Solution Exercice #3, Série 5

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```
Activer les librairies utiles.
library(here)
library(tidyverse)
library(glue)
library(magrittr)
library(glmnet)
Lire la base de données credit.csv.
credit <- load(here("0_data", "freMPL3.rda"))</pre>
\mathbf{a})
data <- freMPL3 %>% filter(ClaimAmount > 0)
b)
gamma_fit <- glm(</pre>
  ClaimAmount ~ LicAge + VehAge + Gender + MariStat + SocioCateg + DrivAge,
  family = Gamma(link = "log"),
  data = data,
  offset = log(Exposure)
)
summary(gamma_fit)
##
## Call:
  glm(formula = ClaimAmount ~ LicAge + VehAge + Gender + MariStat +
       SocioCateg + DrivAge, family = Gamma(link = "log"), data = data,
##
##
       offset = log(Exposure))
##
##
  Deviance Residuals:
##
       Min
                 1Q
                      Median
                                    3Q
                                            Max
##
  -3.8065 -1.1024 -0.5599
                                0.1153
                                         6.0984
##
## Coefficients:
                     Estimate Std. Error t value Pr(>|t|)
##
## (Intercept)
                    7.5710910 0.3240725 23.362 < 2e-16 ***
## LicAge
                   -0.0008203 0.0007727
                                           -1.062 0.288630
## VehAge1
                    0.1294967 0.1574067
                                            0.823 0.410847
                   -0.1471522 0.3170181
                                           -0.464 0.642605
## VehAge10+
## VehAge2
                   -0.0980039 0.1566142 -0.626 0.531586
## VehAge3
                   -0.0448791 0.1695603 -0.265 0.791302
## VehAge4
                   -0.1953874 0.1713965 -1.140 0.254522
## VehAge5
                    0.7147311 0.1907686
                                            3.747 0.000188 ***
## VehAge6-7
                    0.4353038 0.1915252
                                            2.273 0.023210 *
## VehAge8-9
                    0.2439549 0.2466899 0.989 0.322902
```

GenderMale

```
## MariStatOther
                   0.2287443 0.1209635
                                         1.891 0.058859
## SocioCategCSP20 0.0255151 0.8166161 0.031 0.975079
## SocioCategCSP21 -1.4977180 1.6061031 -0.933 0.351255
## SocioCategCSP22 -0.6301074 1.1438103 -0.551 0.581814
## SocioCategCSP26 0.4710595 0.3158833
                                         1.491 0.136156
## SocioCategCSP37 0.0138807 0.4183706 0.033 0.973538
## SocioCategCSP42 -0.1137252 0.3396833 -0.335 0.737835
## SocioCategCSP46 0.0694105 0.3664637
                                          0.189 0.849806
## SocioCategCSP47 -1.1527185 1.1430011 -1.009 0.313414
## SocioCategCSP48 0.1746436 0.3388862
                                          0.515 0.606405
## SocioCategCSP49 -0.1603568  0.4584226 -0.350  0.726548
## SocioCategCSP50 0.3269536 0.2057376
                                         1.589 0.112280
## SocioCategCSP55 0.1298638 0.2479113 0.524 0.600491
## SocioCategCSP6 -0.8686871 0.7359607 -1.180 0.238094
## SocioCategCSP60 -0.0116380 0.2485941 -0.047 0.962668
## SocioCategCSP65 -0.4241183 1.1552812 -0.367 0.713599
## SocioCategCSP66 0.0827468 0.3305943
                                         0.250 0.802400
## SocioCategCSP74 -3.4982896 1.1484951
                                         -3.046 0.002369 **
## SocioCategCSP77 -2.1169095 0.8202913 -2.581 0.009977 **
## DrivAge
                   0.0137474 0.0091838
                                         1.497 0.134671
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
   (Dispersion parameter for Gamma family taken to be 2.500965)
##
       Null deviance: 1882.1 on 1246 degrees of freedom
##
## Residual deviance: 1721.3 on 1216 degrees of freedom
## AIC: 21928
##
## Number of Fisher Scoring iterations: 9
i)
assure_df <- tibble(
  LicAge = 400,
  VehAge = factor(3),
  Gender = factor("Male"),
 MariStat = factor("Alone"),
  SocioCateg = factor("CSP50"),
  DrivAge = 45,
  Exposure = 1
)
as.numeric(predict(gamma_fit, newdata = assure_df, type = "response"))
## [1] 3553.833
ii)
Cette affirmation est fort probablement fausse puisque la valeur-p associée au paramètre DrivAge est trop élevée.
```

Premièrement, créer une base de données avec seulement les variables Exposure, ClaimAmount et DrivAge.

 \mathbf{c}

data_drv_age <- data %>% select(ClaimAmount, Exposure, DrivAge)

i)

Créer une fonction qui prend en entrée la base df et qui renvoie la même base de données avec les k puissances de la variable var.

```
ajout_puissances <- function(df, var, k) {
   if (k == 1) {
      return(df)
   }

new_vars <- map_dfc(2:k, ~ df[[var]] ^ .x) %>%
      setNames(glue("{var}_{2:k}"))

res <- bind_cols(df, new_vars)
   return(res)
}</pre>
```

Ajuster le GLM gamma pour K = 1, ..., 10.

Calculer l'AIC pour chaque modèle ajuster et choisir K tel que l'AIC est le plus petit.

```
(AICs <- map_dbl(gamma_fit_ls, AIC))

## [1] 21327.55 21329.20 21328.50 21330.44 21332.19 21334.04 21336.03

## [8] 21334.10 21333.05 21319.97

which.min(AICs)
```

[1] 10

Avec le critère AIC, on choisit donc K = 10.

ii)

Premièrement, créer une fonction qui prend en entrée la base de données, la variable explicative et le nombre de puissances k et renvoie l'erreur quadratique moyenne de validation croisée.

```
mse_2_folds_gamma <- function(df, var, k) {
    dat <- ajout_puissances(df, var, k)

folds <- seq(1, nrow(dat)) %>%
    cut(breaks = 2, labels = F) %>%
    sample()

responses_ls <- dat %>%
    mutate(folds = folds) %>%
    group_split(folds) %>%
    group_split(folds) %>%
    map(~ pull(., ClaimAmount))

gamma_fit_ls <- map(
    1:2,
    ~ glm(ClaimAmount ~ ., family = Gamma(link = "log"), data = dat[folds != .x, ], offset = log(Exposure))
)

predictions_ls <- map(1:2, ~ predict(gamma_fit_ls[[.x]], newdata = dat[folds == .x, ], type = "response"))</pre>
```

```
res <- map2(responses_ls, predictions_ls, ~ mean((.x - .y) ^ 2)) %>%
  flatten_dbl() %>%
  mean()

return(res)
}
```

Ensuite, ajuster les modèles pour K = 1, ..., 10 et regarder quelle valeur mène à la plus petite erreur quadratique moyenne.

```
(MSEs <- map_dbl(1:10, ~ mse_2_folds_gamma(data_drv_age, var = "DrivAge", k = .x)))
## [1] 4.312187e+06 4.300823e+06 4.301520e+06 4.319982e+06 4.311656e+06
```

```
## [1] 4.312187e+06 4.300823e+06 4.301520e+06 4.319982e+06 4.311656e+06
## [6] 4.329723e+06 4.317668e+06 4.386675e+06 2.145221e+20 4.306240e+06
```

which.min(MSEs)

[1] 2

Avec le critère de l'erreur quadratique moyenne de 2-validation croisée, on choisit donc K=2.

 \mathbf{d}

```
ridge_cv <- cv.glmnet(
    x = as.matrix(data[c("DrivAge", "LicAge")]),
    y = as.matrix(data["ClaimAmount"]),
    nfolds = 10,
    alpha = 0,
    family = "poisson",
    offset = log(as.matrix(data["Exposure"]))
)

coef(ridge_cv)</pre>
```

```
## 3 x 1 sparse Matrix of class "dgCMatrix"

## 1

## (Intercept) 8.042976e+00

## DrivAge 1.168809e-36

## LicAge 2.096173e-38

ridge_cv$lambda.min
```

[1] 18373.48

La valeur de lambda sélectionnée est 1.8373481×10^4 .

 \mathbf{e})

```
ridge_cv_2 <- cv.glmnet(
    x = model.matrix(~ VehUsage - 1, data = data),
    y = as.matrix(data["ClaimAmount"]),
    nfolds = 20,
    alpha = 0,
    family = "poisson",
    offset = log(as.matrix(data["Exposure"]))
)</pre>
```

Les valeurs des paramètres obtenus sont:

```
coef(ridge_cv_2)
```