Análise de dados (EDA)

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1	Leitura de dados	
	brary(tidyverse) brary(hnp)	
da	<pre>dos <- read.csv("Arthritis.txt", sep="",</pre>	
da	dos\$id<- 1:nrow(dados)	
	dos<- tibble(dados) dos	
##	# A tibble: 51 x 9 Sex	

```
<fct> <int> <fct> <int> <int> <int> <int> <int> <int>
                1
##
  1 M
          48 A
                        1
                             1
                                 1
  2 M
          29 A
##
                   1
## 3 M
          59 P
                   1
                                          3
                       1
                             1
                                 1
                                      1
## 4 F
          56 P
                   1
                        1
                             1
                                 1
## 5 M
          33 P
                   1
                       1
                            1
                                 1
                                      1
## 6 M
          61 P
                   1
                       1
                            0
                                1
                       0
                            1 NA
## 7 M
          63 A
                   0
                                          7
                                     NA
                                    1
                       0
                   1
                            1 1
## 8 M
          57 P
## 9 M
          47 P
                   1 1 1
                                0
                                          9
                                     1
                   O O 1 NA
## 10 F
          42 A
                                     0 10
## # ... with 41 more rows
```

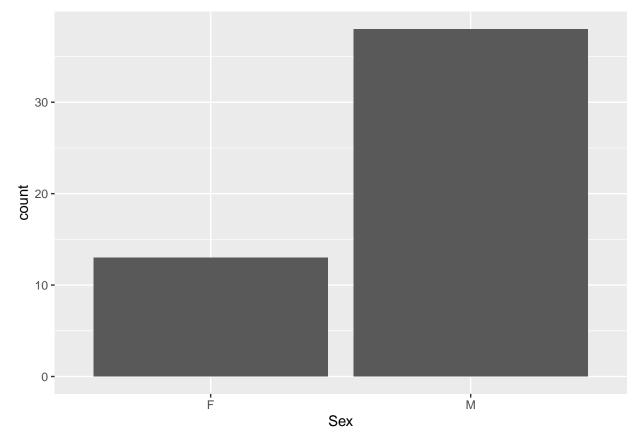
2 Alguns resumos dos dados

```
dados %>%
 group_by(Sex) %>%
summarise(n = n())
## # A tibble: 2 x 2
## Sex
   <fct> <int>
## 1 F
## 2 M
             38
dados %>%
 group_by(Sex) %>%
 summarise(media_Sex = mean(Age))
## # A tibble: 2 x 2
##
   Sex media_Sex
##
   <fct>
              <dbl>
## 1 F
              51.8
## 2 M
               50.2
dados %>%
 group_by(Group) %>%
summarise(n = n())
## # A tibble: 2 x 2
   Group
## <fct> <int>
## 1 A
             27
## 2 P
             24
dados %>%
 group_by(Group) %>%
 summarise( media_Age = mean(Age))
## # A tibble: 2 x 2
## Group media_Age
## <fct>
             <dbl>
## 1 A
              51.0
## 2 P
              50.2
```

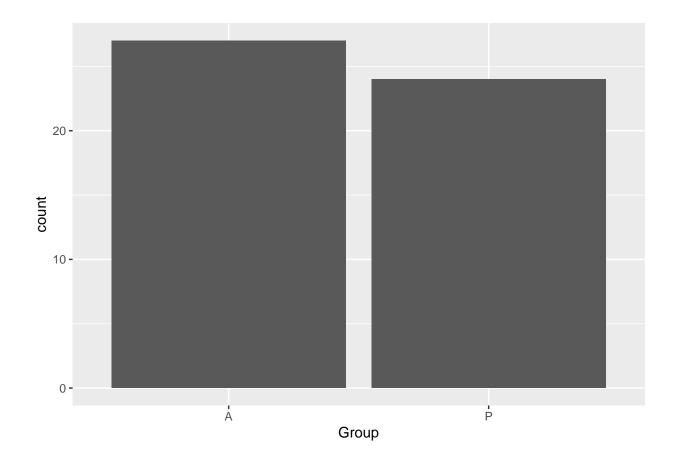
```
dados %>%
 group_by(Sex, Group) %>%
 summarise(n = n())
## # A tibble: 4 x 3
## # Groups: Sex [2]
##
    Sex
         Group
##
    <fct> <fct> <int>
## 1 F
          Α
## 2 F
          Ρ
                    6
## 3 M
                   20
          Α
## 4 M
          Р
                   18
```

3 Gráficos de interesse

```
ggplot(dados, aes(x = Sex)) +
  geom_bar()
```



```
ggplot(dados, aes(x = Group)) +
  geom_bar()
```

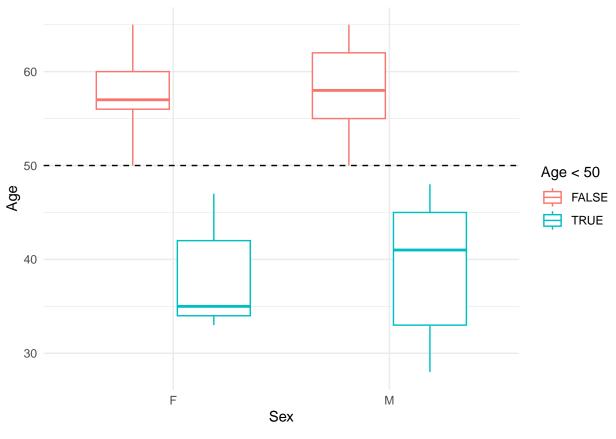


4 Transformando os dados

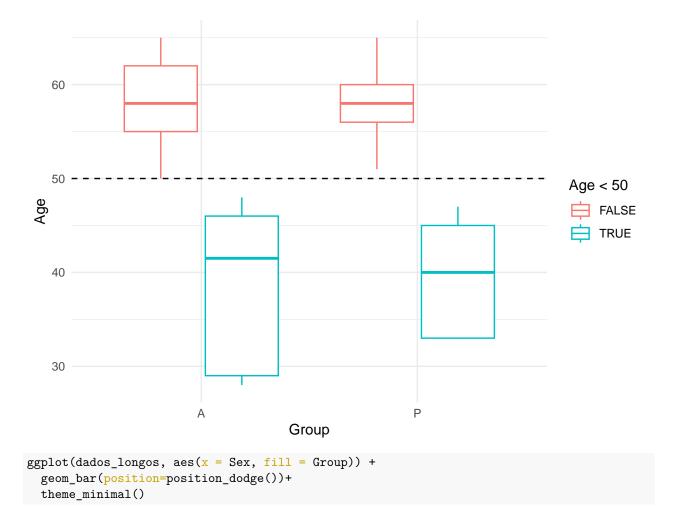
```
dados_longos<- dados %>%
  pivot_longer(
    cols = starts_with("Week"),
    names_to = "week",
    names_prefix = "Week",
    values_to = "Y",
    values_drop_na = TRUE
)
```

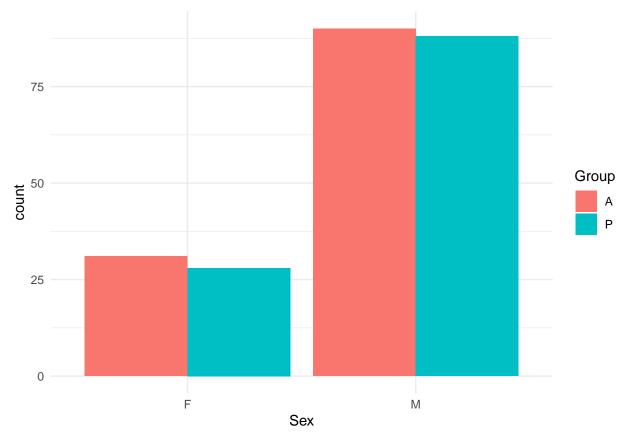
4.1 Gráficos antes de transformar dados

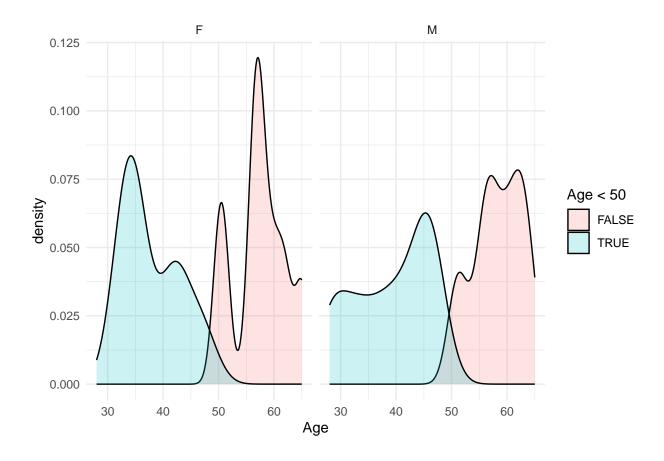
```
ggplot(dados_longos, aes(Sex, Age, col = Age < 50)) +
  geom_boxplot()+
  geom_hline(yintercept = 50, col = "black", linetype = 2)+
  theme_minimal()</pre>
```



```
ggplot(dados_longos, aes(Group, Age, col = Age < 50)) +
  geom_boxplot()+
  geom_hline(yintercept = 50, col = "black", linetype = 2)+
  theme_minimal()</pre>
```







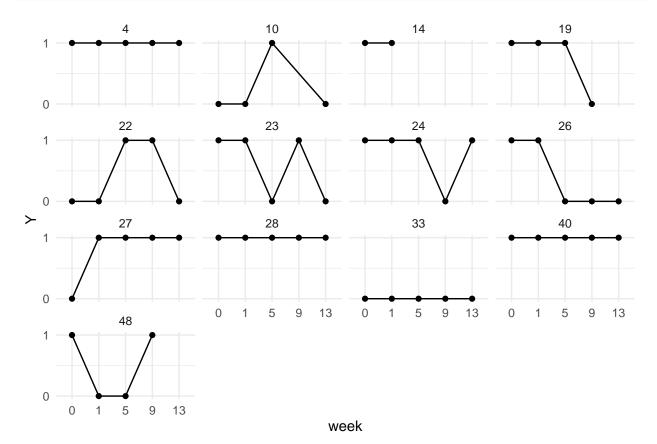
4.2 Transformando dados (seguindo o feito pelo Jalmar)

```
## # A tibble: 5 x 2
## vweek n
## 
## 1 0 51
## 2 1 51
## 3 5 48
## 4 9 45
## 5 13 42
```

5 Gráficos de perfis

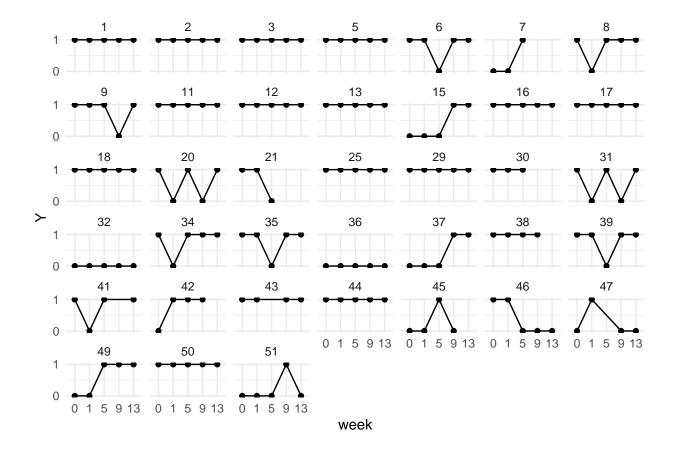
5.1 Sexo Female == 0

```
dados_longos %>% filter(Sex == "0") %>%
  ggplot(aes(week, Y, group = id)) +
  geom_point()+
  geom_line()+
  theme_minimal()+
  scale_y_continuous(breaks = c(0,1))+
  facet_wrap(~id)
```



5.2 Sexo Male == 1

```
dados_longos %>% filter(Sex == "1") %>%
  ggplot(aes(week, Y, group = id)) +
  geom_point()+
  geom_line()+
  theme_minimal()+
  scale_y_continuous(breaks = c(0,1))+
  facet_wrap(~id)
```

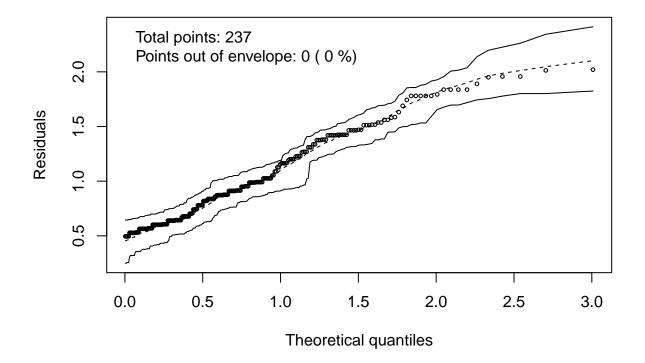


6 Ajuste de modelos

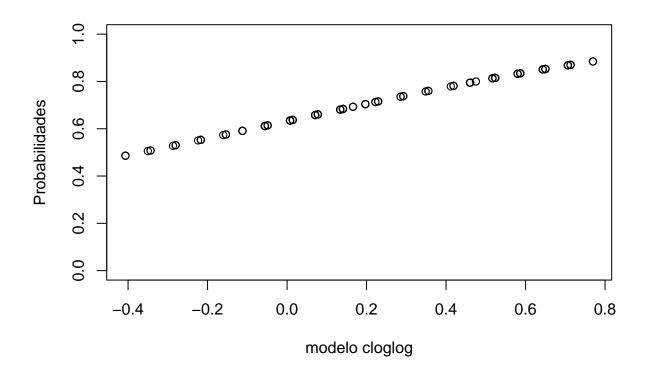
6.1 cloglog

```
modelo_cloglog<- glm(Y ~ Sex +</pre>
               Age +
               Group +
               as.numeric(week),
             family = binomial(link = "cloglog"),
             data= dados_longos)
modelo_cloglog$family
##
## Family: binomial
## Link function: cloglog
summary(modelo_cloglog)
##
## Call:
## glm(formula = Y ~ Sex + Age + Group + as.numeric(week), family = binomial(link = "cloglog"),
##
       data = dados_longos)
##
## Deviance Residuals:
##
       Min
                      Median
                                    3Q
                 1Q
                                            Max
## -2.0202 -1.2688
                      0.6394
                                0.8721
                                          1.2010
```

```
##
## Coefficients:
##
                    Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                                0.27391
                                        -1.715 0.086281 .
                    -0.46985
## Sex1
                     0.29448
                                0.19859
                                          1.483 0.138115
## Age1
                                0.17402
                                          0.324 0.745898
                     0.05639
## Group1
                     0.57257
                                0.16874
                                          3.393 0.000691 ***
## as.numeric(week)
                                0.05979
                                          1.057 0.290470
                     0.06321
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
   (Dispersion parameter for binomial family taken to be 1)
##
       Null deviance: 282.26 on 236 degrees of freedom
##
## Residual deviance: 268.18 on 232 degrees of freedom
## AIC: 278.18
##
## Number of Fisher Scoring iterations: 5
hnp(modelo_cloglog, print.on = TRUE)
```



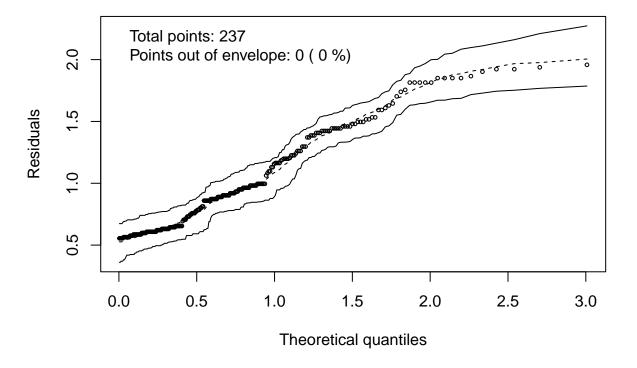
```
plot(predict.glm(modelo_cloglog, type="response")~predict.glm(modelo_cloglog, type="link"),
    ylab = "Probabilidades",
    xlab = "modelo cloglog",
    ylim=c(0,1))
```



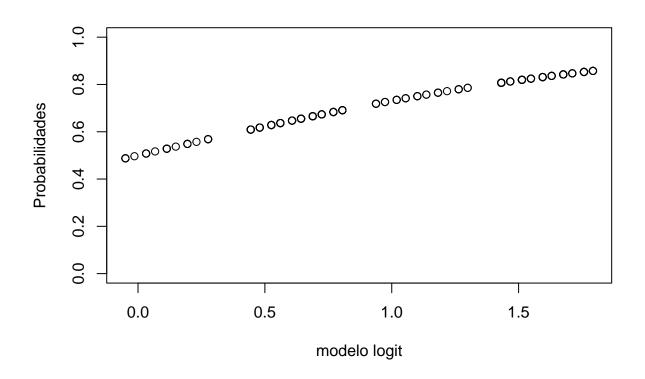
6.2 logit

```
modelo_logit<- glm(Y ~ Sex +</pre>
               Age +
               Group +
               as.numeric(week),
                      family = binomial(link = "logit"),
                      data= dados_longos)
modelo_logit$family
##
## Family: binomial
## Link function: logit
summary(modelo_logit)
##
## Call:
## glm(formula = Y ~ Sex + Age + Group + as.numeric(week), family = binomial(link = "logit"),
##
       data = dados_longos)
##
## Deviance Residuals:
##
       Min
                 1Q
                      Median
                                    3Q
                                             Max
                      0.6309
## -1.9579 -1.2612
                                0.8723
                                          1.1986
##
## Coefficients:
```

```
##
                   Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                   -0.13114
                               0.44776
                                        -0.293 0.76962
## Sex1
                    0.49379
                                0.33640
                                          1.468
                                                0.14214
                    0.03550
                               0.31166
                                                0.90930
## Age1
                                         0.114
## Group1
                     0.98760
                               0.30382
                                         3.251
                                                0.00115 **
## as.numeric(week)
                    0.08148
                               0.10635
                                         0.766
                                                0.44360
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
   (Dispersion parameter for binomial family taken to be 1)
##
##
      Null deviance: 282.26 on 236 degrees of freedom
##
## Residual deviance: 268.79 on 232 degrees of freedom
## AIC: 278.79
##
## Number of Fisher Scoring iterations: 4
hnp(modelo_logit, print.on = TRUE)
```



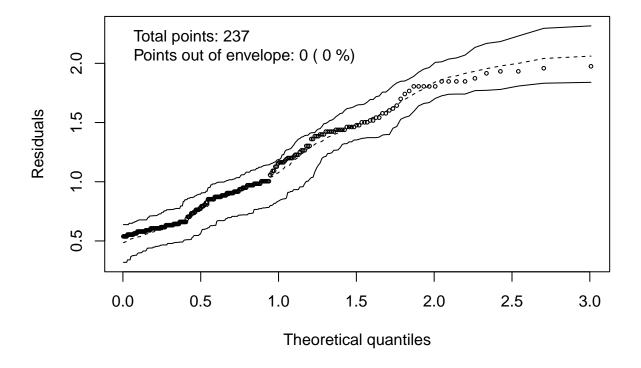
```
plot(predict.glm(modelo_logit, type="response")~predict.glm(modelo_logit, type="link"),
    ylab = "Probabilidades",
    xlab = "modelo logit",
    ylim=c(0,1))
```



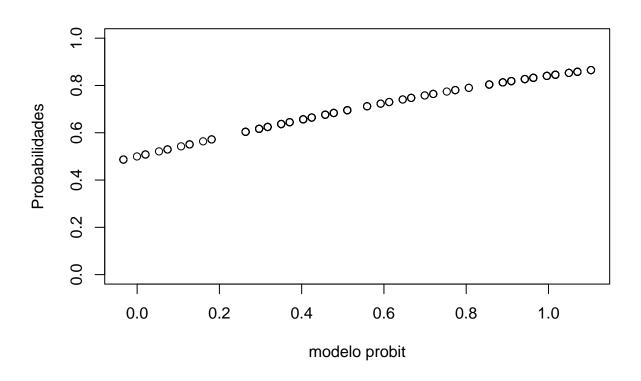
6.3 probit

```
modelo_probit<- glm(Y ~ Sex +</pre>
               Age +
               Group +
               as.numeric(week),
                   family = binomial(link = "probit"),
                    data= dados_longos)
modelo_probit$family
##
## Family: binomial
## Link function: probit
summary(modelo_probit)
##
## Call:
## glm(formula = Y ~ Sex + Age + Group + as.numeric(week), family = binomial(link = "probit"),
##
       data = dados_longos)
##
## Deviance Residuals:
       Min
##
                 1Q
                      Median
                                    3Q
                                            Max
## -1.9750 -1.2650
                      0.6329
                                         1.2001
                                0.8721
##
## Coefficients:
```

```
Estimate Std. Error z value Pr(>|z|)
##
                                        -0.321 0.748510
## (Intercept)
                    -0.08693
                                0.27115
## Sex1
                                          1.468 0.142233
                     0.29695
                                0.20235
                     0.03294
                                0.18530
                                          0.178 0.858905
## Age1
## Group1
                     0.59240
                                0.17866
                                          3.316 0.000914 ***
## as.numeric(week)
                    0.05359
                                0.06329
                                          0.847 0.397121
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
   (Dispersion parameter for binomial family taken to be 1)
##
##
      Null deviance: 282.26 on 236 degrees of freedom
##
## Residual deviance: 268.63 on 232 degrees of freedom
## AIC: 278.63
##
## Number of Fisher Scoring iterations: 4
hnp(modelo_probit, print.on = TRUE)
```



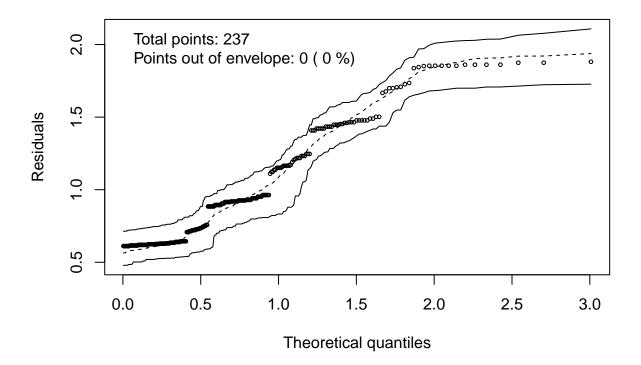
```
plot(predict.glm(modelo_probit, type="response")~predict.glm(modelo_probit, type="link"),
    ylab = "Probabilidades",
    xlab = "modelo probit",
    ylim=c(0,1))
```



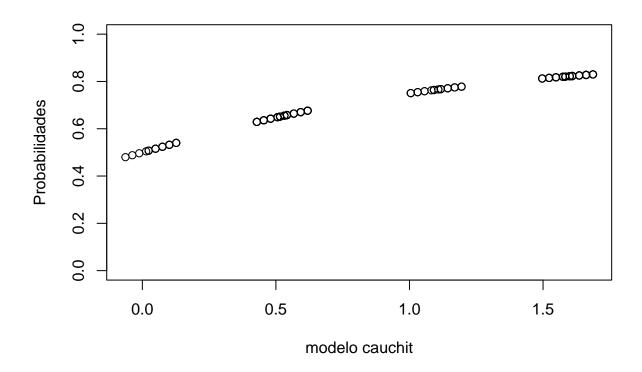
6.4 cauchit

```
modelo_cauchit<- glm(Y ~ Sex +</pre>
               Age +
               Group +
               as.numeric(week),
                    family = binomial(link = "cauchit"),
                    data= dados_longos)
modelo_cauchit$family
##
## Family: binomial
## Link function: cauchit
summary(modelo_cauchit)
##
## Call:
## glm(formula = Y ~ Sex + Age + Group + as.numeric(week), family = binomial(link = "cauchit"),
##
       data = dados_longos)
##
## Deviance Residuals:
##
       Min
                 1Q
                      Median
                                    3Q
                                            Max
                      0.6286
## -1.8815 -1.2323
                                0.8943
                                          1.2118
##
## Coefficients:
```

```
##
                     Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                    -0.002309
                                0.420176
                                         -0.005
                                                   0.9956
## Sex1
                                           1.471
                     0.492029
                                0.334590
                                                   0.1414
                    -0.086834
                                0.321406
                                         -0.270
                                                   0.7870
## Age1
## Group1
                     1.068118
                                0.383931
                                           2.782
                                                   0.0054 **
## as.numeric(week)
                    0.025811
                                0.107747
                                           0.240
                                                   0.8107
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
   (Dispersion parameter for binomial family taken to be 1)
##
##
      Null deviance: 282.26 on 236 degrees of freedom
##
## Residual deviance: 269.49
                             on 232 degrees of freedom
## AIC: 279.49
##
## Number of Fisher Scoring iterations: 6
hnp(modelo_cauchit, print.on = TRUE)
```



```
plot(predict.glm(modelo_cauchit, type="response")~predict.glm(modelo_cauchit, type="link"),
    ylab = "Probabilidades",
    xlab = "modelo cauchit",
    ylim=c(0,1))
```



7 GEE

7.1 independence

```
library(gee)
modelo_gee_1 <- gee(Y ~ Sex + Age + Group + as.numeric(week),</pre>
               data = dados_longos,
               id = id,
               family = binomial(link = "cloglog"),
               corstr = "independence")
##
        (Intercept)
                                 Sex1
                                                   Age1
                                                                  Group1
##
        -0.46985374
                           0.29447851
                                            0.05639087
                                                              0.57257244
## as.numeric(week)
         0.06320692
summary(modelo_gee_1)
##
    GEE: GENERALIZED LINEAR MODELS FOR DEPENDENT DATA
##
##
    gee S-function, version 4.13 modified 98/01/27 (1998)
## Model:
   Link:
                                Cloglog
    Variance to Mean Relation: Binomial
##
  Correlation Structure:
                                Independent
```

```
##
## Call:
## gee(formula = Y ~ Sex + Age + Group + as.numeric(week), id = id,
      data = dados_longos, family = binomial(link = "cloglog"),
      corstr = "independence")
##
## Summary of Residuals:
##
         Min
                           Median
                                          3Q
                                                   Max
## -0.8700482 -0.5528727 0.1848709 0.3163075 0.5138260
##
##
## Coefficients:
                     Estimate Naive S.E.
                                          Naive z Robust S.E.
## (Intercept)
                  ## Sex1
                    ## Age1
                    0.05640169 \quad 0.1753833 \quad 0.321591 \quad 0.24953232 \quad 0.2260296
## Group1
                    0.57257352 0.1700686 3.366721 0.24311673 2.3551383
## as.numeric(week) 0.06321248 0.0602626 1.048950 0.05275103 1.1983173
## Estimated Scale Parameter: 1.015735
## Number of Iterations: 1
## Working Correlation
       [,1] [,2] [,3] [,4] [,5]
## [1,]
              0
                        0
          1
                    0
## [2,]
          0
               1
                    0
## [3,]
          0
               0
                        0
                             0
                    1
## [4,]
          0
               0
                    0
                        1
                             0
## [5,]
          0
     exchangeable
modelo_gee_2 <- gee(Y ~ Sex + Age + Group + as.numeric(week),</pre>
              data = dados longos,
              id = id,
              family = binomial(link = "cloglog"),
              corstr = "exchangeable")
##
       (Intercept)
                                                             Group1
                              Sex1
                                              Age1
                        0.29447851
                                                         0.57257244
##
       -0.46985374
                                         0.05639087
## as.numeric(week)
        0.06320692
summary(modelo_gee_2)
##
   GEE: GENERALIZED LINEAR MODELS FOR DEPENDENT DATA
##
   gee S-function, version 4.13 modified 98/01/27 (1998)
##
## Model:
## Link:
                             Cloglog
   Variance to Mean Relation: Binomial
## Correlation Structure:
                             Exchangeable
##
## Call:
```

```
## gee(formula = Y ~ Sex + Age + Group + as.numeric(week), id = id,
##
       data = dados_longos, family = binomial(link = "cloglog"),
##
       corstr = "exchangeable")
##
## Summary of Residuals:
##
         Min
                                            3Q
                     1Q
                            Median
                                                     Max
## -0.8599897 -0.5530691 0.1870639 0.3276707 0.5032505
##
##
## Coefficients:
                      Estimate Naive S.E.
                                             Naive z Robust S.E.
                                                                   Robust z
## (Intercept)
                   -0.42904864 0.33365054 -1.2859222 0.32034671 -1.3393259
## Sex1
                    0.27285705 0.28277327 0.9649323 0.27849472 0.9797567
## Age1
                    0.08134966 0.24774357 0.3283623 0.24724658 0.3290224
                    0.53829103 0.24201844 2.2241736 0.24310512 2.2142316
## Group1
## as.numeric(week) 0.05314300 0.05066053 1.0490021 0.05083011 1.0455023
## Estimated Scale Parameter: 0.9965314
## Number of Iterations: 4
## Working Correlation
                       [,2]
                                 [,3]
             [,1]
## [1,] 1.0000000 0.2907233 0.2907233 0.2907233 0.2907233
## [2,] 0.2907233 1.0000000 0.2907233 0.2907233 0.2907233
## [3,] 0.2907233 0.2907233 1.0000000 0.2907233 0.2907233
## [4,] 0.2907233 0.2907233 1.0000000 0.2907233
## [5,] 0.2907233 0.2907233 0.2907233 1.0000000
7.3 unstructured
modelo_gee_3 <- gee(Y ~ Sex + Age + Group + as.numeric(week),</pre>
              data = dados_longos,
               id = id,
              family = binomial(link = "cloglog"),
               corstr = "unstructured")
##
        (Intercept)
                                Sex1
                                                                Group1
                                                 Age1
        -0.46985374
                         0.29447851
##
                                          0.05639087
                                                            0.57257244
## as.numeric(week)
        0.06320692
summary(modelo_gee_3)
##
   GEE: GENERALIZED LINEAR MODELS FOR DEPENDENT DATA
##
   gee S-function, version 4.13 modified 98/01/27 (1998)
##
##
## Model:
## Link:
                               Cloglog
## Variance to Mean Relation: Binomial
## Correlation Structure:
                              Unstructured
##
## Call:
## gee(formula = Y ~ Sex + Age + Group + as.numeric(week), id = id,
      data = dados_longos, family = binomial(link = "cloglog"),
```

```
##
    corstr = "unstructured")
##
## Summary of Residuals:
  Min 1Q Median 3Q
                                              Max
## -0.8703654 -0.5230487 0.1650142 0.3603214 0.5205362
##
##
## Coefficients:
##
                    Estimate Naive S.E. Naive z Robust S.E. Robust z
## (Intercept)
               ## Sex1
                 0.279293127  0.2731156  1.02261888  0.27175387  1.02774297
                  0.007621797  0.2397362  0.03179243  0.24396340  0.03124156
## Age1
## Group1
                  0.693890430 0.2333050 2.97417687 0.24223920 2.86448444
## as.numeric(week) 0.041897668 0.0552891 0.75779257 0.04990972 0.83946913
## Estimated Scale Parameter: 1.028027
## Number of Iterations: 7
##
## Working Correlation
         [,1] [,2] [,3] [,4] [,5]
## [1,] 1.0000000 0.64699683 0.19879899 0.3105900 0.2345909
## [2,] 0.6469968 1.00000000 0.02545539 0.2919917 0.1593287
## [3,] 0.1987990 0.02545539 1.00000000 0.1337962 0.2364074
## [4,] 0.3105900 0.29199173 0.13379622 1.0000000 0.1851979
## [5,] 0.2345909 0.15932868 0.23640737 0.1851979 1.0000000
```