Trabalho7

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'C:/Users/Marcelo/OneDrive/Área de Trabalho/ts/trabalho 2 - TS/Trabalho7' Função de Verossimilhança da Professora:

```
[25]: Like <- function(Parametros, y, nb, np, nq, lamb, x, Ind) {
        ny <- length(y)</pre>
        # Separating parameters: beta, fi, theta
        if (nb > 0) {
          beta <- Parametros[1:nb]</pre>
        ar_ma_start <- nb + 1
        if (np > 0 && nq == 0) {
          fi <- Parametros[ar_ma_start:(ar_ma_start + np - 1)]</pre>
        if (nq > 0 && np == 0) {
          theta <- Parametros[ar_ma_start:(ar_ma_start + nq - 1)]
        }
        if (np > 0 && nq > 0) {
          fi <- Parametros[ar_ma_start:(ar_ma_start + np - 1)]</pre>
          theta <- Parametros[(ar_ma_start + np):(ar_ma_start + np + nq - 1)]</pre>
        }
        # Initialize variables
        z < - rep(0, ny)
```

```
w < - rep(0, ny)
mi <- rep(0, ny)
erro <- rep(0, ny)
# First observation: compute w[1]
if (nb == 0) {
 w[1] <- 0
} else if (nb == 1) {
 w[1] <- beta[1]
} else {
 w[1] <- beta[1] + as.numeric(x[1, 1:(nb - 1)] %*% beta[2:nb])
mi[1] < -exp(w[1])
erro[1] \leftarrow (y[1] - mi[1]) / (mi[1]^lamb)
      <-y[1] * w[1] - exp(w[1])
# Loop over t = 2 to ny
for (t in 2:ny) {
  # AR component
 if (np > 0) {
    for (k in 1:min(np, t - 1)) {
      z[t] <- z[t] + fi[k] * (z[t - k] + erro[t - k])
  }
  # MA component
  if (nq > 0) {
   for (k in 1:min(nq, t - 1)) {
      z[t] \leftarrow z[t] + theta[k] * erro[t - k]
  }
  # Compute w[t] based on z[t] and x[t,]
  if (nb == 0) {
   w[t] <_{-} z[t]
  } else if (nb == 1) {
    w[t] \leftarrow beta[1] + z[t]
    w[t] \leftarrow beta[1] + as.numeric(x[t, 1:(nb - 1)] %*% beta[2:nb]) + z[t]
 mi[t] < -exp(w[t])
  erro[t] <- (y[t] - mi[t]) / (mi[t]^lamb)
         <-L + y[t] * w[t] - exp(w[t])
  L
}
```

```
L <- -L # negative log-likelihood

valores <- cbind(erro, z, mi)

if (Ind == 0) return(L)
  if (Ind == 1) return(valores)
}</pre>
```

Função que adiciona faz a estimação, calcula o AIC dado o numero de bases para cada covariável não linear:

```
[26]: MLE_com_splines <- function(y, z_list, k_list,</pre>
                                   other_covariates = NULL,
                                   np = 0, nq = 0, lamb = 0.5,
                                   Ind = 0) {
        nz <- length(z_list)</pre>
         # Construindo as bases splines com seus respectivos ks.
        spline_list <- list()</pre>
        spline_sizes <- integer(nz)</pre>
        for (i in seq_along(z_list)) {
          k_i <- k_list[[i]]
           spline_i <- ns(z_{list[[i]]}, df = k_i) #bases splines para a variável n\tilde{a}o_{l}
        \rightarrow linear numero i
           spline_list[[i]] <- spline_i #lista com as bases</pre>
           spline_sizes[i] <- ncol(spline_i) #numero de bases a variavel não linear i
        }
         # juntando as bases splines usando cbind:
        names(spline_list) <- names(z_list)</pre>
        spline_basis <- do.call(cbind, spline_list)</pre>
        n_spline <- sum(spline_sizes)</pre>
         # Juntando com as outras covariáveis:
        if (!is.null(other_covariates)) {
           x <- cbind(spline_basis, other_covariates)</pre>
          n_other <- ncol(other_covariates)</pre>
        } else {
           x <- spline_basis
          n_other <- 0
        }
        nb <- 1 + n_spline + n_other # intercepto + splines + numero betas das outras_
        ⇔coυ
```

```
n_params <- nb + np + nq
start_vals <- rep(0, n_params)</pre>
 # ajustando o modelo dado um numero fixo de bases para cada covariável n 	ilde{a} 	ilde{o}_{\sqcup}
\rightarrow linear:
opt <- optim(par = start_vals,</pre>
               fn = Like,
               y = y,
               nb = nb,
               np = np,
               nq = nq,
               lamb = lamb,
               x = x,
               Ind = Ind,
               method = "BFGS",
               hessian = FALSE)
logLik <- -opt$value</pre>
AIC <-2 * n_params - 2 * logLik
best_coef <- opt$par</pre>
beta0 <- best_coef[1]</pre>
beta_spline_all <- best_coef[2:(1 + n_spline)]</pre>
 # separando os coeficientes das bases splines para cada variável não linear:
beta_spline_list <- list()</pre>
idx <- 1
for (i in seq_along(z_list)) {
  varname <- names(z_list)[i]</pre>
   len <- spline_sizes[i]</pre>
  beta_spline_list[[varname]] <- beta_spline_all[idx:(idx + len - 1)]</pre>
  idx <- idx + len
}
 # separando os outros coeficientes:
beta_other <- if (n_other > 0)
     {best_coef[(2 + n_spline):(1 + n_spline + n_other)]}
                else {numeric(0)}
fi <- if (np > 0)
     \{best\_coef[(nb + 1):(nb + np)]\}
                else {numeric(0)}
theta < if (nq > 0)
     \{best\_coef[(nb + np + 1):(nb + np + nq)]\}
      else {numeric(0)}
```

```
fit <- list(
   AIC = AIC,
   logLik = logLik,
   beta0 = beta0,
   beta_spline = beta_spline_list,
   beta_other = beta_other,
   fi = fi,
   theta = theta,
   k_list = k_list
)

return(fit)
}</pre>
```

```
[27]: select_best_k_multi <- function(y, z_list, k_ranges,
                                        other_covariates = NULL,
                                        np = 0, nq = 0, lamb = 0.5,
                                        Ind = 0) {
        # Gerando um grid com os possíveis numeros de bases para cada variável spline:
        k_grid <- expand.grid(k_ranges)</pre>
        best_AIC <- Inf
        best_fit <- NULL</pre>
        best_k_list <- NULL</pre>
       #calculando o AIC para cada combinação de número de bases:
        for (i in seq_len(nrow(k_grid)))
        {
          current_k <- as.list(k_grid[i, ])</pre>
          result <- tryCatch({</pre>
                                  MLE_com_splines(
                                    y = y,
                                    z_list = z_list,
                                    k_list = current_k,
                                    other_covariates = other_covariates,
                                    np = np,
                                    nq = nq,
                                    lamb = lamb,
                                    Ind = Ind
                                },
                               error = function(e) {return(NULL)}
```

```
if (!is.null(result) && result$AIC < best_AIC) {
    best_AIC <- result$AIC
    best_fit <- result
    best_k_list <- current_k
    }
}
best_fit$k_list <- best_k_list
    return(best_fit)
}</pre>
```

Pegando o dataset de Doença respiratórias:

```
[28]: df <- read.csv("DR.csv",header = TRUE,sep=';')
```

[29]: summary(df)

```
tempmed
                                                         03
   Data
                   Atendimentos
Length:72
                         : 1.0
                                 Length:72
                                                    Length:72
Class :character
                  1st Qu.:12.0
                                 Class :character
                                                    Class :character
Mode :character
                  Median:21.5
                                 Mode :character
                                                    Mode :character
                  Mean :24.1
                  3rd Qu.:33.0
                  Max. :91.0
    CO
                      NO2
                                         S02
Length:72
                  Length:72
                                     Length:72
Class :character
                  Class :character
                                     Class : character
Mode :character
                  Mode :character
                                     Mode :character
```

```
[30]: df<- df %>% select(Atendimentos, tempmed, 03, C0, N02, S02)
```

Limpando os dados:

```
[31]: df[] <- lapply(df, function(x) {
   if (is.character(x)) as.numeric(gsub(",", ".", x)) else x
})</pre>
```

[32]: summary(df)

```
{\tt tempmed}
 Atendimentos
                                    03
                                                    CO
Min. : 1.0
                                     :16.76
                                              Min. : 295.2
              Min.
                     :18.85
                              Min.
1st Qu.:12.0
               1st Qu.:22.61
                              1st Qu.:26.34
                                              1st Qu.: 645.7
Median:21.5
              Median :23.97
                              Median :32.22
                                              Median: 829.8
Mean :24.1
                     :24.17
                                     :32.72
                                              Mean : 826.8
              Mean
                              Mean
3rd Qu.:33.0
               3rd Qu.:26.02
                              3rd Qu.:38.44
                                              3rd Qu.:1018.4
```

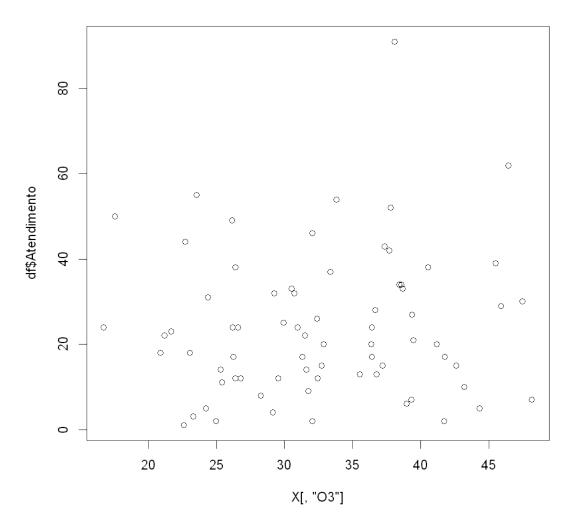
```
Max. :91.0
                    Max.
                          :29.14
                                   Max. :48.15 Max.
                                                        :1616.3
          NO2
                          S02
     Min.
                    Min.
                          : 5.959
            :11.19
      1st Qu.:19.33
                    1st Qu.: 9.361
     Median :24.45
                    Median :11.140
     Mean
           :24.42
                     Mean
                          :11.809
      3rd Qu.:28.70
                    3rd Qu.:13.695
     Max.
            :38.72
                     Max.
                           :20.420
[33]: df<- df %>% drop_na()
     Montando o X:
[34]: X <- as.matrix(df[ , setdiff(names(df), "Atendimentos")])
     Ajustando o modelo sem os splines:
[35]: fit =glarma(dfAtendimentos, X=X, phiLags = c(1), thetaLags = NULL, type = \Box
      →"Poi")
     summary(fit)
     Call: glarma(y = df$Atendimentos, X = X, type = "Poi", phiLags = c(1),
         thetaLags = NULL)
     Pearson Residuals:
        Min
                  10
                      Median
                                   ЗQ
                                           Max
     -5.1470 -1.5949 -0.2822
                               1.8875 11.4303
     GLARMA Coefficients:
          Estimate Std.Error z-ratio Pr(>|z|)
     phi_1 0.064316  0.007945  8.095 6.66e-16 ***
     Linear Model Coefficients:
              Estimate Std.Error z-ratio Pr(>|z|)
     tempmed 0.0626788 0.0080746 7.763 8.44e-15 ***
             03
     CO
             0.0018250 0.0001737 10.508 < 2e-16 ***
     NO2
            -0.0078196 0.0055289 -1.414
                                            0.157
     S02
            -0.0994414 0.0113291 -8.778 < 2e-16 ***
         Null deviance: 795.53 on 71 degrees of freedom
     Residual deviance: 612.48 on 66 degrees of freedom
     AIC: 943.4291
     Number of Fisher Scoring iterations: 30
     LRT and Wald Test:
     Alternative hypothesis: model is a GLARMA process
```

```
Null hypothesis: model is a GLM with the same regression structure
              Statistic p-value
                  75.71 < 2e-16 ***
     LR Test
     Wald Test
                  65.53 5.55e-16 ***
     Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
[36]: | fit =glarma(df$Atendimentos, X=X[,c('tempmed','03','C0','S02')] , phiLags =
      \rightarrowc(1,2), thetaLags = NULL, type = "Poi")
     summary(fit)
     Call: glarma(y = df$Atendimentos, X = X[, c("tempmed", "03", "CO",
         "SO2")], type = "Poi", phiLags = c(1, 2), thetaLags = NULL)
     Pearson Residuals:
        Min
                  1Q Median
                                   ЗQ
                                          Max
     -5.2529 -1.7941 -0.1555 1.4626 8.5222
     GLARMA Coefficients:
          Estimate Std.Error z-ratio Pr(>|z|)
     phi 1 0.082932 0.008198 10.116 < 2e-16 ***
     phi_2 0.046149 0.008895 5.188 2.13e-07 ***
     Linear Model Coefficients:
              Estimate Std.Error z-ratio Pr(>|z|)
     tempmed 0.0639349 0.0083392 7.667 1.75e-14 ***
     03
             0.0364563 0.0032981 11.054 < 2e-16 ***
             0.0013487 0.0001142 11.814 < 2e-16 ***
     CO
            S02
        Null deviance: 795.53 on 71 degrees of freedom
     Residual deviance: 578.17 on 66 degrees of freedom
     AIC: 917.5981
     Number of Fisher Scoring iterations: 30
     LRT and Wald Test:
     Alternative hypothesis: model is a GLARMA process
     Null hypothesis: model is a GLM with the same regression structure
              Statistic p-value
     LR Test
                  106.0 <2e-16 ***
                 112.5 <2e-16 ***
     Wald Test
     Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

A partir do AR(2) os coeficientes do phi não são significativos em todos os níveis de confiança.

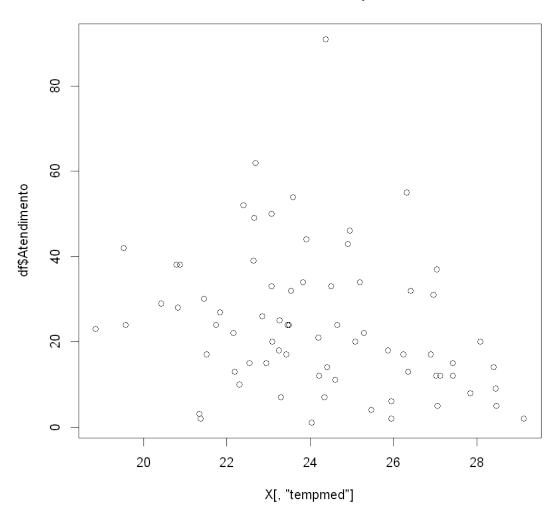
```
[37]: plot(X[, "03"], df$Atendimento, main = "Atendimento X 03")
```

Atendimento X O3



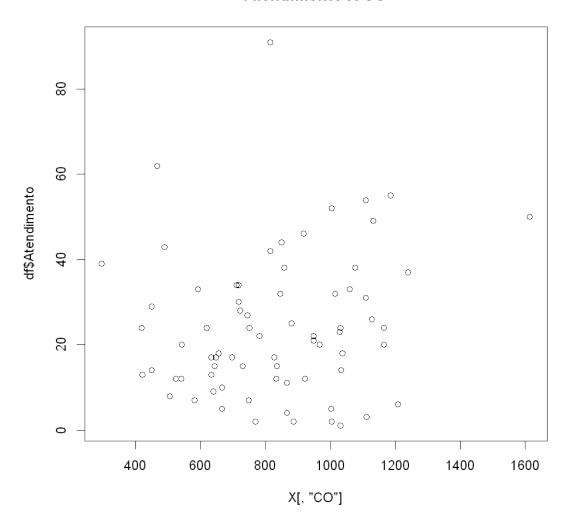
```
[38]: plot(X[, "tempmed"], df$Atendimento, main = "Atendimento X tempmed")
```

Atendimento X tempmed



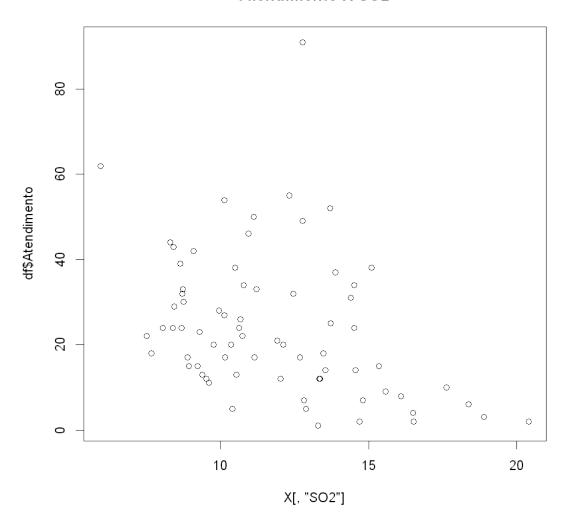
```
[39]: plot(X[, "CO"], df$Atendimento, main = "Atendimento X CO")
```

Atendimento X CO



```
[40]: plot(X[, "SO2"], df$Atendimento, main = "Atendimento X SO2")
```

Atendimento X SO2



O3 aparentemente tem relação não linear com o número de atendimentos Rodando nossa regressão com os splines:

\$AIC -7840.30738737551

\$logLik 3942.15369368775

\$beta0 2.76306610689278

\$beta_spline \$O3 1. 0.351727607570428 2. 0.748339770830553 3. 0.0514625420616618 4. 1.07794204788271 5. -0.521010265442694 6. 0.572686123411616 7. 0.636101002121954

\$tempmed 1. -0.10411575379499 2. -0.342784661481924 3. -0.0206866784576695 4. -0.127113092415123 5. -0.622252363904484 6. 0.0563834096290639 7. -1.64067137163042 8. -1.7303685099724

\$beta_other 1. 0.0284734533890816 2. 0.000740302083696139 3. -0.0866522606064693

\$fi 1. 0.0433467203032375 2. 0.00227101251663183 3. -0.0660772354515024

\$theta

\$k_list \$O3 7

\$temp_med 8

 $[\]:$