegLin2WB.uncentered2$summary, 3)
##
                                    25%
                                                   75%
                                                          97.5% Rhat n.eff
            mean
                    sd
                          2.5%
                                            50%
## beta1 257.549 53.600 145.800 221.000 255.700 291.400 364.500 1.074
## beta2 -0.125 0.027 -0.178 -0.142 -0.124 -0.107 -0.069 1.068
                                                                         34
## sigma
          0.721 0.128
                         0.523
                                 0.632
                                         0.704
                                                 0.791
                                                         1.019 1.001 15000
cor(RegLin2WB.uncentered2$sims.list$beta1, RegLin2WB.uncentered2$sims.list$beta2)
## [1] -0.999995
```

## Convergence plots

