

# TFM\_UOC

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```
# Librerías
library(knitr)
library(kableExtra)
library(fastDummies)
library(xgboost)
library(tidyverse)
library(naniar)
library(corrplot)
library(caret)
library(DMwR2) #imputación knn
library(reshape2)
library(FactoMineR)
library(factoextra)
library(ggrepel)
library(MASS) # LDA
library(car)
library(multcomp)
library(pROC)
library(fastshap)

#Otras posibles
#library(readxl)
#library(psych)
#library(DataExplorer)
#library(GGally)
#library(ROSE)
#library(ggfortify)
```

## OBTENCIÓN DEL DATASET FINAL

Tras reunión de coordinación, seleccionaremos demográficas y clínicas relevantes.

La columna del espectro clínico de la primera infección tiene NAs, con una proporción de 10 asintomáticos, 35 hospitalizados, 185 en mild-moderated y 7 en UCI. La imputación que generaré es con el valor más común.

```
#Seleccionamos el espectro clínico de la primera infección
column_spec <- bddAECM$spec1
selected_spec <- data.frame(spec1 = column_spec)

#Imputación de valores faltantes en la columna 'spec1'
most_frequent <- names(sort(table(selected_spec), decreasing = TRUE))[1]

#Imputamos los valores faltantes con el más frecuente
```

```
selected_spec[is.na(selected_spec)] <- most_frequent
```

```
#Generamos dummies solo para 'spec1'  
selected_spec1_dummy <- dummy_cols(selected_spec,  
                                   remove_first_dummy = TRUE,  
                                   remove_selected_columns = TRUE)
```

Ahora seleccionamos las demográficas de interés.

```
#Variables sociodemográficas indicadas en la reunión de coordinación.  
selected_sociodemographic <- bbddAECM %>%  
  dplyr::select(sex, ag, el, ptg19)  
  
#Creamos dummies para las variables demográficas seleccionadas.  
selected_sociodemographic_dummies <- dummy_cols(  
  selected_sociodemographic,  
  remove_first_dummy = TRUE,  
  remove_selected_columns = TRUE  
)
```

Ahora seleccionamos las variables neuropsicológicas de interés.

```
#Realizamos selección de tests neuropsicológicos de interés.  
selected_neuropsychological <- bbddAECM %>%  
  dplyr::select(tn46, tn52, tn36,  
               tn38, tn40, tn42,  
               tn22, tn44, tn14,  
               tn24, tn12, tn6,  
               tn30, tn34, tn8,  
               tn48, tn50)
```

Y ahora que ya tenemos todas las variables de interés las combinamos en un único dataset.

```
#Combinamos las variables seleccionadas en un solo dataset.  
df_def <- bind_cols(selected_sociodemographic_dummies, selected_spec1_dummy, selected_neuropsychological)
```

Resta crear la variable Target sobre la que generaremos el estudio.

```
# Seleccionamos NoCOVID y No LongCOVID (que será un grupo) vs LongCovid Cog  
column_cluster <- bbddAECM$cluster  
selected_cluster <- data.frame(cluster = column_cluster)  
  
#Creamos la nueva variable Target  
selected_cluster$cluster <- ifelse(  
  selected_cluster$cluster %in% c("No COVID", "No LongCOVID"), "No_LC",  
  ifelse(selected_cluster$cluster == "LongCOVID Cog", "LC_Cog", NA) # Excluir "LongCOVID NoCog"  
)  
  
#Convertimos Target a factor  
selected_cluster$cluster <- factor(selected_cluster$cluster, levels = c("No_LC", "LC_Cog"))  
  
# Convertir a numérico restando 1 para que los niveles sean 0 y 1  
selected_cluster$cluster_binary <- as.numeric(selected_cluster$cluster) - 1
```

Finalmente unimos todas las variables definitivamente seleccionadas.

```
#Unimos la columna de los targets al dataframe final  
df_def <- cbind(df_def, Target = selected_cluster$cluster_binary)
```

```

#-----
##COMPROBAMOS QUE LOS INDICES COINCIDEN EN LA BBDD ORIGINAL Y EN LA RE-CREADA
#-----

#Obtenemos los índices de las observaciones donde cluster es "LongCOVID NoCog"
indices <- which(bbddAECM$cluster == "LongCOVID NoCog")

#Verificamos
print(indices)

[1] 1 22 42 55 58 60 61 62 63 64 66 68 184 192 195 199 205 209 211
[20] 230 234 237 238 241 243 250 263

#Obtenemos los índices de las observaciones donde Target es NA en el nuevo dataset
indices_na <- which(is.na(df_def$Target))

#Verificamos
print(indices_na)

[1] 1 22 42 55 58 60 61 62 63 64 66 68 184 192 195 199 205 209 211
[20] 230 234 237 238 241 243 250 263

# ELIMINAMOS TODAS LAS OBSERVACIONES NAS EN TARGET
df_def <- df_def[!is.na(df_def$Target), ]

#Dado que los índices coinciden, df_def será el dataset que utilizaremos para realizar el TFM

#Aquí remodificamos los nombres para evitar problemas
df_def <- df_def %>% rename_with(make.names)

#-----
# Análisis general del dataset de trabajo final generado
#-----

# Ver tipos de variables y estructura
str(df_def)

'data.frame': 241 obs. of 29 variables:
 $ ag          : num 26.9 60.5 33.1 44.6 61.9 43 55.6 53.6 46.3 51.2 ...
 $ ptg19       : num 38.2 26.3 18.2 24.1 32.8 ...
 $ sex_Woman   : int 1 0 1 1 0 1 1 1 1 1 ...
 $ el_Elementary : int 0 0 0 0 0 0 0 0 0 0 ...
 $ el_High.School : int 1 0 0 0 0 0 0 0 0 0 ...
 $ el_Secondary : int 0 0 0 0 0 0 0 0 0 0 ...
 $ el_Specialist.Master : int 0 1 0 0 1 1 1 0 1 0 ...
 $ el_University.Deg. : int 0 0 0 1 0 0 0 1 0 1 ...
 $ spec1_Hospitalization: int 0 1 0 0 1 0 0 0 0 1 ...
 $ spec1_Mild.Moderated : int 1 0 1 1 0 1 1 1 1 0 ...
 $ spec1_UCI      : int 0 0 0 0 0 0 0 0 0 0 ...
 $ tn46           : num 0.3 1 0.7 0 0 0.3 -0.3 -0.3 -0.7 -0.3 ...
 $ tn52           : num 1 0 2 -0.6 0 ...
 $ tn36           : num -1 0 1 0.3 0.3 -0.7 -0.3 -0.3 -1.7 0 ...
 $ tn38           : num -0.3 0.3 0.3 0 -0.3 -0.3 -0.7 0 -2.3 -0.3 ...
 $ tn40           : num 0 0 1 1 0.3 0.3 -0.3 -0.3 -0.3 -0.3 ...
 $ tn42           : num -1.3 0 0.3 1.3 -0.3 -0.3 -1 -1.7 -0.7 0.3 ...

```

```

$ tn22      : num  0.4 0.4 -0.5 -0.9 -0.2 1 0.4 0.3 0.3 0.1 ...
$ tn44      : num  -0.7 -1.3 -1 1 -0.3 0 -0.7 1 -2 -0.7 ...
$ tn14      : num  -1.7 1 0.3 -0.3 1.3 0.7 0.3 0.7 -0.3 0.7 ...
$ tn24      : num  -3 0 -1 -0.7 -0.3 -0.7 0.4 -0.7 -2.7 -0.7 ...
$ tn12      : num  -2 0 1.3 -0.3 1.3 0 1 0 0.7 0 ...
$ tn6       : num  -2.06 0.86 1.7 -0.82 -0.57 1.38 -1.18 -0.2 -0.01 -0.32 ...
$ tn30      : num  -2.3 0.7 0.32 -0.43 0.4 1.71 -1.22 0.03 -0.07 0.03 ...
$ tn34      : num  -1 0.3 -1 -1 0.3 0 -0.3 -1 1 -2 ...
$ tn8       : num  2 -0.3 0.3 -0.7 -0.3 2.7 0 -0.3 -0.3 0.3 ...
$ tn48      : num  1 3.1 3.1 1.3 3.1 1.3 1.3 3.1 1 0.8 ...
$ tn50      : num  -1.7 1.3 -0.3 -0.3 -0.7 0.7 1.3 0.7 0.3 0.7 ...
$ Target    : num  1 1 0 1 1 1 1 1 1 1 ...

```

```
# Resumen estadístico
```

```
summary(df_def)
```

ag	ptg19	sex_Woman	el_Elementary
Min. :25.40	Min. :16.05	Min. :0.0000	Min. :0.00000
1st Qu.:42.40	1st Qu.:23.01	1st Qu.:1.0000	1st Qu.:0.00000
Median :48.80	Median :26.19	Median :1.0000	Median :0.00000
Mean :48.72	Mean :27.69	Mean :0.8008	Mean :0.04149
3rd Qu.:55.40	3rd Qu.:30.60	3rd Qu.:1.0000	3rd Qu.:0.00000
Max. :70.80	Max. :87.31	Max. :1.0000	Max. :1.00000
	NA's :6		
el_High.School	el_Secondary	el_Specialist.Master	el_University.Deg.
Min. :0.0000	Min. :0.00000	Min. :0.0000	Min. :0.0000
1st Qu.:0.0000	1st Qu.:0.00000	1st Qu.:0.0000	1st Qu.:0.0000
Median :0.0000	Median :0.00000	Median :0.0000	Median :0.0000
Mean :0.3278	Mean :0.03734	Mean :0.1494	Mean :0.3776
3rd Qu.:1.0000	3rd Qu.:0.00000	3rd Qu.:0.0000	3rd Qu.:1.0000
Max. :1.0000	Max. :1.00000	Max. :1.0000	Max. :1.0000
spec1_Hospitalization	spec1_Mild.Moderated	spec1_UCI	tn46
Min. :0.0000	Min. :0.0000	Min. :0.0000	Min. : -2.7000
1st Qu.:0.0000	1st Qu.:1.0000	1st Qu.:0.0000	1st Qu.: -0.7000
Median :0.0000	Median :1.0000	Median :0.0000	Median : 0.0000
Mean :0.1411	Mean :0.7967	Mean :0.0249	Mean : -0.1021
3rd Qu.:0.0000	3rd Qu.:1.0000	3rd Qu.:0.0000	3rd Qu.: 0.3000
Max. :1.0000	Max. :1.0000	Max. :1.0000	Max. : 2.7000
			NA's :3
tn52	tn36	tn38	tn40
Min. : -2.30000	Min. : -3.0000	Min. : -2.3000	Min. : -3.0000
1st Qu.: -0.40000	1st Qu.: -0.7000	1st Qu.: -0.7000	1st Qu.: -0.9250
Median : 0.00000	Median : -0.3000	Median : -0.3000	Median : -0.3000
Mean : 0.02143	Mean : -0.2453	Mean : -0.1378	Mean : -0.2483
3rd Qu.: 0.40000	3rd Qu.: 0.3000	3rd Qu.: 0.3000	3rd Qu.: 0.3000
Max. : 2.00000	Max. : 2.3000	Max. : 2.7000	Max. : 2.7000
NA's :3	NA's :3	NA's :3	NA's :3
tn42	tn22	tn44	tn14
Min. : -3.1000	Min. : -2.3000	Min. : -3.000	Min. : -2.3000
1st Qu.: -1.3000	1st Qu.: -0.1000	1st Qu.: -0.700	1st Qu.: -0.3000
Median : -0.7000	Median : 0.5000	Median : -0.300	Median : 0.3000
Mean : -0.7286	Mean : 0.3982	Mean : -0.363	Mean : 0.4055
3rd Qu.: 0.0000	3rd Qu.: 0.9000	3rd Qu.: 0.225	3rd Qu.: 1.0000
Max. : 2.3000	Max. : 3.0000	Max. : 2.300	Max. : 2.7000

NA's :3 tn24	NA's :4 tn12	NA's :3 tn6	NA's :3 tn30
Min. : -3.0000	Min. : -2.0000	Min. : -5.9400	Min. : -2.53000
1st Qu.: -1.0000	1st Qu.: -0.3000	1st Qu.: -1.0500	1st Qu.: -0.78000
Median : -0.3000	Median : 0.1500	Median : -0.1900	Median : 0.03000
Mean : -0.4887	Mean : 0.2479	Mean : -0.1951	Mean : 0.03378
3rd Qu.: 0.0000	3rd Qu.: 1.0000	3rd Qu.: 0.8575	3rd Qu.: 0.97000
Max. : 2.3000	Max. : 2.3000	Max. : 2.2700	Max. : 1.73000
NA's :3 tn34	NA's :3 tn8	NA's :3 tn48	NA's :3 tn50
Min. : -2.7000	Min. : -2.0000	Min. : 0.300	Min. : -2.0000
1st Qu.: -0.7000	1st Qu.: -0.7000	1st Qu.: 1.000	1st Qu.: 0.0000
Median : -0.3000	Median : -0.3000	Median : 1.300	Median : 0.3000
Mean : -0.3571	Mean : -0.1202	Mean : 1.615	Mean : 0.2294
3rd Qu.: 0.0000	3rd Qu.: 0.3000	3rd Qu.: 3.000	3rd Qu.: 0.7000
Max. : 2.7000	Max. : 2.7000	Max. : 3.100	Max. : 3.0000
NA's :3 Target			
Min. : 0.000			
1st Qu.: 0.000			
Median : 1.000			
Mean : 0.722			
3rd Qu.: 1.000			
Max. : 1.000			

```
# if (!requireNamespace("webshot", quietly = TRUE)) {
#   install.packages("webshot")
#   webshot::install_phantomjs()
# }
# library(webshot)
#
# # Crear un data frame a partir del summary
# summary_table <- as.data.frame(summary(df_def))
#
# save_kable(
#   kable(summary_table, format = "html", caption = "Resumen de df_def") %>%
#     kable_styling(full_width = FALSE, bootstrap_options = c("striped", "hover", "condensed")),
#   file = "tabla_resumen.html"
# )
#
# # Convertir el archivo HTML a JPG
# webshot("tabla_resumen.html", file = "tabla_resumen.jpg", vwidth = 800, vheight = 600)
#
# Ver las dimensiones y primeras filas
cat("Dimensiones del dataset:", dim(df_def), "\n")
```

Dimensiones del dataset: 241 29

Vamos a verificar si hay algún valor faltante

```
#-----
# Valores faltantes
#-----

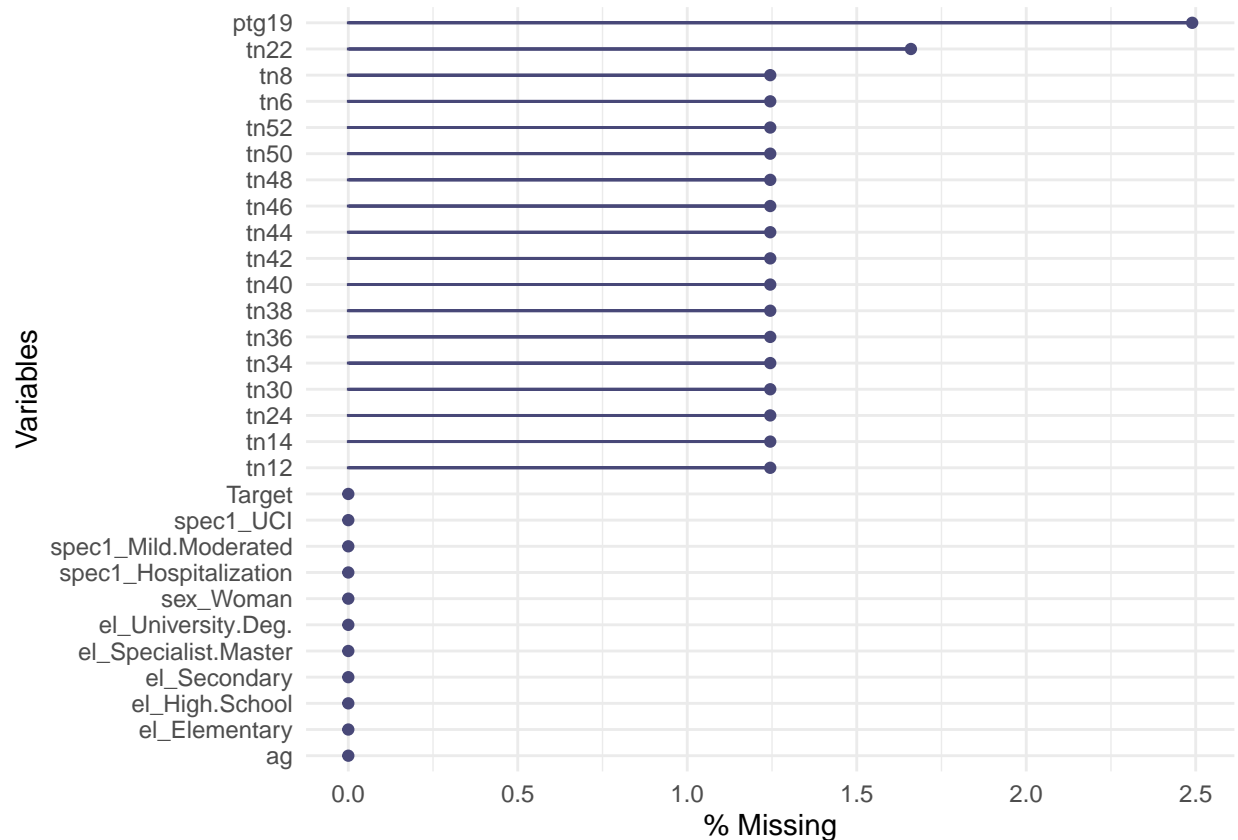
pdf("outputs/images/01_NAs.pdf", width = 16, height = 10)
```

```
# Resumen de valores faltantes
gg_miss_var(df_def, show_pct = TRUE)

dev.off()
```

pdf  
2

```
gg_miss_var(df_def, show_pct = TRUE)
```



Como no tenemos una idea global del tipo de valores faltantes, vamos a hacer una imputación con kNN para estos.

```
#Excluimos columnas específicas
excluded_cols <- c("Target")

#Separamos las columnas excluidas
excluded_data <- df_def %>%
  dplyr::select(all_of(excluded_cols))

#Convertimos todas las columnas numéricas a tipo double
data_for_imputation_selected <- df_def %>% dplyr::select(-all_of(excluded_cols))
data_for_imputation_selected <- data_for_imputation_selected %>%
  mutate(across(where(is.numeric), as.numeric))

#Aplicamos la imputación
selected_data_imputed <- knnImputation(data_for_imputation_selected, k = 10)
```

```
#Combinamos las columnas imputadas con las excluidas
df_def_clean <- bind_cols(selected_data_imputed, excluded_data)

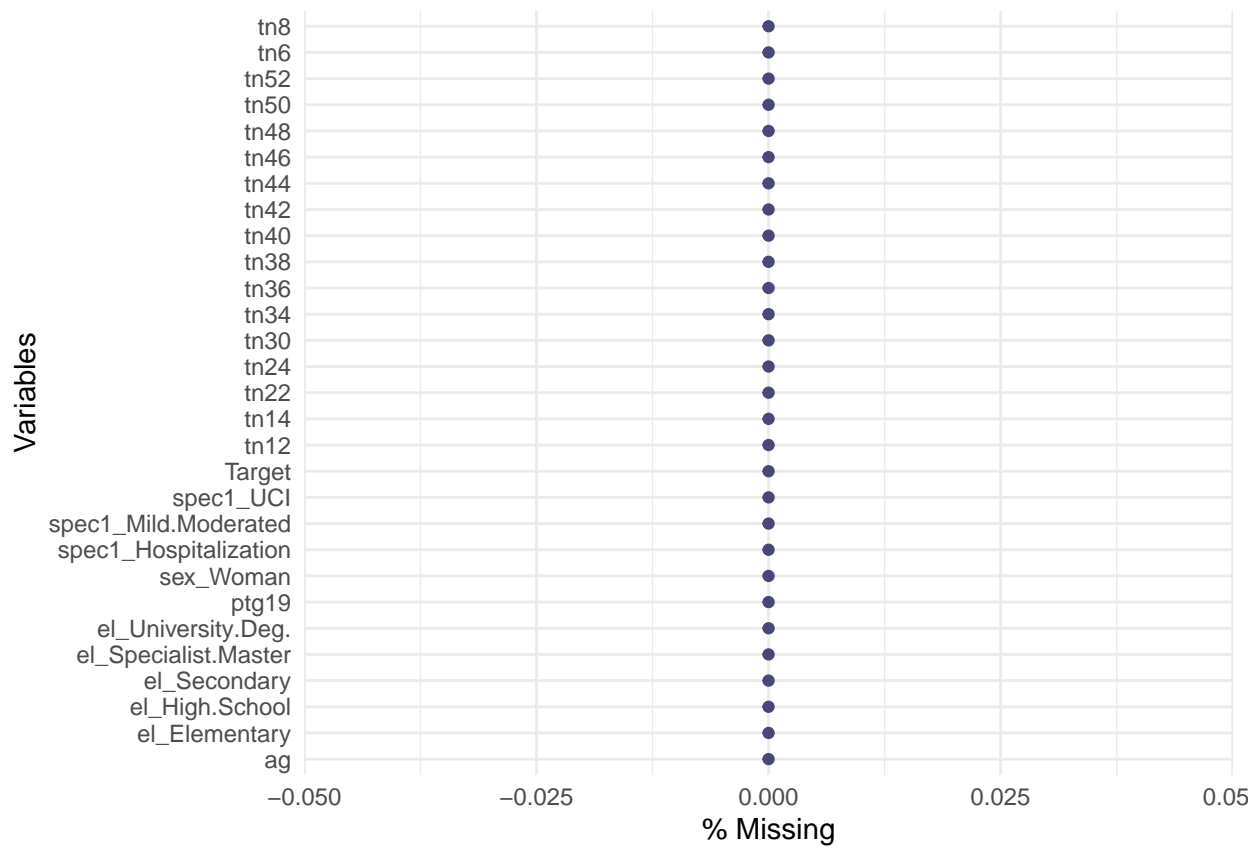
# Gráfico tras imputación
pdf("outputs/images/02_NAs_tras_imputacion.pdf", width = 16, height = 10)

gg_miss_var(df_def_clean, show_pct = TRUE)

dev.off()
```

```
pdf
  2
```

```
gg_miss_var(df_def_clean, show_pct = TRUE)
```



Vamos a dejar el código del balanceo del dataset y guardamos el dataset balanceado por si fuera de interés para utilizarlo posteriormente.

```
# -----
# Balancear el dataset
# -----

table(df_def_clean$Target) #Hay 64 vs 174
```

```
0    1
67 174
```

```

#Si aplicmos ROSE: OJO: DATOS SINTÉTICOS
df_def_clean_balanced_ROSE <- ROSE(Target ~ ., data = df_combined_clean, seed = 123, method = "under")

# Oversampling manual
minority_class <- df_def_clean %>% filter(Target == "0")
oversampled <- minority_class %>% sample_n(size = nrow(df_def_clean[df_def_clean$Target == "1", ]), rep)

#Combinamos con la clase mayoritaria
df_def_clean_balanced <- bind_rows(
  df_def_clean %>% filter(Target == "1"),
  oversampled
)

table(df_def_clean_balanced$Target)

```

```

0    1
174 174

```

Análisis univariado de variables numéricas

```

# -----
# Análisis univariado
# -----

# a) Seleccionar variables numéricas
numeric_vars_def <- df_def_clean %>% dplyr::select(where(is.numeric))

pdf("outputs/images/03_univariado_num.pdf", width = 16, height = 10)

# Histogramas para variables numéricas
numeric_vars_def %>%
  gather(key = "Variable", value = "Value") %>%
  ggplot(aes(x = Value)) +
  geom_histogram(bins = 30, fill = "skyblue", color = "black") +
  facet_wrap(~ Variable, scales = "free") +
  theme_minimal() +
  labs(title = "Distribuciones de variables numéricas en conjunto de datos final")

dev.off()

```

```

pdf
2

```

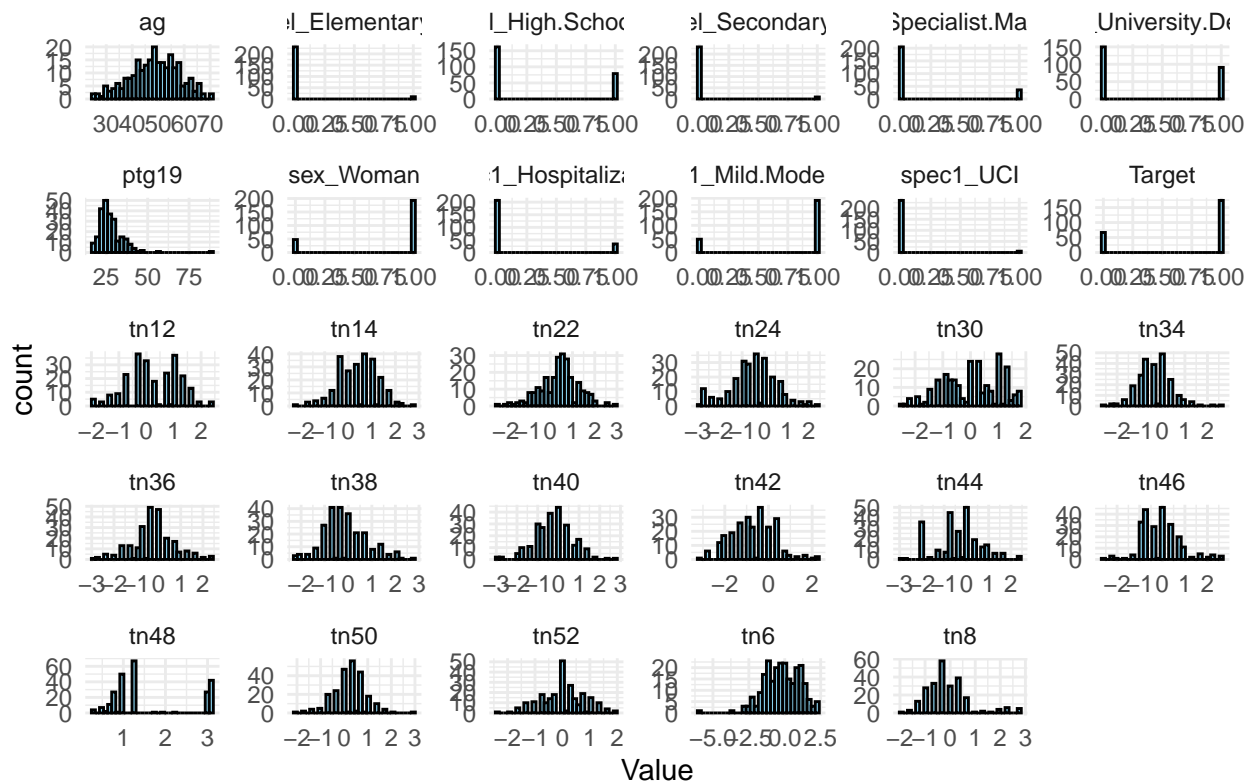
```

# Histogramas para variables numéricas
numeric_vars_def %>%
  gather(key = "Variable", value = "Value") %>%
  ggplot(aes(x = Value)) +
  geom_histogram(bins = 30, fill = "skyblue", color = "black") +
  facet_wrap(~ Variable, scales = "free") +
  theme_minimal() +
  labs(title = "Distribuciones de variables numéricas en conjunto de datos final")

```



## Distribuciones de variables numéricas en conjunto de datos final



## Análisis univariado de variables categóricas

# b) Variables categóricas

# Seleccionar columnas específicas y transformarlas a factor: mantenemos el nombre cambiado  
# para evitar afectar el dataset generado con columnas numéricas que usaremos en los modelos

```
df_def_clean_factors <- df_def_clean %>%
  mutate(across(
    where(~ is.numeric(.) && all(. %in% c(0, 1)) && n_distinct(.) == 2),
    as.factor
  ))
```

# b.1) Selección de variables categóricas

```
categorical_vars_def <- df_def_clean_factors %>% dplyr::select(where(is.factor))
```

```
pdf("outputs/images/04_categóricas.pdf", width = 16, height = 10)
```

# Gráficos de barras para variables categóricas

```
categorical_vars_def %>%
  gather(key = "Variable", value = "Value") %>%
  ggplot(aes(x = Value, fill = Variable)) +
  geom_bar() +
  facet_wrap(~ Variable, scales = "free") +
  theme_minimal() +
  labs(title = "Distribuciones de variables categóricas (dataset final)")
```

```
dev.off()
```

pdf

```
# Gráficos de barras para variables categóricas
```

```
categorical_vars_def %>%
```

```
  gather(key = "Variable", value = "Value") %>%
```

```
  ggplot(aes(x = Value, fill = Variable)) +
```

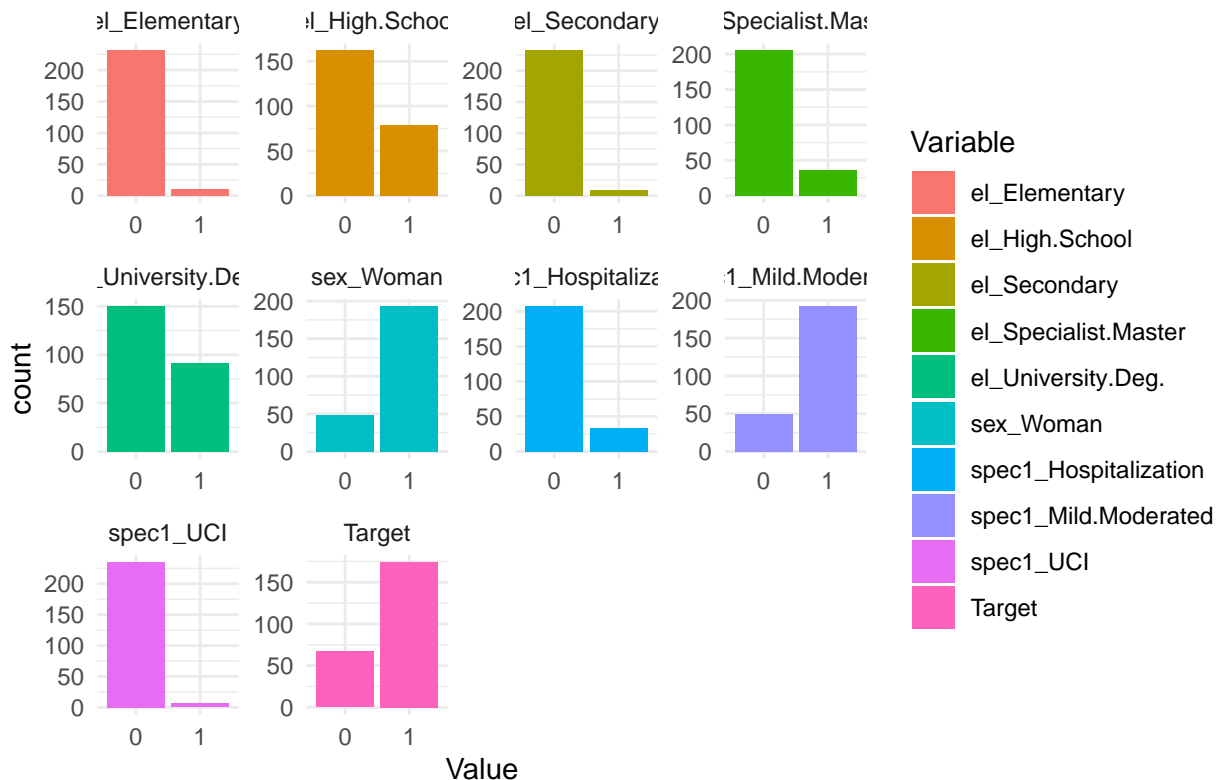
```
  geom_bar() +
```

```
  facet_wrap(~ Variable, scales = "free") +
```

```
  theme_minimal() +
```

```
  labs(title = "Distribuciones de variables categóricas (dataset final)")
```

## Distribuciones de variables categóricas (dataset final)



## Correlación y relaciones multivariadas

```
# -----
```

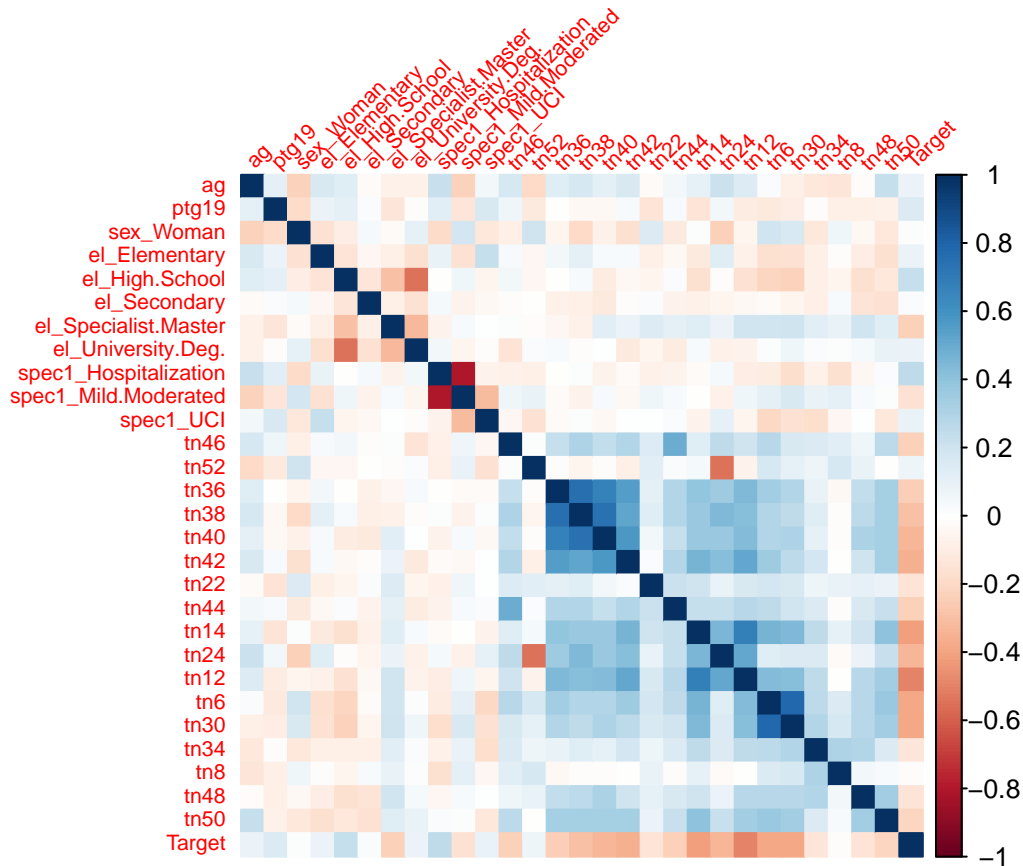
```
# Correlación y relaciones multivariadas
```

```
# -----
```

```
# a) Correlación entre variables numéricas
```

```
cor_matrix_def <- cor(numeric_vars_def, use = "pairwise.complete.obs")
```

```
corrplot(cor_matrix_def, method = "color", type = "full", tl.cex = 0.7, tl.srt = 45)
```



```
# Filtrar correlaciones mayores o iguales a |0.6| (sin incluir la diagonal)
threshold_def <- 0.6
high_correlations_def <- which(abs(cor_matrix_def) >= threshold_def & abs(cor_matrix_def) < 1, arr.ind = TRUE)

# Crear un data.frame con las variables altamente correlacionadas
cor_df_def <- data.frame(
  Var1 = rownames(cor_matrix_def)[high_correlations_def[, 1]],
  Var2 = colnames(cor_matrix_def)[high_correlations_def[, 2]],
  Correlation = cor_matrix_def[high_correlations_def]
)

# Eliminar duplicados (porque la matriz de correlación es simétrica)
cor_df_def <- cor_df_def[!duplicated(t(apply(cor_df_def, 1, sort))), ]
#write.csv(cor_df, "./outputs/correlaciones.csv")

# Mostrar las correlaciones
print(cor_df_def)
```

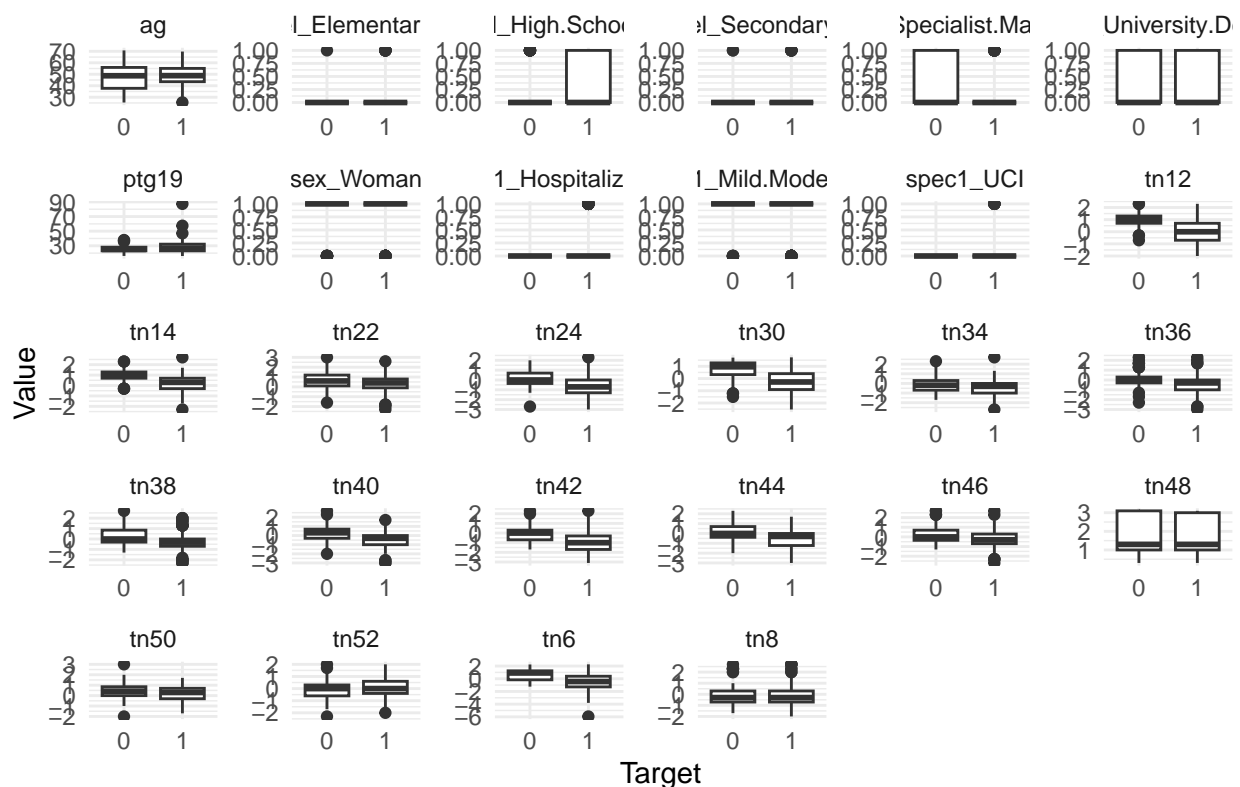
	Var1	Var2	Correlation
1	el_High.School	el_High.School	1.0000000
2	el_Secondary	el_Secondary	1.0000000
3	spec1_Mild.Moderated	spec1_Hospitalization	-0.8022447
5	tn52	tn52	1.0000000
6	tn38	tn36	0.7554991
7	tn40	tn36	0.6749011
9	tn38	tn38	1.0000000
10	tn40	tn38	0.7465677

13	tn12	tn14	0.6849510
15	tn6	tn6	1.0000000
16	tn30	tn6	0.7811306
18	tn30	tn30	1.0000000
19	tn34	tn34	1.0000000
20	tn50	tn50	1.0000000

Relaciones numéricas-categorías

```
# b) Relaciones numéricas-categorías
# Ajustando "Target" como variable objetivo
if ("Target" %in% colnames(df_def_clean)) {
  df_def_clean %>%
    mutate(Target = as.factor(Target)) %>% #Considerando que Target es binaria
    gather(key = "Variable", value = "Value", -Target) %>%
    filter(!is.na(Value)) %>%
    ggplot(aes(x = Target, y = Value)) +
    geom_boxplot() +
    facet_wrap(~ Variable, scales = "free") +
    theme_minimal() +
    labs(title = "Relaciones entre variables y Target")
}
```

## Relaciones entre variables y Target



Reducción de la dimensionalidad

```
# a) Eliminar variables con baja variabilidad: esto es "automático" con el paquete caret
nzv <- nearZeroVar(df_def_clean, saveMetrics = TRUE)
low_variability_vars <- rownames(nzv[nzv$nzv == TRUE, ])
cat("Variables con baja variabilidad eliminadas:", low_variability_vars, "\n")
```

```

Variables con baja variabilidad eliminadas: el_Elementary el_Secondary spec1_UCI
df_def_clean_low_var <- df_def_clean %>% dplyr::select(-all_of(low_variability_vars))

```

```

# b) Análisis de componentes principales (PCA)

```

```

pca_data_scaled <- scale(df_def_clean)

```

```

pca <- prcomp(pca_data_scaled, center = TRUE)
summary(pca) # Proporción de la varianza explicada

```

Importance of components:

	PC1	PC2	PC3	PC4	PC5	PC6	PC7
Standard deviation	2.4283	1.7280	1.36641	1.25357	1.22288	1.15161	1.14405
Proportion of Variance	0.2033	0.1030	0.06438	0.05419	0.05157	0.04573	0.04513
Cumulative Proportion	0.2033	0.3063	0.37067	0.42486	0.47643	0.52216	0.56729
	PC8	PC9	PC10	PC11	PC12	PC13	PC14
Standard deviation	1.09425	1.06641	1.01097	0.98501	0.97596	0.92702	0.88838
Proportion of Variance	0.04129	0.03922	0.03524	0.03346	0.03284	0.02963	0.02721
Cumulative Proportion	0.60858	0.64780	0.68304	0.71650	0.74934	0.77898	0.80619
	PC15	PC16	PC17	PC18	PC19	PC20	PC21
Standard deviation	0.8617	0.82799	0.78399	0.73707	0.71217	0.68631	0.64483
Proportion of Variance	0.0256	0.02364	0.02119	0.01873	0.01749	0.01624	0.01434
Cumulative Proportion	0.8318	0.85543	0.87663	0.89536	0.91285	0.92909	0.94343
	PC22	PC23	PC24	PC25	PC26	PC27	PC28
Standard deviation	0.61088	0.5491	0.50475	0.46039	0.42703	0.39811	0.30577
Proportion of Variance	0.01287	0.0104	0.00879	0.00731	0.00629	0.00547	0.00322
Cumulative Proportion	0.95630	0.9667	0.97548	0.98279	0.98908	0.99454	0.99776
	PC29						
Standard deviation	0.25462						
Proportion of Variance	0.00224						
Cumulative Proportion	1.00000						

Necesitamos unas 13 componentes para explicar el 80% de la varianza. Las 2 primeras solo explican un 50% de la varianza.

```

# Visualización más sencilla

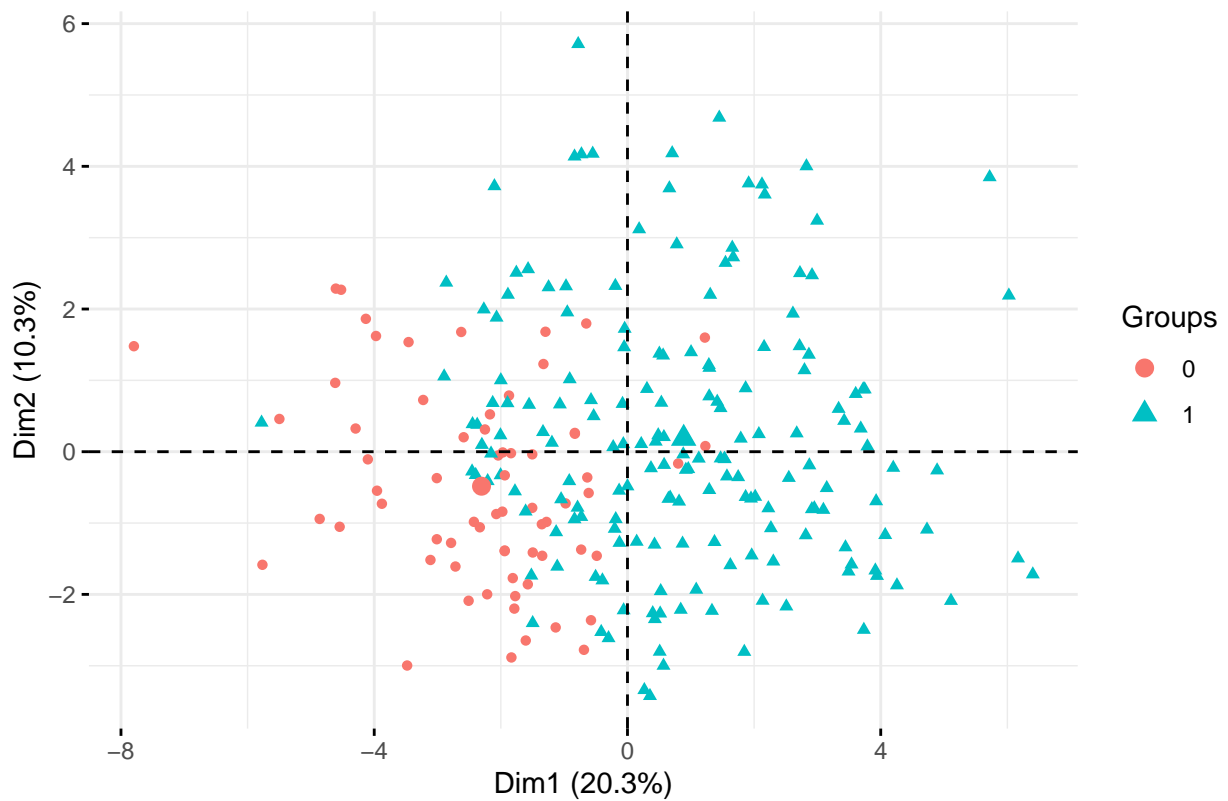
```

```

fviz_pca_ind(pca,
  habillage = df_def_clean$Target,
  label = "none",
  repel = TRUE,
  title = "Gráfico de PCA (Target como variable suplementaria)")

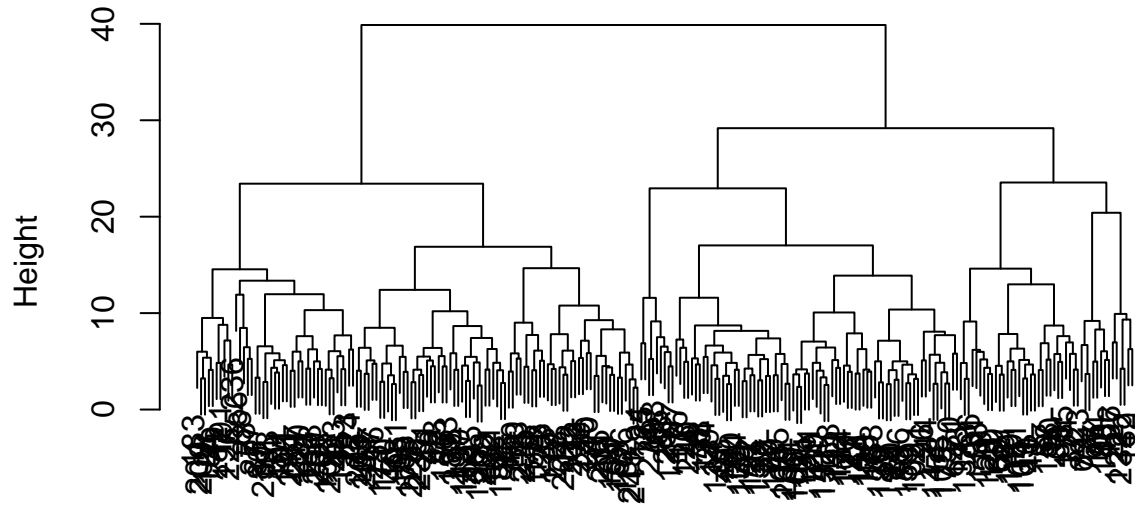
```

Gráfico de PCA (Target como variable suplementaria)



```
# -----
# Clustering para patrones
# -----
# Matriz de distancias y dendrograma
dist_matrix <- dist(scale(numeric_vars_def))
hclust_model <- hclust(dist_matrix, method = "ward.D2")
plot(hclust_model, main = "Dendrograma de clustering Jerárquico")
```

## Dendrograma de clustering Jerárquico



```
dist_matrix
hclust (*, "ward.D2")
```

```
# Cortar el dendrograma en 2 clústeres
clusters <- cutree(hclust_model, k = 2) # Cambiar "k" al número de clústeres sobre el que se cortará

# Ver asignaciones de clústeres para las observaciones
print(clusters)
```

```
[1] 1 1 2 1 1 2 2 2 1 2 1 2 1 2 2 1 1 2 1 1 2 1 2 2 2 1 1 1 2 1 1 2 1
[38] 1 1 2 1 1 1 1 2 2 2 2 2 2 2 2 2 2 2 2 1 1 1 1 1 1 2 1 1 2 1 1 1 1 1
[75] 2 1 1 1 1 2 1 1 1 1 2 1 1 1 1 1 1 1 1 2 1 1 1 1 2 1 2 1 1 1 2 1 1 1
[112] 1 2 1 2 1 2 2 1 1 1 1 1 2 1 1 1 2 1 1 2 2 2 1 1 1 1 1 1 1 1 1 2 1 1 1
[149] 1 2 2 1 2 1 2 1 1 1 1 1 1 2 1 1 1 2 1 1 1 2 2 2 1 1 1 1 2 1 2 2 2 2 2
[186] 2 2 1 2 2 2 2 2 1 2 2 2 1 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2
[223] 2 2 2 2 1 2 2 2 2 1 1 2 2 2 2 2 2 2 1 2
```

Verificar como funciona el clúster

```
# Asociar observaciones con cada clúster
cluster_observations <- split(row.names(numeric_vars_def), clusters)

# Imprimir observaciones de cada clúster
print("Observaciones en el clúster 1:")
```

```
[1] "Observaciones en el clúster 1:"
```

```
print(cluster_observations[[1]])
```

```
[1] "1"  "2"  "4"  "5"  "10" "12" "14" "17" "18" "20" "21" "23"
[13] "25" "29" "30" "31" "32" "34" "35" "37" "38" "39" "41" "42"
[25] "43" "44" "58" "59" "60" "61" "62" "63" "64" "66" "67" "69"
[37] "70" "71" "72" "73" "74" "76" "77" "78" "79" "81" "82" "83"
```

```
[49] "84" "86" "87" "88" "89" "90" "91" "92" "93" "94" "96" "97"
[61] "98" "99" "100" "102" "104" "105" "106" "107" "109" "110" "111" "112"
[73] "114" "116" "119" "120" "121" "122" "123" "125" "126" "127" "129" "130"
[85] "134" "135" "136" "137" "138" "139" "140" "141" "142" "143" "144" "146"
[97] "147" "148" "149" "152" "154" "156" "157" "158" "159" "160" "161" "162"
[109] "164" "165" "166" "168" "169" "170" "174" "175" "176" "177" "179" "188"
[121] "194" "198" "221" "227" "232" "233" "240"
```

```
print("Observaciones en el clúster 2:")
```

```
[1] "Observaciones en el clúster 2:"
```

```
print(cluster_observations[[2]])
```

```
[1] "3" "6" "7" "8" "9" "11" "13" "15" "16" "19" "22" "24"
[13] "26" "27" "28" "33" "36" "40" "45" "46" "47" "48" "49" "50"
[25] "51" "52" "53" "54" "55" "56" "57" "65" "68" "75" "80" "85"
[37] "95" "101" "103" "108" "113" "115" "117" "118" "124" "128" "131" "132"
[49] "133" "145" "150" "151" "153" "155" "163" "167" "171" "172" "173" "178"
[61] "180" "181" "182" "183" "184" "185" "186" "187" "189" "190" "191" "192"
[73] "193" "195" "196" "197" "199" "200" "201" "202" "203" "204" "205" "206"
[85] "207" "208" "209" "210" "211" "212" "213" "214" "215" "216" "217" "218"
[97] "219" "220" "222" "223" "224" "225" "226" "228" "229" "230" "231" "234"
[109] "235" "236" "237" "238" "239" "241"
```

```
# Filtrar datos por clústeres si se desea trabajar con subconjuntos
```

```
cluster_1 <- numeric_vars_def[clusters == 1, ]
```

```
cluster_2 <- numeric_vars_def[clusters == 2, ]
```

```
# Mostrar los subconjuntos
```

```
print("Datos del clúster 1:")
```

```
[1] "Datos del clúster 1:"
```

```
print(cluster_1)
```

	ag	ptg19	sex_Woman	el_Elementary	el_High.School	el_Secondary
1	26.9	38.23	1	0	1	0
2	60.5	26.34	0	0	0	0
4	44.6	24.14	1	0	0	0
5	61.9	32.80	0	0	0	0
10	51.2	28.66	1	0	0	0
12	36.4	28.45	1	0	1	0
14	48.8	30.98	1	0	0	0
17	54.7	35.75	1	0	0	0
18	47.5	25.44	1	0	1	0
20	46.3	33.01	1	0	1	0
21	33.5	25.10	1	0	0	0
23	53.9	26.57	0	1	0	0
25	55.4	22.68	1	0	0	1
29	39.7	28.13	1	0	0	0
30	46.1	26.41	1	0	0	0
31	63.0	26.47	0	0	0	0
32	42.6	22.21	1	0	0	0
34	63.0	19.70	1	0	0	0
35	39.2	27.71	1	0	1	0
37	59.7	22.10	1	0	1	0



38	43.0	26.25	0	0	0	0
39	44.7	22.62	1	0	1	0
41	50.4	39.94	1	0	0	0
42	56.6	25.80	1	0	0	0
43	46.7	21.95	1	0	0	0
44	48.8	24.01	1	0	1	0
58	47.0	21.29	1	0	1	0
59	51.4	23.05	0	0	1	0
60	67.2	32.21	1	0	1	0
61	53.8	25.71	1	0	1	0
62	55.3	28.67	0	0	1	0
63	57.8	39.08	1	0	0	0
64	57.9	25.44	1	1	0	0
66	56.7	29.97	0	0	1	0
67	40.3	26.89	1	0	0	0
69	65.7	36.88	1	1	0	0
70	43.6	17.62	1	0	0	0
71	48.5	32.25	0	0	0	0
72	41.8	23.07	1	0	1	0
73	69.8	33.24	0	1	0	0
74	54.4	32.81	1	0	1	0
76	48.1	28.52	1	0	0	1
77	45.5	28.19	1	1	0	0
78	48.3	34.36	1	0	0	0
79	53.7	21.85	1	0	0	0
81	61.8	27.21	1	0	0	0
82	54.9	37.80	1	1	0	0
83	49.8	23.65	1	0	1	0
84	50.3	29.34	1	0	1	0
86	42.0	25.64	1	0	0	1
87	49.5	31.92	1	0	0	0
88	51.4	30.61	1	0	0	1
89	55.6	22.83	1	0	1	0
90	55.6	35.77	0	0	0	0
91	44.8	21.33	1	0	1	0
92	50.2	40.32	1	0	0	0
93	43.3	16.05	1	0	0	0
94	37.9	33.67	1	0	0	1
96	26.4	42.40	1	0	1	0
97	56.4	26.71	0	0	1	0
98	53.3	33.21	0	0	1	0
99	62.6	36.92	0	1	0	0
100	39.8	20.56	1	0	1	0
102	48.2	24.35	1	0	1	0
104	47.2	17.38	1	0	0	0
105	41.5	22.81	1	0	1	0
106	49.1	29.40	0	0	0	0
107	42.2	57.46	1	0	0	0
109	47.5	29.26	1	0	1	0
110	56.9	24.44	0	0	1	0
111	54.3	33.45	1	0	1	0
112	40.5	25.64	0	0	1	0
114	48.6	38.29	1	0	0	0
116	37.9	41.27	0	0	1	0

119	56.5	27.24	0	1	0	0
120	53.2	34.17	1	0	0	0
121	50.7	27.21	1	0	0	1
122	61.3	30.59	0	0	0	0
123	55.4	25.75	0	0	1	0
125	39.1	33.57	0	0	0	1
126	46.0	44.23	0	0	1	0
127	47.6	33.87	1	0	0	0
129	59.9	28.93	1	0	1	0
130	40.0	20.00	1	0	0	0
134	59.6	37.97	1	0	1	0
135	44.6	20.90	1	0	0	0
136	48.4	20.62	1	0	1	0
137	52.4	36.31	1	0	0	0
138	48.8	30.21	0	0	1	0
139	48.4	22.31	1	0	1	0
140	48.4	27.97	1	0	1	0
141	51.7	17.94	1	0	1	0
142	53.9	21.15	1	0	1	0
143	45.2	24.16	0	0	1	0
144	41.9	28.95	1	0	0	0
146	50.6	22.32	1	1	0	0
147	37.0	25.30	1	0	1	0
148	57.6	32.78	1	0	0	0
149	57.7	37.72	1	0	0	0
152	54.5	23.84	1	0	1	0
154	49.0	28.76	1	0	0	0
156	50.6	22.50	1	0	1	0
157	43.2	29.30	1	0	0	0
158	57.2	29.34	1	0	0	0
159	54.2	22.82	1	0	1	0
160	52.3	30.35	1	0	0	0
161	39.8	22.34	1	0	0	0
162	54.9	19.54	1	0	1	0
164	44.3	30.51	0	1	0	0
165	40.6	27.79	0	0	0	0
166	54.8	29.79	1	0	0	0
168	47.3	36.06	1	0	1	0
169	50.2	24.01	1	0	0	0
170	38.2	19.14	1	0	0	0
174	44.2	28.28	1	0	0	0
175	39.5	28.26	1	0	1	0
176	46.9	22.47	1	0	0	0
177	35.2	26.40	1	0	1	0
179	53.1	21.21	1	0	0	0
188	31.7	23.18	1	0	1	0
194	43.5	19.43	0	0	0	0
198	67.5	21.88	1	0	0	1
221	34.5	33.24	1	0	0	1
227	35.8	47.02	1	0	1	0
232	54.5	39.36	1	0	0	0
233	48.6	35.23	1	0	0	0
240	51.1	23.83	0	0	0	0

el\_Specialist.Master el\_University.Deg. spec1\_Hospitalization

1	0	0	0
2	1	0	1
4	0	1	0
5	1	0	1
10	0	1	1
12	0	0	0
14	0	1	0
17	0	1	0
18	0	0	0
20	0	0	0
21	0	1	0
23	0	0	1
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30	1	0	0
31	0	1	1
32	0	1	0
34	0	1	1
35	0	0	1
37	0	0	1
38	0	1	0
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41	0	1	1
42	0	1	0
43	0	1	0
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59	0	0	1
60	0	0	1
61	0	0	0
62	0	0	1
63	0	1	1
64	0	0	0
66	0	0	1
67	0	1	0
69	0	0	1
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73	0	0	1
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76	0	0	0
77	0	0	0
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84	0	0	0
86	0	0	0
87	0	1	1
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89	0	0	0
90	0	1	1

91	0	0	0
92	0	1	0
93	0	1	0
94	0	0	0
96	0	0	0
97	0	0	0
98	0	0	1
99	0	0	0
100	0	0	0
102	0	0	0
104	0	1	1
105	0	0	0
106	1	0	0
107	0	1	1
109	0	0	0
110	0	0	1
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112	0	0	0
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121	0	0	0
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157	0	1	0
158	1	0	1
159	0	0	0
160	0	1	0
161	0	1	1
162	0	0	0

164	0	0	0
165	0	0	0
166	0	1	0
168	0	0	0
169	0	1	1
170	0	1	0
174	0	1	1
175	0	0	0
176	0	1	0
177	0	0	0
179	0	1	0
188	0	0	0
194	0	1	0
198	0	0	0
221	0	0	0
227	0	0	0
232	0	1	0
233	0	1	0
240	0	1	0
	spec1_Mild.Moderated	spec1_UCI	tn46 tn52 tn36 tn38 tn40 tn42 tn22 tn44
1	1	0 0.3 1.0 -1.00 -0.3 0.0 -1.3 0.4 -0.7	
2	0	0 1.0 0.0 0.00 0.3 0.0 0.0 0.4 -1.3	
4	1	0 0.0 -0.6 0.30 0.0 1.0 1.3 -0.9 1.0	
5	0	0 0.0 0.0 0.30 -0.3 0.3 -0.3 -0.2 -0.3	
10	0	0 -0.3 0.7 0.00 -0.3 -0.3 0.3 0.1 -0.7	
12	1	0 -0.7 0.7 -2.00 -1.0 -1.0 -1.7 0.1 0.0	
14	1	0 -0.7 -0.3 -0.30 -0.3 -1.0 -0.7 1.3 -0.7	
17	1	0 -0.7 -2.0 -2.00 -1.3 -1.3 -1.0 -0.5 -2.0	
18	1	0 -0.7 0.0 1.30 2.0 0.3 -1.3 -0.9 -0.7	
20	1	0 -1.0 0.0 0.00 -0.3 -0.3 -0.3 0.2 0.0	
21	1	0 0.3 1.0 -0.70 -0.3 -0.7 -2.7 -0.1 0.0	
23	0	0 -2.0 0.0 0.30 1.0 1.3 0.3 0.2 -1.3	
25	1	0 0.7 0.0 -0.30 -0.3 -0.3 -0.7 0.8 -0.7	
29	1	0 -0.7 -1.3 0.70 -0.7 -0.7 -2.0 0.3 1.0	
30	1	0 0.7 0.0 0.70 -0.7 0.3 -2.0 1.2 1.7	
31	0	0 -0.7 -1.0 -0.70 0.0 -0.7 -0.7 -0.2 1.0	
32	1	0 0.3 -0.3 1.70 -0.3 -0.7 -0.7 -0.2 -2.0	
34	0	0 0.0 -1.7 -0.30 0.3 1.0 -0.3 0.7 -0.3	
35	0	0 1.3 0.0 0.70 1.7 0.0 0.3 -1.2 0.0	
37	0	0 0.0 -0.7 1.30 0.3 0.0 0.0 0.2 0.0	
38	0	0 0.0 0.7 0.30 -0.3 0.0 -0.3 -1.3 -1.3	
39	1	0 0.3 0.6 0.00 0.0 -0.3 -1.0 1.4 -0.7	
41	0	0 -0.3 0.4 -0.30 -0.3 -0.3 -1.0 0.5 -0.7	
42	1	0 -0.3 0.4 -1.70 -1.0 -1.0 -2.0 0.5 -0.3	
43	1	0 -1.0 1.4 -1.00 -1.0 -1.0 -2.7 -0.6 -0.7	
44	1	0 0.3 1.0 0.00 -0.3 -1.3 -0.3 1.4 0.0	
58	1	0 0.3 0.0 -1.00 -0.7 -1.3 -2.7 1.6 0.0	
59	0	0 0.0 -0.7 -0.30 -0.7 0.0 0.0 -0.1 0.3	
60	0	0 0.3 0.0 -0.70 -0.7 -1.3 -1.7 0.7 -0.7	
61	1	0 -0.7 0.6 -2.30 -2.0 -2.0 -2.3 0.7 -2.0	
62	0	0 0.0 0.4 0.00 0.3 -0.3 -1.7 0.6 -0.3	
63	0	0 -0.3 -0.6 1.00 0.7 0.0 0.0 -0.2 0.3	
64	1	0 0.0 0.0 -0.70 -0.7 -0.3 -1.0 -0.6 -2.7	
66	0	0 -0.3 1.0 0.00 1.3 1.0 0.3 -0.1 1.0	

67	1	0	-1.0	1.3	0.30	0.0	-0.7	-1.0	0.5	-0.7
69	0	0	0.3	-1.3	-0.70	-0.3	-0.7	-2.0	0.5	-1.0
70	1	0	-1.0	-1.0	1.30	0.3	0.0	-0.3	-1.8	-2.0
71	1	0	0.3	0.3	-0.70	-0.7	-0.7	-2.3	2.6	0.0
72	0	0	-0.7	-0.7	0.70	0.3	-0.7	-0.7	0.4	0.0
73	0	0	0.7	-0.4	-1.00	0.0	-1.0	-1.3	0.7	-0.7
74	1	0	0.0	-0.3	-1.70	-1.0	-1.0	-0.7	-0.1	0.0
76	1	0	-1.0	-0.3	-1.00	-0.7	-1.0	-2.3	1.3	-2.0
77	1	0	-0.7	1.0	0.00	0.7	0.7	-1.0	1.0	-1.7
78	1	0	0.3	-0.3	-0.30	-0.7	-1.0	0.3	-2.3	-0.7
79	1	0	0.0	-0.3	-1.00	-0.7	-0.7	-0.7	0.0	-0.7
81	0	0	-0.7	0.4	-0.70	-0.7	-0.3	-2.3	0.5	-3.0
82	0	1	0.0	0.0	0.00	-1.0	-1.3	-0.3	-0.8	-0.3
83	1	0	0.3	0.0	-0.70	-0.3	0.0	-1.7	0.3	-0.7
84	1	0	-0.7	0.7	-0.30	-1.0	-0.7	-0.3	-0.1	-0.7
86	1	0	-0.7	2.0	0.00	-0.7	-0.7	-0.3	0.7	-0.7
87	0	0	1.7	0.7	0.70	-0.3	-0.7	-1.3	0.3	1.7
88	0	0	0.0	0.3	0.00	0.7	0.3	0.7	0.5	-0.3
89	1	0	0.0	0.7	-0.30	0.0	0.0	-0.3	0.6	-0.3
90	0	0	-2.7	-0.4	-0.30	0.3	0.0	-0.7	0.9	-2.0
91	1	0	-2.3	0.7	-0.30	-1.0	-2.0	-1.0	0.5	0.0
92	1	0	-0.3	-0.6	0.00	0.0	-0.3	-1.0	-0.5	-2.0
93	1	0	-1.0	-0.3	-1.70	-1.0	-1.0	-2.7	0.4	-0.7
94	1	0	-1.0	0.6	-2.30	-2.3	-3.0	-2.7	0.3	0.7
96	1	0	0.7	0.4	-0.70	-1.0	-1.7	-2.0	0.5	0.0
97	1	0	0.0	-0.7	0.00	0.0	0.0	0.0	-0.9	-0.3
98	0	0	0.0	-1.0	-1.00	-0.7	-1.7	0.0	-0.5	0.3
99	1	0	0.7	0.0	1.00	1.3	1.0	-0.3	-1.1	0.3
100	1	0	-2.3	0.3	-2.30	-1.8	-2.7	-1.0	0.0	-0.7
102	1	0	-1.0	0.7	0.00	0.0	0.0	-2.0	1.3	0.0
104	0	0	-0.7	0.3	-0.30	-0.3	-1.0	-1.3	1.0	-0.7
105	1	0	0.3	0.0	-1.30	-1.0	-1.0	-2.0	-0.9	-2.0
106	0	1	-1.3	-1.0	-1.07	-1.0	-1.3	-1.3	1.1	-1.0
107	0	0	-1.3	-0.7	-1.30	-1.0	-1.0	-2.0	-0.4	0.0
109	1	0	0.3	0.0	-1.00	-0.7	-1.3	-1.7	0.6	-0.7
110	0	0	-0.3	0.0	0.00	-0.7	-0.3	-0.3	-1.0	-2.0
111	1	0	1.0	-0.7	-1.70	-1.0	-1.7	-2.0	0.5	0.3
112	1	0	-1.0	0.3	-1.00	-0.7	-1.0	-1.0	-0.1	-2.0
114	1	0	-1.3	0.3	-0.30	-2.0	-0.3	-2.3	-0.9	-2.0
116	1	0	-0.7	-1.3	0.00	-0.3	-0.7	-1.3	-2.0	0.0
119	0	1	0.3	-1.3	0.30	0.7	0.0	0.3	1.8	0.3
120	1	0	-0.7	-0.3	-1.30	-1.0	-1.0	-3.1	-0.6	-2.0
121	1	0	2.7	0.6	0.30	0.0	0.0	0.3	1.8	1.3
122	0	1	0.0	-0.6	0.30	1.7	0.3	-1.0	1.0	0.3
123	0	0	-0.3	0.3	-0.70	-0.3	-0.3	-1.3	0.6	-2.0
125	0	0	-1.0	-0.7	-1.30	-1.7	-2.0	-2.3	-1.2	-2.0
126	1	0	-1.0	-1.0	-1.70	-1.7	-1.7	-1.7	-0.3	-2.0
127	1	0	-0.7	0.3	-0.30	-0.7	0.0	-0.3	0.7	-0.7
129	1	0	-0.3	0.4	-0.70	-1.0	-0.7	-2.0	-0.6	0.0
130	1	0	-1.0	0.0	-0.70	0.0	0.0	-0.3	-0.9	-2.0
134	1	0	0.0	0.3	-1.70	-1.0	-1.7	-1.3	0.0	-1.3
135	1	0	-0.7	0.3	0.70	-0.3	0.0	-1.0	0.6	-2.0
136	1	0	-0.7	-0.6	-0.30	-0.3	-1.0	-1.0	0.7	-2.0
137	1	0	-0.7	0.0	0.30	0.3	-0.3	-1.0	0.4	-0.7

138	1	0	-1.0	-1.4	-0.30	-0.3	-0.3	-0.3	-0.8	-2.0
139	1	0	-1.0	-0.4	0.70	1.3	0.3	-1.3	0.2	0.0
140	0	0	-0.7	0.0	-1.70	-0.7	-1.0	-2.3	0.7	-2.0
141	1	0	-0.3	-0.4	-2.00	-1.0	-1.3	-2.3	0.6	0.0
142	1	0	0.0	1.0	-0.70	-1.0	-0.3	0.3	0.7	0.3
143	1	0	-0.7	-0.7	0.00	0.3	0.0	-1.0	0.3	0.0
144	1	0	-1.0	0.0	-2.70	-2.3	-3.0	-1.0	-0.8	-2.0
146	1	0	0.3	-0.4	1.30	1.7	0.3	1.0	-0.3	1.7
147	1	0	-0.7	0.3	-0.30	-0.7	-1.7	-1.3	0.4	-0.7
148	0	0	-0.7	-1.0	-0.70	-1.3	-0.7	-0.7	-0.5	-0.3
149	0	0	-0.7	0.0	-1.30	-1.3	-2.0	-2.3	-0.1	-1.7
152	1	0	-0.3	0.0	-1.70	-1.0	-0.7	-2.0	0.2	0.3
154	1	0	-1.0	0.0	-2.70	-1.3	-1.0	-2.7	0.0	-2.0
156	1	0	-0.3	0.3	-1.30	-1.3	-1.3	-2.3	0.8	-0.7
157	1	0	-1.0	-0.4	-0.30	-0.7	0.0	-1.0	-0.8	-2.0
158	0	0	1.0	0.7	-0.30	0.3	-0.3	-0.3	1.7	1.3
159	1	0	0.0	-0.4	0.00	0.7	0.0	-1.0	1.2	0.3
160	1	0	-0.7	-1.6	-1.30	-0.7	-0.7	-2.0	0.5	-0.7
161	0	0	0.3	0.7	0.00	-0.3	0.0	0.7	0.7	-0.7
162	1	0	-0.3	0.0	-1.00	-0.7	-0.3	-1.7	0.3	-0.7
164	1	0	0.7	1.0	-0.70	0.0	-1.0	-1.3	-0.9	0.3
165	1	0	-1.0	1.3	-0.70	-0.3	0.0	-1.3	-0.3	-1.0
166	1	0	-0.3	0.6	-0.70	-0.3	-0.3	-0.7	0.0	-2.0
168	1	0	-0.7	0.7	-1.30	-1.0	-0.7	-1.7	0.8	-2.0
169	0	0	-0.7	0.4	-0.70	-0.7	-0.7	-1.3	2.1	0.0
170	1	0	0.0	1.7	1.00	0.0	0.0	-1.0	1.5	0.0
174	0	0	-1.0	-0.3	0.70	1.3	0.3	-1.3	-0.1	0.0
175	1	0	0.3	0.3	-0.30	-0.3	-0.3	-2.0	0.7	0.0
176	1	0	0.7	0.7	-1.30	-1.7	-1.0	-2.0	0.4	-2.0
177	1	0	-1.0	0.0	-2.30	-2.0	-2.7	-2.3	0.3	-2.0
179	1	0	-0.3	-0.7	0.00	0.0	0.0	-2.3	0.3	-2.0
188	1	0	-1.0	0.0	0.00	0.3	0.3	-0.3	-1.5	0.0
194	1	0	-0.7	-0.4	-0.30	0.0	-1.0	-1.7	-0.2	-0.7
198	0	0	-0.3	-1.4	0.00	0.3	0.3	0.3	-1.1	-0.7
221	1	0	-1.0	-1.0	-1.00	0.0	-0.3	-0.3	1.4	-2.0
227	0	1	-0.7	-1.0	-1.70	-1.0	-1.3	-2.0	-1.4	-0.7
232	0	1	-0.3	-0.4	-0.30	0.3	1.0	0.3	1.0	0.0
233	1	0	-0.7	1.0	-0.70	-0.7	-1.0	-1.0	0.0	-2.0
240	1	0	0.7	0.0	-3.00	-1.0	-0.3	0.3	-1.8	0.0

	tn14	tn24	tn12	tn6	tn30	tn34	tn8	tn48	tn50	Target
1	-1.7	-3.0	-2.0	-2.06	-2.30	-1.0	2.0	1.0	-1.7	1
2	1.0	0.0	0.0	0.86	0.70	0.3	-0.3	3.1	1.3	1
4	-0.3	-0.7	-0.3	-0.82	-0.43	-1.0	-0.7	1.3	-0.3	1
5	1.3	-0.3	1.3	-0.57	0.40	0.3	-0.3	3.1	-0.7	1
10	0.7	-0.7	0.0	-0.32	0.03	-2.0	0.3	0.8	0.7	1
12	0.3	-1.0	-0.3	-1.03	-0.03	0.0	-0.3	3.1	-0.7	1
14	0.7	-1.7	-0.7	-2.22	-1.14	-1.0	0.3	1.3	0.7	1
17	-0.3	0.7	0.0	0.04	-0.28	-0.3	0.0	0.7	-1.0	1
18	-0.3	-0.3	0.0	0.01	-0.42	-0.7	-0.3	1.0	-0.7	1
20	-0.3	-0.3	0.0	-1.52	-1.14	-0.7	-0.3	1.3	0.0	1
21	-0.7	-2.0	-1.3	-2.36	-1.46	0.0	0.3	0.8	-0.7	1
23	0.0	0.0	0.0	-1.80	-0.59	0.3	0.3	1.3	-1.3	1
25	1.0	-0.3	1.0	-0.93	-0.90	-0.7	-0.3	0.5	0.7	1
29	0.7	-1.0	0.3	-1.39	-0.75	-1.3	-0.3	1.0	-0.7	1

30	-0.7	-0.7	-0.7	-1.87	-0.78	0.0	0.3	1.3	0.0	1
31	0.7	-0.7	0.7	-0.96	-1.60	-0.7	-2.0	0.7	1.3	1
32	0.3	-0.7	0.7	0.24	-0.42	-0.3	-0.3	1.3	0.7	1
34	1.0	1.0	1.3	0.07	0.06	0.0	-0.3	0.8	0.7	1
35	1.3	1.0	1.0	0.98	0.40	1.3	0.7	0.3	0.3	1
37	1.0	0.0	1.0	0.79	0.03	0.0	0.3	3.0	0.7	1
38	0.7	-1.0	0.7	-1.48	-1.60	-0.3	-0.3	1.3	-0.3	1
39	0.7	-0.3	1.0	0.22	-0.78	-0.3	-0.3	1.0	0.0	1
41	-0.3	-0.7	-0.3	0.04	0.03	-0.3	-0.3	1.3	0.0	1
42	-0.3	-1.7	-0.3	-0.93	-0.90	-0.3	0.0	1.0	-0.3	1
43	-0.3	-2.7	-1.3	-1.29	-1.50	0.0	2.3	1.0	0.0	1
44	-0.3	-2.3	-1.0	0.80	1.35	1.3	0.7	3.0	-1.0	1
58	-1.0	-1.0	-0.7	-0.59	-0.07	-0.3	0.7	3.1	0.3	1
59	-0.3	-0.3	-1.0	-2.91	-1.21	-0.3	-1.3	1.3	0.0	1
60	-0.3	-0.3	-1.0	-0.31	-0.93	-1.7	-0.3	0.3	0.3	1
61	0.3	-1.3	-0.7	-1.18	-0.90	-0.7	-0.3	0.5	0.5	1
62	0.0	-0.7	-1.3	-0.69	-0.90	-1.3	0.0	0.8	0.7	1
63	1.3	0.3	2.0	0.42	0.65	0.3	-0.7	3.1	1.7	1
64	0.0	-1.0	-0.3	-0.93	-0.59	-2.3	-0.7	0.5	-0.3	1
66	1.3	0.7	1.0	-0.36	-1.86	-1.0	-1.0	1.0	-0.7	1
67	-0.7	-2.3	-0.3	-0.71	-0.43	-1.0	-0.3	3.0	0.0	1
69	-0.3	0.0	-1.0	-1.22	-0.93	-1.0	0.0	0.7	-0.7	1
70	-0.3	-1.0	0.3	-0.01	-0.07	-1.0	0.7	0.8	0.3	1
71	-1.7	-1.0	-0.7	-0.24	-0.07	-0.7	0.7	0.9	0.3	1
72	0.3	-1.0	0.0	0.22	1.00	-1.3	0.0	1.3	0.0	1
73	0.0	0.7	-0.3	-1.22	-0.93	-0.3	-0.3	0.7	0.3	1
74	-0.3	-2.0	-2.0	-1.43	-0.59	-1.7	-1.0	0.7	0.3	1
76	-0.3	-1.0	-0.7	0.33	-0.42	-0.7	0.7	0.7	-1.7	1
77	-0.3	-1.7	0.0	-0.24	0.28	-1.3	-0.7	3.1	0.3	1
78	-0.7	0.0	-0.7	1.03	1.35	-1.0	-0.7	1.0	0.3	1
79	-0.3	-0.7	0.7	-0.44	-0.59	0.0	-0.3	1.0	-0.3	1
81	0.3	-0.7	0.0	-1.22	-0.60	-0.7	0.3	3.1	0.7	1
82	-0.3	0.0	-0.3	-0.32	0.03	-0.7	-0.3	1.0	-1.0	1
83	0.0	-0.7	-0.3	0.33	-0.07	-2.0	-0.3	0.8	0.0	1
84	0.3	-0.7	-0.3	0.54	0.03	-0.3	0.3	1.3	0.7	1
86	0.7	-2.7	0.3	1.50	1.35	0.0	0.7	1.0	-0.3	1
87	0.3	0.3	0.0	1.84	1.00	0.0	0.3	1.3	0.3	1
88	0.7	-0.3	-0.3	-0.44	-0.59	-1.0	1.7	1.3	0.3	1
89	0.0	-1.7	-0.3	-0.32	-1.22	-0.3	0.3	1.3	0.0	1
90	1.3	-0.3	1.7	0.54	0.34	0.0	-0.3	3.1	1.3	1
91	-0.3	-2.0	-0.3	-1.05	-0.78	-1.7	-1.0	1.3	0.3	1
92	-0.3	0.3	-0.3	1.03	0.34	-0.7	-0.3	0.8	0.8	1
93	0.0	-1.0	-0.3	-0.59	0.28	0.7	0.0	1.3	-0.7	1
94	0.0	-2.3	-0.7	-1.27	-0.75	-0.3	-1.0	0.7	-0.7	1
96	-1.0	-2.7	-2.0	-0.70	0.30	-0.3	0.3	1.0	0.0	1
97	1.0	0.7	0.3	-1.80	-1.21	0.3	-0.3	1.0	0.7	1
98	-0.7	-0.7	2.0	-1.80	-1.53	-0.3	-1.0	1.0	-0.7	1
99	-1.0	0.7	1.0	-2.77	-0.93	0.0	0.0	1.0	0.3	1
100	0.0	-1.3	-0.7	-1.03	-0.39	0.0	0.3	0.6	-0.7	1
102	0.3	0.0	-0.3	-1.40	-0.42	-1.0	-0.3	0.8	-0.7	1
104	0.0	0.0	-0.7	-1.75	-1.85	-1.0	-1.0	1.3	0.7	1
105	0.3	-1.0	-0.3	-2.33	-0.78	0.3	0.7	0.8	0.0	1
106	-0.3	-1.0	0.0	-1.40	-0.78	-0.7	-1.3	1.0	-0.7	1
107	-0.3	-1.0	-0.3	-0.36	1.00	0.0	-1.3	1.0	-0.7	1



109	-0.3	-1.0	-1.3	-0.36	0.28	-0.7	0.3	1.3	0.3	1
110	1.0	-0.7	0.7	-1.18	-0.59	0.3	-0.7	1.0	0.7	1
111	-1.3	-0.3	-0.3	-2.66	-1.84	-0.3	-0.3	1.3	-0.7	1
112	0.0	-1.0	-1.3	-1.63	-1.14	-1.3	-1.0	1.0	0.0	1
114	1.0	-1.0	0.3	-1.52	-1.14	-0.3	-0.7	1.0	-1.3	1
116	-2.3	0.0	-0.7	-0.67	-0.03	-0.7	-1.0	1.3	-0.3	1
119	1.0	2.3	0.0	-1.30	-0.90	-1.0	0.3	1.3	0.0	1
120	-1.3	-2.0	-2.0	-1.06	-0.59	-1.7	-1.7	0.7	0.0	1
121	0.3	-0.3	1.0	-0.07	0.34	-0.7	1.3	1.3	0.7	1
122	0.7	0.3	0.7	-1.48	-0.93	-1.0	-0.3	3.1	0.3	1
123	0.3	-1.0	-0.3	-2.91	-1.84	-1.0	-1.0	0.8	-0.3	1
125	-1.3	-2.0	-2.0	-1.63	-1.82	-1.3	-1.7	1.0	0.3	1
126	-0.3	-0.7	-0.3	-1.17	0.28	0.3	0.7	0.5	-1.0	1
127	0.7	-0.3	-1.0	-0.12	0.64	-0.3	-0.7	0.8	0.0	1
129	0.0	-1.7	-0.7	-2.54	-2.15	-2.7	0.0	0.7	0.0	1
130	-0.3	-1.0	-1.0	-1.63	-1.46	0.0	-0.3	3.0	-0.3	1
134	1.0	0.0	0.3	-1.80	-0.90	0.0	0.3	1.0	0.3	1
135	1.0	0.0	-1.0	-0.12	0.28	0.0	0.7	1.0	0.7	1
136	-0.7	-0.7	-0.7	-1.29	-0.07	-0.7	0.3	0.3	-1.3	1
137	0.0	-0.3	-0.3	-1.43	-2.15	-0.3	-0.7	1.3	0.3	1
138	0.0	0.7	0.7	-2.33	-0.78	-0.3	-0.3	1.3	0.0	1
139	0.0	-0.3	-0.7	0.80	1.00	-0.7	-0.3	1.3	0.0	1
140	-1.0	-1.0	-1.3	-0.94	-0.07	-0.3	-0.3	1.3	0.0	1
141	-0.7	-1.3	-1.7	-2.29	-1.53	-1.0	-0.3	0.8	0.0	1
142	-0.7	-2.7	0.0	-0.19	-0.90	-1.3	-0.3	1.0	-0.3	1
143	-0.3	0.7	1.0	-5.94	-1.50	0.0	-1.0	1.0	0.0	1
144	-0.7	-2.7	-1.3	-2.33	-1.85	-0.7	-0.3	0.8	-0.7	1
146	0.7	0.7	0.0	-0.69	-1.21	0.0	0.0	1.3	0.0	1
147	-0.3	-1.3	-0.3	-1.03	-0.39	-0.7	-0.3	3.0	-0.3	1
148	1.7	0.0	-0.7	-1.80	-1.53	-1.0	-1.3	1.3	1.0	1
149	0.7	-0.3	0.0	-0.56	0.03	-0.7	-0.7	0.8	0.3	1
152	-1.3	-1.7	-0.7	-1.92	-0.59	-1.7	-1.0	0.7	-0.3	1
154	-0.3	-1.3	-1.7	-1.52	0.07	0.0	0.3	1.3	0.3	1
156	-0.3	-1.0	-0.3	-1.43	-0.59	0.0	-0.7	1.0	0.0	1
157	-0.7	-1.3	-0.7	-1.40	-1.14	-0.7	0.3	1.3	0.3	1
158	1.3	-0.7	0.0	2.27	1.59	0.3	-0.7	3.0	0.0	1
159	-0.3	-0.3	-0.3	0.54	-1.21	-1.3	-0.3	1.0	0.0	1
160	0.3	-0.7	0.0	-0.56	-0.90	-0.3	0.0	3.0	0.0	1
161	-0.3	-1.7	0.0	-1.03	-2.53	-0.7	-0.7	3.0	-0.3	1
162	0.0	0.3	-0.7	-1.18	-0.90	-0.3	-0.7	1.3	0.3	1
164	0.0	0.0	0.0	-1.05	-1.50	-0.3	-0.3	0.8	-0.7	0
165	0.3	-2.3	-1.0	-0.12	1.00	0.0	0.0	1.0	0.3	1
166	-0.3	-1.3	-0.3	1.65	0.34	0.0	0.0	1.0	0.3	1
168	-1.7	-2.7	-1.3	-0.82	-1.50	-1.0	-1.0	3.1	0.3	1
169	0.3	-0.7	-0.3	0.42	-0.59	-0.7	-1.0	0.8	0.0	1
170	-0.3	-1.7	-0.7	-0.19	-0.39	-0.3	0.3	1.0	0.3	1
174	-1.0	-0.7	-0.7	0.10	1.35	-1.0	-1.3	3.0	1.0	1
175	0.3	-1.0	-0.3	0.53	1.39	-0.7	0.3	1.0	0.7	1
176	-1.0	-2.7	-1.0	-1.52	-0.78	-1.3	-0.3	1.0	0.3	1
177	-0.7	-2.7	-1.7	-2.60	-2.17	0.0	0.7	0.5	-0.7	1
179	0.3	0.0	-0.3	-0.69	-0.28	-0.7	0.0	1.3	0.7	1
188	0.7	0.0	0.7	1.13	0.68	-1.3	-1.0	0.8	-0.3	0
194	0.0	-1.3	0.7	-1.05	-0.78	0.0	-1.0	3.0	0.0	1
198	0.3	0.7	1.0	0.07	1.06	0.0	-0.3	0.8	0.0	0

```

221 -0.3  1.3  1.3 -0.79 -0.39 -1.3 -0.7  1.0 -2.0      0
227 -0.7 -0.3 -0.3 -3.80 -2.17 -2.3 -1.0  0.5 -1.3      1
232  0.0 -0.3 -0.3 -2.29 -0.90 -1.3  0.3  3.1  1.0      1
233  0.0 -2.7 -0.7  0.10  0.64  0.3  0.7  0.8  0.0      1
240  0.0  0.3  0.0 -0.32  0.65 -0.3 -0.3  1.3  0.0      1

```

```
print("Datos del clúster 2:")
```

```
[1] "Datos del clúster 2:"
```

```
print(cluster_2)
```

	ag	ptg19	sex_Woman	el_Elementary	el_High.School	el_Secondary
3	33.1	18.24000	1	0	0	0
6	43.0	32.70000	1	0	0	0
7	55.6	25.44000	1	0	0	0
8	53.6	18.24000	1	0	0	0
9	46.3	23.23000	1	0	0	0
11	58.7	26.01000	1	0	1	0
13	42.7	19.37000	1	0	0	0
15	48.7	28.31000	1	0	0	0
16	53.6	20.97000	1	0	0	0
19	55.5	23.65000	1	0	0	0
22	64.6	30.90000	1	0	0	0
24	46.6	20.51000	1	0	0	0
26	29.9	17.54000	1	0	0	0
27	56.9	27.41000	1	0	1	0
28	47.6	23.67000	0	0	1	0
33	45.3	23.53000	1	0	0	0
36	61.6	23.87000	1	0	0	0
40	48.7	25.28000	1	0	0	0
45	36.5	29.40000	1	0	0	0
46	65.1	21.64000	1	0	0	0
47	62.8	24.40000	1	0	0	0
48	52.3	19.43000	1	0	0	0
49	53.0	25.40000	0	0	0	0
50	57.6	23.28000	1	0	1	0
51	51.1	27.57000	1	0	0	0
52	45.2	26.25000	1	0	0	0
53	65.9	24.69000	1	0	0	0
54	31.1	22.97000	1	0	0	0
55	42.6	24.58000	0	0	0	0
56	33.3	27.19237	0	0	0	0
57	61.6	29.72000	0	0	0	0
65	55.2	27.70000	1	0	1	0
68	57.2	45.79000	0	0	1	0
75	25.5	22.21000	1	0	0	0
80	44.8	28.12000	1	0	1	0
85	65.9	25.92000	1	0	1	0
95	33.7	39.51000	1	0	0	0
101	52.7	35.62000	1	0	1	0
103	42.7	20.95000	1	0	0	0
108	39.9	35.13000	1	0	0	0
113	58.3	23.06000	1	0	1	0
115	44.3	27.55000	1	0	0	0

117	47.1	24.17000	1	0	0	0
118	50.8	22.18000	1	0	1	0
124	57.0	40.61000	0	0	1	0
128	40.1	30.63000	1	0	0	0
131	46.1	30.76000	1	0	0	0
132	58.3	22.97000	1	0	1	0
133	59.7	24.46000	1	0	1	0
145	52.0	24.17000	1	0	0	0
150	43.1	21.91000	1	0	0	0
151	45.9	41.29000	1	0	1	0
153	41.9	20.42000	1	0	0	0
155	59.2	25.38000	1	0	0	0
163	45.8	27.87000	1	0	0	0
167	30.2	24.14000	1	0	0	0
171	38.4	24.19000	1	0	0	0
172	49.6	25.37000	0	0	0	0
173	58.7	28.55000	1	0	0	0
178	38.2	30.56000	1	0	0	0
180	29.4	17.95000	1	0	0	0
181	42.3	33.41000	1	0	0	0
182	41.5	23.57000	1	0	0	0
183	29.4	24.12000	1	0	0	0
184	40.8	25.43000	1	0	0	0
185	65.3	35.22000	0	0	1	0
186	56.1	23.42000	1	0	0	0
187	33.8	20.78000	1	0	0	0
189	61.0	37.46000	1	0	0	0
190	50.7	25.96000	1	0	0	0
191	62.6	28.57000	0	0	0	0
192	54.3	23.95000	1	0	0	0
193	28.9	26.75000	1	0	0	0
195	65.8	28.06000	1	0	1	0
196	48.7	22.63000	1	0	1	0
197	70.8	24.39000	0	0	0	0
199	40.7	27.42241	1	0	0	0
200	64.6	27.85000	0	0	1	0
201	55.3	26.19000	0	0	1	0
202	54.1	24.85000	0	0	0	0
203	57.8	19.17000	1	0	1	0
204	30.1	21.73000	1	0	0	0
205	57.7	38.63000	1	0	1	0
206	31.9	21.80000	1	0	0	0
207	41.1	27.36000	1	0	0	0
208	63.5	29.96000	1	0	0	0
209	40.9	26.15119	1	0	0	0
210	48.3	25.08000	0	0	0	0
211	49.0	23.54000	1	0	0	0
212	50.1	30.64000	0	0	0	0
213	49.0	22.04000	1	0	0	0
214	55.8	34.02000	1	0	1	0
215	43.3	23.44000	1	0	0	0
216	35.9	20.18000	1	0	0	0
217	51.6	22.79000	1	0	0	0
218	51.8	36.18000	1	0	0	0

219	34.8	28.10000	1	0	0	0
220	51.4	35.75000	1	0	1	0
222	52.6	21.08000	1	0	0	0
223	50.7	25.11000	1	0	0	0
224	45.4	37.82000	1	0	0	0
225	33.0	20.02000	1	0	0	0
226	57.5	30.02000	1	0	0	0
228	37.9	24.93343	1	0	0	0
229	34.7	24.72016	1	0	0	0
230	42.4	26.62936	1	0	0	0
231	46.3	23.24000	1	0	1	0
234	37.0	21.33000	1	0	0	0
235	37.6	24.42000	1	0	0	0
236	51.0	87.31000	0	0	1	0
237	62.1	29.67000	1	0	0	0
238	63.2	27.35000	0	0	1	0
239	47.1	24.10000	0	0	0	0
241	25.4	16.14000	1	0	0	0

	el_Specialist.Master	el_University.Deg.	spec1_Hospitalization
3	0	0	0
6	1	0	0
7	1	0	0
8	0	1	0
9	1	0	0
11	0	0	0
13	0	1	0
15	0	1	0
16	1	0	0
19	1	0	0
22	1	0	0
24	1	0	0
26	1	0	0
27	0	0	0
28	0	0	0
33	1	0	0
36	0	1	0
40	0	1	0
45	0	1	0
46	0	0	0
47	0	0	0
48	0	0	0
49	0	1	0
50	0	0	0
51	0	0	0
52	0	1	0
53	1	0	0
54	1	0	0
55	1	0	0
56	1	0	0
57	0	1	0
65	0	0	0
68	0	0	0
75	0	1	0
80	0	0	0

85	0	0	0
95	0	1	0
101	0	0	0
103	0	1	0
108	0	1	0
113	0	0	0
115	0	1	0
117	1	0	0
118	0	0	0
124	0	0	0
128	0	0	0
131	0	1	0
132	0	0	0
133	0	0	0
145	0	1	0
150	0	1	0
151	0	0	0
153	1	0	0
155	0	1	0
163	0	1	0
167	0	0	0
171	0	1	0
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173	1	0	0
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184	0	1	0
185	0	0	0
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187	1	0	0
189	0	1	0
190	1	0	0
191	0	1	0
192	0	1	0
193	1	0	0
195	0	0	0
196	0	0	0
197	1	0	0
199	0	1	0
200	0	0	0
201	0	0	0
202	1	0	0
203	0	0	0
204	0	1	0
205	0	0	0
206	0	1	0
207	1	0	0
208	0	1	0
209	0	1	0
210	0	1	0
211	1	0	0

212	0	1	0		
213	0	1	0		
214	0	0	0		
215	1	0	0		
216	0	0	0		
217	0	0	0		
218	1	0	0		
219	1	0	0		
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222	0	1	0		
223	0	0	0		
224	0	0	0		
225	1	0	0		
226	1	0	0		
228	0	0	0		
229	0	0	0		
230	0	0	0		
231	0	0	0		
234	0	1	0		
235	1	0	0		
236	0	0	0		
237	0	1	0		
238	0	0	0		
239	1	0	0		
241	1	0	0		
	spec1_Mild.Moderated	spec1_UCI	tn46	tn52	tn36
3	1	0	0.7000000	2.000000000	1.0000000
6	1	0	0.3000000	0.400000010	-0.7000000
7	1	0	-0.3000000	-0.699999990	-0.3000000
8	1	0	-0.3000000	0.400000010	-0.3000000
9	1	0	-0.7000000	1.000000000	-1.7000000
11	1	0	0.0000000	-0.400000010	0.3000000
13	1	0	-0.7000000	1.000000000	-1.7000000
15	1	0	0.0000000	0.699999990	0.0000000
16	1	0	-0.3000000	-1.700000000	-0.3000000
19	1	0	-0.7000000	0.300000010	-0.3000000
22	1	0	-0.3000000	0.000000000	-0.7000000
24	1	0	0.3000000	1.300000000	0.0000000
26	1	0	-0.7000000	0.699999990	-1.3000000
27	1	0	0.0000000	1.000000000	0.0000000
28	1	0	0.3000000	-0.699999990	0.0000000
33	0	0	0.3000000	1.000000000	-1.3000000
36	1	0	0.0000000	0.699999990	1.0000000
40	1	0	0.3000000	-0.400000010	0.0000000
45	1	0	-0.7000000	1.400000000	-0.7000000
46	1	0	-0.3000000	0.000000000	0.0000000
47	1	0	0.7000000	1.400000000	-0.3000000
48	1	0	0.3000000	0.300000010	-0.7000000
49	1	0	1.3000000	-0.699999990	1.7000000
50	1	0	0.0000000	0.000000000	-0.3000000
51	1	0	2.0000000	0.000000000	-0.7000000
52	1	0	-0.7000000	0.000000000	-0.3000000
53	1	0	0.7000000	-2.300000000	0.0000000
54	1	0	0.0000000	-0.600000020	-0.3000000

55	1	0	-1.0000000	-1.300000000	-1.7000000
56	1	0	2.3000000	-0.400000010	-0.3000000
57	1	0	0.0000000	0.300000010	-0.3000000
65	1	0	1.0000000	1.300000000	1.7000000
68	1	0	0.7000000	0.899999980	0.0000000
75	1	0	1.7000000	-0.300000010	-1.3000000
80	1	0	-0.7000000	-1.700000000	0.7000000
85	1	0	0.0000000	0.600000020	-0.7000000
95	1	0	-1.0000000	0.600000020	-0.7000000
101	1	0	-0.3000000	-0.800000010	0.3000000
103	1	0	0.7000000	0.300000010	-0.3000000
108	1	0	0.7000000	-0.300000010	-0.3000000
113	1	0	0.7000000	0.000000000	2.3000000
115	1	0	-1.0000000	0.000000000	0.7000000
117	1	0	-1.0000000	1.700000000	-0.3000000
118	1	0	1.0000000	-0.400000010	0.3000000
124	1	0	2.3000000	0.300000010	0.0000000
128	1	0	-1.0000000	-0.600000020	1.3000000
131	1	0	-0.7000000	1.000000000	0.0000000
132	1	0	0.0000000	1.400000000	1.0000000
133	1	0	-1.7000000	-0.600000020	-0.7000000
145	1	0	1.0000000	0.300000010	0.3000000
150	1	0	1.7000000	0.300000010	-0.3000000
151	1	0	-0.7000000	-0.600000020	0.7000000
153	1	0	0.0000000	0.000000000	-2.0000000
155	1	0	0.3000000	0.000000000	0.7000000
163	1	0	0.0000000	0.000000000	0.7000000
167	1	0	0.0000000	0.400000010	-0.7000000
171	1	0	0.0000000	1.600000000	0.0000000
172	1	0	-1.3000000	0.000000000	-0.7000000
173	1	0	0.3000000	0.699999990	-1.3000000
178	1	0	0.7000000	0.300000010	-2.3000000
180	0	0	-1.0000000	0.600000020	0.3000000
181	1	0	0.3000000	0.000000000	-0.3000000
182	1	0	0.3000000	0.300000010	0.0000000
183	1	0	0.3000000	0.300000010	2.0000000
184	1	0	0.0000000	0.000000000	1.3000000
185	1	0	0.3000000	0.000000000	0.0000000
186	1	0	-0.3000000	1.700000000	0.3000000
187	1	0	-1.0000000	0.300000010	-0.7000000
189	1	0	0.7000000	-0.400000010	-0.3000000
190	1	0	0.7000000	-0.699999990	0.0000000
191	1	0	-0.7000000	-0.699999990	1.0000000
192	1	0	-0.3000000	0.000000000	0.7000000
193	1	0	-1.0000000	-0.600000020	0.3000000
195	1	0	2.7000000	-1.000000000	1.7000000
196	1	0	1.7000000	1.000000000	0.0000000
197	1	0	0.0000000	0.400000010	0.7000000
199	1	0	-2.3000000	0.300000010	0.3000000
200	1	0	0.3000000	-0.300000010	0.7000000
201	1	0	1.0000000	-1.000000000	1.7000000
202	1	0	1.0000000	-0.699999990	0.0000000
203	0	0	-0.3000000	1.000000000	0.0000000
204	1	0	0.0000000	-0.699999990	0.3000000

205	1	0	-0.3000000	0.400000010	2.0000000	
206	1	0	1.0000000	0.699999990	-0.3000000	
207	1	0	0.0000000	0.300000010	-0.3000000	
208	0	0	0.7000000	0.000000000	-0.3000000	
209	1	0	-0.7000000	-0.300000010	1.3000000	
210	1	0	-1.0000000	-1.700000000	0.0000000	
211	1	0	-1.0000000	-1.000000000	-0.3000000	
212	1	0	2.3000000	0.300000010	2.3000000	
213	1	0	-1.0000000	0.000000000	-0.3000000	
214	1	0	1.0000000	0.000000000	1.3000000	
215	0	0	0.0000000	1.300000000	0.0000000	
216	1	0	0.0000000	-0.300000010	-0.3000000	
217	1	0	2.0000000	-1.400000000	-0.7000000	
218	1	0	-0.3000000	0.400000010	-0.3000000	
219	1	0	-0.1936254	0.008326257	-0.3344785	
220	1	0	2.3000000	-0.300000010	0.3000000	
222	0	0	0.7000000	-0.300000010	-0.3000000	
223	1	0	1.0000000	-0.699999990	0.0000000	
224	0	0	0.0000000	0.699999990	2.3000000	
225	1	0	0.0000000	0.400000010	0.3000000	
226	1	0	-0.7000000	0.000000000	1.7000000	
228	1	0	0.3389087	0.154617766	-0.3216190	
229	1	0	0.0000000	1.300000000	-0.3000000	
230	1	0	0.0000000	0.000000000	0.0000000	
231	1	0	-0.7000000	-1.300000000	0.0000000	
234	1	0	-0.7000000	0.699999990	-1.7000000	
235	1	0	0.0000000	-0.300000010	0.0000000	
236	1	0	1.7000000	0.000000000	0.7000000	
237	1	0	0.0000000	-0.300000010	0.0000000	
238	1	0	2.7000000	-1.300000000	0.0000000	
239	1	0	0.0000000	0.000000000	-0.3000000	
241	1	0	-0.4050728	0.211006487	-0.5695441	
	tn38	tn40	tn42	tn22	tn44	tn14
3	0.3000000	1.00000000	0.3000000	-0.5000000	-1.0000000	0.3000000
6	-0.3000000	0.30000000	-0.3000000	1.0000000	0.0000000	0.7000000
7	-0.7000000	-0.30000000	-1.0000000	0.4000000	-0.7000000	0.3000000
8	0.0000000	-0.30000000	-1.7000000	0.3000000	1.0000000	0.7000000
9	-2.3000000	-0.30000000	-0.7000000	0.3000000	-2.0000000	-0.3000000
11	0.3000000	0.70000000	1.0000000	-0.3000000	-1.3000000	1.3000000
13	-1.3000000	-1.00000000	-1.7000000	0.9000000	0.0000000	1.3000000
15	-0.3000000	-1.00000000	-1.7000000	0.3000000	0.0000000	1.0000000
16	-0.7000000	0.30000000	0.0000000	-0.7000000	0.0000000	0.3000000
19	-0.7000000	0.00000000	-0.7000000	0.0000000	1.0000000	1.3000000
22	-0.7000000	-0.70000000	-1.3000000	0.2000000	1.0000000	1.0000000
24	0.7000000	0.70000000	-1.0000000	0.2000000	-0.7000000	-0.3000000
26	0.0000000	-1.00000000	-1.0000000	1.5000000	-0.7000000	1.7000000
27	-0.3000000	1.00000000	-0.3000000	-0.6000000	-0.3000000	0.7000000
28	1.3000000	1.30000000	-0.7000000	-0.1000000	0.0000000	0.7000000
33	-0.3000000	-0.30000000	0.0000000	0.7000000	-0.7000000	0.3000000
36	0.7000000	0.30000000	0.3000000	1.0000000	-1.3000000	1.0000000
40	-0.7000000	0.00000000	-2.0000000	1.0000000	1.0000000	0.3000000
45	-0.7000000	0.00000000	-0.3000000	0.4000000	-0.7000000	1.0000000
46	-0.7000000	-0.70000000	-0.3000000	1.8000000	-0.7000000	1.0000000
47	0.7000000	0.00000000	-0.3000000	-0.1000000	0.0000000	1.7000000



48	0.0000000	0.3000000	0.3000000	-0.7000000	-1.0000000	1.0000000
49	1.3000000	1.0000000	2.0000000	0.2000000	2.3000000	0.7000000
50	0.0000000	0.3000000	0.3000000	0.7000000	-1.3000000	1.3000000
51	0.0000000	0.0000000	-0.7000000	0.6000000	0.7000000	0.3000000
52	-0.3000000	-1.0000000	-1.7000000	0.6000000	0.0000000	1.0000000
53	0.3000000	-0.3000000	-1.0000000	1.2000000	0.3000000	0.7000000
54	0.0000000	0.0000000	0.0000000	1.0000000	-0.3000000	2.0000000
55	-0.7000000	-1.7000000	-1.0000000	1.6000000	-1.0000000	0.7000000
56	1.0000000	0.0000000	-0.3000000	0.6331074	2.3000000	0.7000000
57	0.7000000	-0.7000000	-0.3000000	1.5000000	0.3000000	0.0000000
65	2.0000000	1.7000000	-0.3000000	1.2000000	1.3000000	1.0000000
68	0.7000000	-1.7000000	0.3000000	0.9000000	1.7000000	1.3000000
75	-0.3000000	-1.0000000	-1.7000000	1.6000000	0.0000000	0.3000000
80	0.0000000	0.0000000	0.0000000	0.3000000	-0.7000000	1.3000000
85	-0.3000000	-0.3000000	-1.3000000	0.6000000	-0.3000000	0.3000000
95	-1.0000000	-0.3000000	-2.0000000	0.6000000	0.7000000	0.7000000
101	-0.3000000	0.0000000	-0.7000000	1.8000000	0.0000000	0.7000000
103	-0.3000000	-1.7000000	-1.3000000	0.3000000	-2.0000000	1.0000000
108	-1.0000000	0.3000000	-1.3000000	-0.1000000	0.7000000	-0.3000000
113	2.0000000	1.3000000	2.3000000	1.3000000	0.3000000	2.7000000
115	0.7000000	1.0000000	0.3000000	1.2000000	0.0000000	1.0000000
117	-1.0000000	0.0000000	-1.3000000	0.8000000	-0.3000000	0.3000000
118	-0.3000000	0.3000000	-0.3000000	1.7000000	1.3000000	1.3000000
124	1.0000000	0.3000000	-0.3000000	0.6000000	0.3000000	0.7000000
128	1.3000000	0.3000000	0.0000000	-0.4000000	-1.0000000	2.3000000
131	-0.3000000	-0.3000000	-0.3000000	0.8000000	-0.7000000	0.7000000
132	0.7000000	0.3000000	-0.3000000	1.0000000	0.0000000	1.7000000
133	-0.7000000	-0.3000000	-0.7000000	0.3000000	-1.3000000	1.0000000
145	0.3000000	0.0000000	-0.3000000	0.3000000	0.7000000	0.0000000
150	-0.7000000	-0.3000000	-1.7000000	1.0000000	1.0000000	1.0000000
151	0.0000000	0.0000000	0.3000000	1.3000000	-0.7000000	0.7000000
153	-2.0000000	-1.0000000	-1.3000000	0.4000000	-0.3000000	0.3000000
155	1.3000000	0.0000000	-1.0000000	1.0000000	-0.7000000	1.3000000
163	0.7000000	1.0000000	-0.3000000	1.3000000	0.0000000	0.7000000
167	-1.3000000	-1.0000000	-1.3000000	0.5000000	-0.3000000	1.7000000
171	-0.7000000	-0.7000000	-0.7000000	0.9000000	0.3000000	0.3000000
172	-0.7000000	-0.3000000	-1.0000000	0.2700000	1.0000000	0.7000000
173	-1.3000000	-1.3000000	-0.7000000	1.6000000	0.0000000	1.0000000
178	-0.7000000	-0.3000000	0.0000000	0.6000000	-0.3000000	1.3000000
180	0.0000000	0.0000000	-1.3000000	0.5000000	-0.3000000	2.3000000
181	0.0000000	0.3000000	-1.7000000	-0.4000000	-2.0000000	0.7000000
182	0.3000000	0.3000000	-1.3000000	0.7000000	-0.7000000	1.0000000
183	2.7000000	2.7000000	-1.0000000	0.8000000	0.0000000	1.3000000
184	0.0000000	0.0000000	0.7000000	-0.4000000	1.0000000	2.0000000
185	0.7000000	0.3000000	0.0000000	-0.7000000	0.0000000	0.0000000
186	0.3000000	0.7000000	0.3000000	0.2000000	0.0000000	1.3000000
187	-0.7000000	-0.7000000	-1.7000000	2.0000000	-0.3000000	0.3000000
189	-0.7000000	-0.7000000	-0.7000000	1.6000000	0.7000000	1.7000000
190	0.0000000	0.3000000	0.3000000	0.2000000	0.0000000	0.7000000
191	2.0000000	1.3000000	0.7000000	-1.6000000	-1.3000000	2.0000000
192	1.3000000	1.3000000	-1.7000000	0.4000000	-0.7000000	0.7000000
193	1.0000000	1.7000000	-0.3000000	1.2000000	-0.3000000	0.7000000
195	1.0000000	0.7000000	0.3000000	0.5000000	0.0000000	0.0000000
196	0.7000000	1.0000000	-1.0000000	0.6000000	1.0000000	1.7000000

197	0.7000000	1.0000000	0.0000000	0.0000000	1.0000000	1.7000000	
199	1.3000000	1.0000000	-0.7000000	0.2000000	0.0000000	0.7000000	
200	0.0000000	-1.0000000	-0.7000000	0.8000000	0.0000000	0.7000000	
201	1.7000000	0.7000000	0.3000000	0.8000000	0.3000000	1.0000000	
202	0.0000000	-0.7000000	-1.0000000	1.1000000	0.7000000	0.3000000	
203	1.0000000	-0.3000000	0.3000000	1.6000000	0.3000000	1.0000000	
204	0.0000000	0.3000000	0.3000000	1.8000000	0.0000000	0.7000000	
205	0.7000000	0.0000000	0.0000000	1.4000000	-0.3000000	1.3000000	
206	-0.3000000	-0.7000000	0.3000000	1.8000000	0.0000000	0.7000000	
207	-0.3000000	0.7000000	-0.7000000	-0.4000000	-0.3000000	-0.3000000	
208	0.0000000	0.0000000	0.0000000	0.1000000	1.0000000	1.0000000	
209	0.7000000	1.0000000	1.0000000	1.4000000	-0.7000000	1.7000000	
210	0.3000000	0.3000000	0.3000000	2.0000000	-2.0000000	1.0000000	
211	-0.7000000	1.0000000	0.7000000	0.4000000	-0.3000000	0.3000000	
212	2.3000000	2.3000000	2.3000000	1.0000000	2.3000000	1.7000000	
213	1.0000000	0.7000000	-0.7000000	0.8000000	-2.0000000	1.7000000	
214	2.0000000	1.3000000	0.0000000	0.6000000	-0.3000000	0.3000000	
215	-0.3000000	-0.3000000	-1.7000000	0.2000000	0.7000000	1.3000000	
216	-1.3000000	-2.0000000	-1.3000000	0.8000000	0.7000000	1.0000000	
217	1.0000000	1.3000000	0.3000000	1.4000000	0.7000000	0.7000000	
218	0.7000000	1.0000000	0.0000000	0.8000000	-2.0000000	0.0000000	
219	-0.4843276	0.36919106	-0.3884471	0.6442551	-0.2315443	0.5955351	
220	1.3000000	0.3000000	0.3000000	2.6000000	1.3000000	1.0000000	
222	0.3000000	-0.3000000	0.0000000	0.1000000	-0.7000000	0.0000000	
223	-1.0000000	-1.0000000	0.3000000	1.2000000	0.7000000	1.7000000	
224	2.0000000	1.3000000	1.7000000	3.0000000	-0.3000000	1.3000000	
225	0.0000000	0.7000000	-0.3000000	0.4000000	-0.3000000	1.0000000	
226	0.7000000	1.0000000	0.7000000	1.2000000	-0.7000000	1.0000000	
228	-0.2380424	-0.29804675	-0.3265751	0.3526449	-0.2024081	1.2087006	
229	0.7000000	0.3000000	0.0000000	0.6000000	-0.3000000	1.3000000	
230	-0.3000000	-0.3000000	0.0000000	0.8000000	0.7000000	-0.3000000	
231	0.0000000	0.0000000	0.0000000	0.8000000	0.0000000	0.7000000	
234	-0.7000000	-1.7000000	-1.3000000	0.6000000	0.0000000	1.0000000	
235	-1.0000000	0.7000000	1.7000000	-0.2000000	-0.3000000	1.3000000	
236	0.0000000	1.3000000	1.7000000	-0.8000000	1.3000000	0.7000000	
237	-0.3000000	-0.3000000	-0.3000000	0.6000000	0.3000000	1.0000000	
238	1.3000000	0.7000000	1.3000000	-0.6000000	1.7000000	1.0000000	
239	0.0000000	0.7000000	0.0000000	1.8000000	1.7000000	0.7000000	
241	-0.2685230	-0.03704518	-0.5917289	0.9404058	-0.4714624	1.0037005	
	tn24	tn12	tn6	tn30	tn34	tn8	tn48
3	-1.0000000	1.3000000	1.7000000	0.3200000	-1.0000000	0.3000000	3.100000
6	-0.7000000	0.0000000	1.3800000	1.7100000	0.0000000	2.7000000	1.300000
7	0.4000000	1.0000000	-1.1800000	-1.2200000	-0.3000000	0.0000000	1.300000
8	-0.7000000	0.0000000	-0.2000000	0.0300000	-1.0000000	-0.3000000	3.100000
9	-2.7000000	0.7000000	-0.0100000	-0.0700000	1.0000000	-0.3000000	1.000000
11	-0.3000000	1.0000000	0.4200000	-0.5900000	1.0000000	0.3000000	1.300000
13	-0.3000000	1.0000000	1.2600000	1.7100000	-0.7000000	2.7000000	0.800000
15	-1.0000000	0.3000000	1.0300000	1.3600000	0.0000000	-0.7000000	3.100000
16	1.0000000	0.0000000	-0.8100000	0.6500000	-0.7000000	-0.3000000	0.800000
19	-1.0000000	1.0000000	-0.3200000	-1.2100000	-0.3000000	-0.7000000	1.300000
22	-0.3000000	1.3000000	0.5900000	0.4000000	-0.3000000	0.0000000	3.000000
24	-1.3000000	1.0000000	0.9100000	1.0000000	0.0000000	-1.0000000	3.000000
26	0.3000000	0.7000000	-0.1500000	-0.1300000	0.0000000	0.3000000	3.100000
27	-1.3000000	1.0000000	0.1700000	0.3400000	-0.3000000	0.3000000	3.100000

28	0.00000000	0.00000000	0.92000000	0.28000000	0.30000000	2.70000000	3.10000000
33	-1.00000000	0.70000000	0.22000000	1.00000000	0.00000000	-0.30000000	1.30000000
36	0.00000000	1.00000000	0.46000000	1.73000000	-0.70000000	0.00000000	1.30000000
40	-0.30000000	0.00000000	-1.05000000	-0.78000000	-0.70000000	-1.00000000	3.00000000
45	-1.70000000	-0.30000000	-0.43000000	1.03000000	0.00000000	2.70000000	1.30000000
46	-0.30000000	1.00000000	0.98000000	0.06000000	-0.70000000	-0.30000000	3.00000000
47	-2.70000000	2.00000000	1.25000000	0.73000000	-1.00000000	-0.70000000	1.00000000
48	-1.00000000	-0.70000000	1.53000000	0.96000000	-0.70000000	-1.30000000	3.00000000
49	0.70000000	0.00000000	1.16000000	0.34000000	-0.70000000	-0.70000000	1.30000000
50	1.70000000	1.70000000	0.79000000	0.34000000	0.30000000	-0.70000000	3.00000000
51	-0.30000000	1.00000000	-0.07000000	-1.21000000	-0.70000000	-1.30000000	1.30000000
52	0.00000000	0.70000000	0.91000000	0.28000000	0.00000000	-0.30000000	1.30000000
53	1.30000000	1.00000000	0.98000000	0.40000000	-1.00000000	-0.70000000	0.80000000
54	0.30000000	1.70000000	-0.67000000	0.32000000	-1.30000000	-1.00000000	1.00000000
55	2.00000000	-0.30000000	-0.59000000	-0.78000000	0.00000000	0.30000000	1.00000000
56	0.70000000	1.30000000	1.98000000	1.39000000	2.30000000	2.30000000	3.10000000
57	-1.30000000	0.30000000	0.59000000	1.40000000	0.70000000	2.00000000	3.10000000
65	0.00000000	1.00000000	-0.05000000	-0.59000000	-1.00000000	0.00000000	1.30000000
68	0.00000000	1.70000000	-0.56000000	-0.90000000	1.30000000	1.30000000	3.10000000
75	-0.70000000	0.00000000	-1.52000000	-0.13000000	1.00000000	0.30000000	0.80000000
80	1.00000000	1.30000000	0.92000000	1.35000000	-1.00000000	-1.00000000	3.10000000
85	-1.30000000	1.00000000	0.85000000	1.73000000	0.00000000	-0.30000000	3.10000000
95	-1.30000000	0.00000000	0.89000000	1.03000000	0.30000000	0.00000000	1.30000000
101	1.80000000	0.70000000	-0.81000000	-1.22000000	0.30000000	0.30000000	3.10000000
103	-0.30000000	-0.30000000	1.50000000	1.35000000	0.70000000	0.30000000	1.00000000
108	-0.70000000	-0.30000000	0.53000000	0.32000000	1.00000000	0.30000000	3.00000000
113	-0.30000000	0.70000000	2.27000000	1.28000000	0.30000000	0.30000000	3.10000000
115	0.00000000	0.70000000	-0.71000000	-0.07000000	-1.30000000	-0.70000000	1.30000000
117	-2.70000000	-0.70000000	0.10000000	0.28000000	-0.70000000	-1.00000000	0.70000000
118	0.70000000	1.30000000	1.03000000	1.59000000	0.30000000	-0.30000000	1.00000000
124	0.00000000	0.30000000	-1.06000000	-0.28000000	-0.70000000	-0.70000000	1.00000000
128	0.30000000	1.70000000	0.69000000	1.35000000	0.70000000	0.00000000	3.10000000
131	0.00000000	0.70000000	-0.36000000	-0.42000000	2.70000000	0.70000000	3.00000000
132	0.30000000	1.70000000	0.04000000	0.03000000	-0.70000000	-0.30000000	1.30000000
133	0.30000000	1.00000000	-0.32000000	0.65000000	0.00000000	-0.30000000	0.60000000
145	-2.30000000	0.30000000	1.90000000	0.65000000	-0.70000000	0.00000000	1.00000000
150	-1.30000000	-0.30000000	0.10000000	0.64000000	0.00000000	-0.30000000	3.00000000
151	0.30000000	0.30000000	1.38000000	1.35000000	0.70000000	-0.70000000	1.30000000
153	-1.00000000	-0.30000000	1.03000000	1.35000000	0.00000000	2.30000000	3.00000000
155	0.00000000	1.70000000	1.28000000	1.59000000	-0.30000000	-0.70000000	1.30000000
163	0.00000000	0.70000000	-0.59000000	0.28000000	0.00000000	0.00000000	3.00000000
167	-1.70000000	1.30000000	1.01000000	0.32000000	-1.00000000	0.30000000	1.30000000
171	-2.30000000	1.00000000	1.25000000	1.03000000	-0.30000000	-1.00000000	3.00000000
172	-1.00000000	0.30000000	-0.83000000	1.35000000	0.00000000	-1.30000000	3.00000000
173	-1.00000000	1.00000000	1.03000000	0.96000000	-1.30000000	2.00000000	0.30000000
178	-0.30000000	1.70000000	1.73000000	1.39000000	0.00000000	-1.00000000	1.30000000
180	-0.30000000	0.50000000	0.94000000	1.17000000	-1.00000000	2.30000000	1.30000000
181	1.00000000	0.70000000	0.56000000	1.35000000	-1.00000000	0.30000000	3.00000000
182	-0.30000000	1.00000000	0.10000000	0.64000000	-0.30000000	0.30000000	1.00000000
183	0.00000000	1.00000000	-0.15000000	1.60000000	0.70000000	-0.30000000	3.10000000
184	1.00000000	1.70000000	2.19000000	1.71000000	0.70000000	-1.00000000	3.10000000
185	-1.30000000	-0.30000000	-0.31000000	-0.06000000	-0.30000000	-0.70000000	1.00000000
186	-0.70000000	1.30000000	0.17000000	0.97000000	-0.30000000	-0.30000000	3.10000000
187	-0.30000000	1.30000000	0.89000000	1.03000000	-0.70000000	0.00000000	1.30000000

189	-0.30000000	1.30000000	0.72000000	1.40000000	-1.70000000	-1.70000000	0.50000000
190	0.70000000	0.30000000	-0.19000000	-0.59000000	0.30000000	0.00000000	1.30000000
191	2.00000000	2.30000000	-0.83000000	0.40000000	-0.70000000	0.70000000	3.10000000
192	-0.70000000	0.70000000	2.14000000	0.96000000	0.30000000	-0.70000000	0.80000000
193	0.30000000	1.00000000	-1.24000000	0.73000000	-1.30000000	-1.30000000	1.30000000
195	0.30000000	0.30000000	-0.18000000	1.06000000	-1.00000000	0.00000000	1.00000000
196	0.00000000	1.70000000	0.91000000	1.28000000	-0.70000000	-1.30000000	0.80000000
197	0.30000000	2.30000000	2.25000000	1.66000000	-0.30000000	-1.30000000	3.00000000
199	-1.30000000	0.00000000	0.57000000	1.35000000	0.00000000	-1.30000000	1.00000000
200	0.00000000	1.00000000	1.25000000	0.73000000	0.00000000	1.00000000	1.30000000
201	0.30000000	0.30000000	1.40000000	0.97000000	0.70000000	-0.70000000	1.00000000
202	-0.30000000	0.70000000	1.28000000	0.97000000	-0.70000000	-0.70000000	3.10000000
203	-1.30000000	0.70000000	1.65000000	0.03000000	0.30000000	-0.70000000	1.00000000
204	0.70000000	1.30000000	1.37000000	1.39000000	0.30000000	0.00000000	1.30000000
205	-0.70000000	2.30000000	-0.07000000	0.97000000	-0.70000000	-0.30000000	1.00000000
206	-0.70000000	0.30000000	-0.19000000	1.04000000	0.70000000	0.70000000	0.80000000
207	-1.30000000	0.70000000	1.26000000	1.71000000	-0.30000000	0.30000000	3.10000000
208	-0.30000000	1.00000000	1.24000000	1.40000000	0.00000000	0.30000000	1.30000000
209	0.30000000	1.30000000	1.38000000	1.71000000	0.00000000	-1.00000000	1.30000000
210	1.00000000	1.30000000	-1.20000000	-0.78000000	-0.70000000	0.00000000	1.30000000
211	0.00000000	1.00000000	-0.01000000	0.28000000	1.30000000	0.00000000	3.10000000
212	1.70000000	1.70000000	1.53000000	1.28000000	0.30000000	0.00000000	3.10000000
213	-0.30000000	1.00000000	-1.05000000	0.29000000	0.30000000	2.70000000	1.30000000
214	-0.30000000	1.00000000	0.17000000	0.97000000	-0.30000000	0.00000000	3.10000000
215	-1.30000000	0.30000000	1.61000000	1.71000000	1.00000000	2.30000000	3.10000000
216	1.00000000	1.30000000	-0.07000000	0.67000000	0.00000000	-1.00000000	1.00000000
217	1.70000000	1.30000000	1.16000000	0.96000000	0.00000000	-0.30000000	3.00000000
218	-0.70000000	0.00000000	1.53000000	0.65000000	-1.30000000	-0.30000000	3.00000000
219	-0.41269192	0.8447977	0.3163209	0.9025117	-0.3208913	0.3926552	1.753575
220	2.00000000	1.30000000	-0.69000000	-0.90000000	0.00000000	-0.30000000	1.00000000
222	-0.70000000	0.00000000	-0.44000000	0.03000000	-0.70000000	0.00000000	3.10000000
223	0.70000000	1.30000000	2.02000000	1.59000000	0.00000000	2.00000000	1.30000000
224	0.00000000	1.00000000	1.03000000	1.00000000	0.00000000	-1.00000000	1.30000000
225	0.30000000	1.70000000	1.13000000	1.39000000	1.70000000	2.30000000	3.10000000
226	0.00000000	1.30000000	0.66000000	0.97000000	-1.00000000	-0.70000000	3.10000000
228	-0.11222345	1.2872673	1.0044066	0.8061191	-0.1814728	-0.3374685	1.937584
229	-1.00000000	1.00000000	0.65000000	1.03000000	1.00000000	-0.70000000	3.10000000
230	-1.00000000	1.00000000	1.38000000	0.64000000	0.30000000	-0.70000000	3.10000000
231	0.00000000	0.30000000	-0.70000000	0.29000000	-1.30000000	-1.30000000	3.10000000
234	0.00000000	0.70000000	0.77000000	1.03000000	-0.30000000	-0.30000000	0.80000000
235	0.30000000	1.30000000	1.13000000	1.03000000	-0.70000000	0.30000000	1.30000000
236	0.70000000	0.00000000	0.30000000	-0.28000000	0.30000000	-0.30000000	1.30000000
237	1.30000000	1.70000000	0.98000000	0.40000000	-1.30000000	0.70000000	1.30000000
238	1.30000000	1.00000000	1.24000000	0.40000000	0.70000000	0.70000000	1.30000000
239	0.70000000	0.00000000	-1.05000000	-0.78000000	0.30000000	-1.00000000	3.10000000
241	-0.09916293	1.0844352	0.2887386	0.6956717	-0.1530272	0.3005667	2.201543

tn50 Target

3	-0.30000000	0
6	0.70000000	1
7	1.30000000	1
8	0.70000000	1
9	0.30000000	1
11	0.30000000	1
13	0.00000000	1

15	0.3000000	1
16	0.7000000	1
19	0.7000000	0
22	-0.3000000	1
24	0.7000000	1
26	-0.3000000	1
27	0.3000000	1
28	1.3000000	1
33	0.0000000	1
36	0.7000000	1
40	0.3000000	1
45	0.0000000	1
46	0.3000000	0
47	1.3000000	0
48	0.3000000	0
49	0.3000000	0
50	1.0000000	1
51	0.0000000	0
52	0.0000000	0
53	-0.7000000	0
54	0.3000000	0
55	-1.0000000	0
56	3.0000000	0
57	1.0000000	1
65	0.7000000	1
68	0.7000000	1
75	-0.3000000	1
80	0.3000000	1
85	1.0000000	1
95	-0.3000000	1
101	0.3000000	1
103	0.7000000	1
108	0.3000000	1
113	1.3000000	1
115	0.7000000	1
117	0.3000000	1
118	0.0000000	1
124	0.3000000	1
128	1.0000000	0
131	0.7000000	1
132	0.3000000	1
133	0.0000000	1
145	1.3000000	1
150	0.0000000	1
151	-0.3000000	1
153	1.0000000	1
155	0.3000000	1
163	0.7000000	1
167	0.0000000	1
171	0.0000000	1
172	0.3000000	0
173	0.3000000	0
178	1.0000000	0
180	1.0000000	0

181	0.7000000	0
182	0.7000000	0
183	1.0000000	0
184	0.7000000	0
185	0.7000000	0
186	0.7000000	0
187	0.3000000	0
189	0.3000000	0
190	0.7000000	0
191	1.7000000	0
192	-0.3000000	0
193	0.0000000	0
195	0.0000000	0
196	0.3000000	0
197	2.0000000	0
199	0.3000000	1
200	1.3000000	0
201	1.0000000	0
202	1.7000000	1
203	0.3000000	0
204	0.0000000	0
205	0.3000000	1
206	-0.3000000	0
207	0.7000000	0
208	1.0000000	0
209	0.7000000	0
210	0.3000000	0
211	0.7000000	0
212	1.3000000	0
213	-0.3000000	0
214	0.0000000	0
215	-0.7000000	0
216	0.7000000	0
217	0.7000000	0
218	0.3000000	1
219	0.4238885	0
220	1.0000000	0
222	0.7000000	0
223	0.7000000	0
224	0.7000000	0
225	-0.3000000	0
226	1.7000000	0
228	0.5523615	0
229	0.3000000	0
230	1.0000000	0
231	0.7000000	1
234	1.0000000	1
235	1.0000000	0
236	0.7000000	1
237	0.0000000	0
238	1.0000000	0
239	0.3000000	0
241	0.1655717	0

```
# Agregar la asignación de clústeres como columna en los datos originales
numeric_vars_def$cluster <- clusters
```

```
# Visualizar los datos con la columna de clúster
print("Datos originales con asignación de clúster:")
```

```
[1] "Datos originales con asignación de clúster:"
```

```
print(head(numeric_vars_def))
```

```
      ag ptg19 sex_Woman el_Elementary el_High.School el_Secondary
1 26.9 38.23      1         0             1             0
2 60.5 26.34      0         0             0             0
3 33.1 18.24      1         0             0             0
4 44.6 24.14      1         0             0             0
5 61.9 32.80      0         0             0             0
6 43.0 32.70      1         0             0             0
      el_Specialist.Master el_University.Deg. spec1_Hospitalization
1              0              0              0
2              1              0              1
3              0              0              0
4              0              1              0
5              1              0              1
6              1              0              0
      spec1_Mild.Moderated spec1_UCI tn46 tn52 tn36 tn38 tn40 tn42 tn22 tn44 tn14
1              1              0 0.3 1.0 -1.0 -0.3 0.0 -1.3 0.4 -0.7 -1.7
2              0              0 1.0 0.0 0.0 0.3 0.0 0.0 0.4 -1.3 1.0
3              1              0 0.7 2.0 1.0 0.3 1.0 0.3 -0.5 -1.0 0.3
4              1              0 0.0 -0.6 0.3 0.0 1.0 1.3 -0.9 1.0 -0.3
5              0              0 0.0 0.0 0.3 -0.3 0.3 -0.3 -0.2 -0.3 1.3
6              1              0 0.3 0.4 -0.7 -0.3 0.3 -0.3 1.0 0.0 0.7
      tn24 tn12  tn6  tn30 tn34  tn8 tn48 tn50 Target cluster
1 -3.0 -2.0 -2.06 -2.30 -1.0 2.0 1.0 -1.7      1      1
2 0.0 0.0 0.86 0.70 0.3 -0.3 3.1 1.3      1      1
3 -1.0 1.3 1.70 0.32 -1.0 0.3 3.1 -0.3      0      2
4 -0.7 -0.3 -0.82 -0.43 -1.0 -0.7 1.3 -0.3      1      1
5 -0.3 1.3 -0.57 0.40 0.3 -0.3 3.1 -0.7      1      1
6 -0.7 0.0 1.38 1.71 0.0 2.7 1.3 0.7      1      2
```

```
# Opcional: Resumen por clúster
```

```
cluster_summary <- aggregate(. ~ cluster, data = numeric_vars_def, mean)
print("Resumen por clúster:")
```

```
[1] "Resumen por clúster:"
```

```
print(cluster_summary)
```

```
      cluster      ag      ptg19 sex_Woman el_Elementary el_High.School el_Secondary
1          1 49.16929 28.42748 0.7716535 0.07874016 0.4094488 0.07086614
2          2 48.21404 26.79841 0.8333333 0.00000000 0.2368421 0.00000000
      el_Specialist.Master el_University.Deg. spec1_Hospitalization
1          0.03937008          0.3937008          0.2677165
2          0.27192982          0.3596491          0.0000000
      spec1_Mild.Moderated spec1_UCI      tn46      tn52      tn36      tn38
1          0.6692913 0.04724409 -0.3299213 -0.02440945 -0.50448819 -0.3881890
2          0.9385965 0.00000000 0.1521071 0.07521009 0.03924876 0.1360448
```

```

      tn40      tn42      tn22      tn44      tn14      tn24      tn12
1 -0.5763780 -1.0795276 0.1574803 -0.67165354 -0.02519685 -0.7598425 -0.2503937
2  0.1239833 -0.3298838 0.6749159 -0.01759136  0.89919242 -0.1791586  0.8247061
      tn6      tn30      tn34      tn8      tn48      tn50      Target
1 -0.8196063 -0.4996850 -0.5622047 -0.2188976 1.319685 -0.01732283 0.9685039
2  0.5199076  0.6482834 -0.1250473 -0.0038969 1.953445  0.50826159 0.4473684

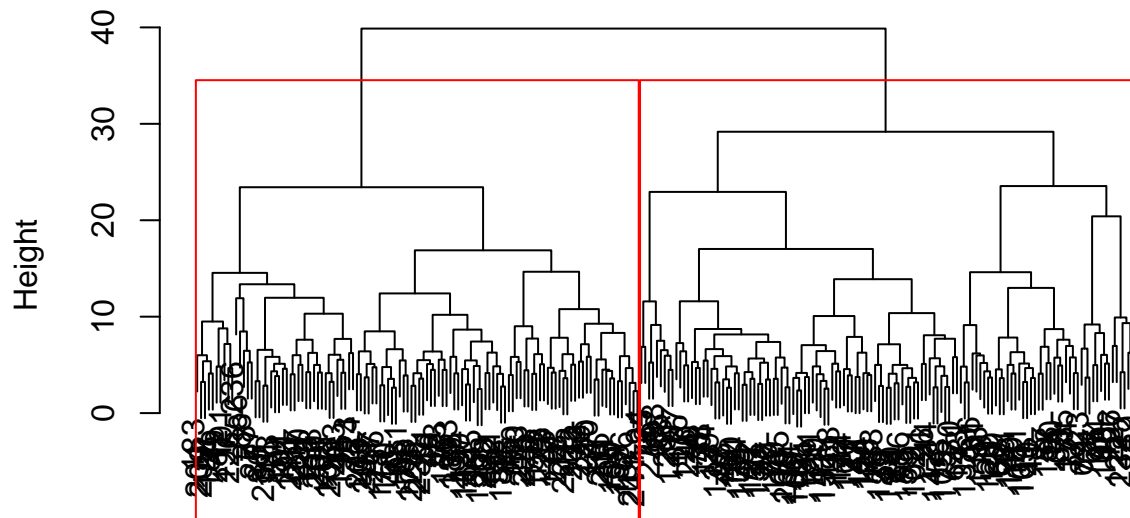
```

```

# Opcional: Agregar rectángulos al dendrograma para visualizar los clústeres
plot(hclust_model, main = "Dendrograma con clústeres")
rect.hclust(hclust_model, k = 2, border = "red") # Cambia "k" si es necesario

```

## Dendrograma con clústeres



```

dist_matrix
hclust (*, "ward.D2")

```

La organización por clústeres no parece ser de utilidad inicial.

## Estadística descriptiva e inferial básica.

Vamos a seleccionar las variables que son factores (reales) y las vamos a contrastar con chi-cuadrado

```

df_chi <- df_def_clean %>%
  dplyr::select(where(~ is.numeric(.) && all(. %in% c(0, 1)))) %>%
  mutate(Target = as.factor(df_def_clean$Target)) # Agregar Target como factor

```

```

# -----
# FILTRADO POR FRECUENCIA Y PRUEBAS DE CHI-CUADRADO
# -----

# 1. Calcular las frecuencias mínimas para cada variable dummy
low_freq_vars <- sapply(names(df_chi), function(var) {
  freq_table <- table(df_chi[[var]]) # Calcular la tabla de frecuencias
  min(freq_table) # Obtener la frecuencia mínima para cada variable
})

```



```

# 2. Definir el umbral de frecuencia mínima
threshold <- 5 # Cambiar el umbral si es necesario

# 3. Seleccionar variables con frecuencias mínimas mayores o iguales al umbral
valid_vars <- names(low_freq_vars[low_freq_vars >= threshold])

# 4. Filtrar el dataset para mantener solo las variables con frecuencias aceptables
df_chi_filtered <- df_chi %>%
  dplyr::select(all_of(valid_vars)) # Seleccionar las variables válidas

# 5. Realizar pruebas de chi-cuadrado para las variables filtradas contra Target
chi_results_filtered <- sapply(names(df_chi_filtered), function(var) {
  table_var <- table(df_chi_filtered[[var]], df_def$Target) # Crear tabla de contingencia
  chi_test <- chisq.test(table_var) # Realizar prueba de chi-cuadrado
  return(chi_test$p.value) # Extraer el valor p
})

# 6. Convertir los resultados a un data.frame
chi_results_filtered_df <- data.frame(
  Variable = names(chi_results_filtered), # Nombre de las variables
  P_Value = chi_results_filtered # Valores p
)

# 7. Ordenar los resultados por valor p
chi_results_filtered_df <- chi_results_filtered_df %>%
  arrange(P_Value)

# -----
# RESULTADOS Y VISUALIZACIÓN
# -----

# Mostrar el resumen de los resultados
print(chi_results_filtered_df)

```

	Variable	P_Value
Target	Target	2.866752e-53
spec1_Hospitalization	spec1_Hospitalization	2.176275e-04
el_High.School	el_High.School	4.464665e-04
el_Specialist.Master	el_Specialist.Master	6.144517e-04
spec1_Mild.Moderated	spec1_Mild.Moderated	2.872846e-02
el_University.Deg.	el_University.Deg.	2.598910e-01
spec1_UCI	spec1_UCI	2.810929e-01
el_Elementary	el_Elementary	3.560693e-01
sex_Woman	sex_Woman	9.553274e-01
el_Secondary	el_Secondary	9.987447e-01

```

# Guardar los resultados en un archivo CSV si es necesario
# write.csv(chi_results_filtered_df, "chi_squared_results_filtered.csv", row.names = FALSE)

# Opcional: Filtrar variables significativas (p < 0.05)
significant_vars_chi <- chi_results_filtered_df %>%
  filter(P_Value < 0.05)

# Mostrar las variables significativas

```

```
print(significant_vars_chi)
```

	Variable	P_Value
Target	Target	2.866752e-53
spec1_Hospitalization	spec1_Hospitalization	2.176275e-04
el_High.School	el_High.School	4.464665e-04
el_Specialist.Master	el_Specialist.Master	6.144517e-04
spec1_Mild.Moderated	spec1_Mild.Moderated	2.872846e-02

Sobre Chi-cuadrado Un p-valor bajo significa que la distribución de las categorías de la variable explicativa difiere significativamente entre las categorías de Target. Es decir, las frecuencias de las categorías no son independientes entre sí, lo que sugiere una relación entre la variable explicativa y Target.

Por ejemplo, si spec1\_Hospitalization tiene un p-valor bajo, esto implica que las frecuencias de hospitalización están distribuidas de manera diferente según la categoría de Target (No\_LC vs. LC\_Cog), lo que podría ser relevante para el modelo.

```
# Librerías necesarias
```

```
# Seleccionar variables numéricas continuas (excluyendo las dummy)
```

```
numeric_vars_log <- df_def_clean %>%
```

```
  dplyr::select(where(is.numeric)) %>% # Selecciona variables numéricas
```

```
  dplyr::select_if(~ length(unique(.)) > 2) # Excluye las dummies (solo deja aquellas con más de 2 val
```

```
numeric_vars_log$Target <- as.factor(df_def_clean$Target)
```

```
# Ajustar el modelo de regresión logística
```

```
logistic_model <- glm(Target ~ ., data = numeric_vars_log, family = binomial)
```

```
# Resumen del modelo
```

```
summary(logistic_model)
```

Call:

```
glm(formula = Target ~ ., family = binomial, data = numeric_vars_log)
```

Coefficients:

	Estimate	Std. Error	z value	Pr(> z )
(Intercept)	-2.32422	1.42136	-1.635	0.102004
ag	0.04644	0.02229	2.084	0.037188 *
ptg19	0.06075	0.03643	1.668	0.095369 .
tn46	-0.27194	0.26750	-1.017	0.309334
tn52	0.40342	0.34525	1.169	0.242605
tn36	0.56483	0.34420	1.641	0.100803
tn38	-0.21690	0.40333	-0.538	0.590730
tn40	-0.56926	0.38874	-1.464	0.143088
tn42	-0.37778	0.29118	-1.297	0.194483
tn22	-0.12377	0.23784	-0.520	0.602794
tn44	-0.26662	0.25533	-1.044	0.296398
tn14	-0.15135	0.40507	-0.374	0.708666
tn24	-0.29628	0.31218	-0.949	0.342588
tn12	-1.24769	0.35217	-3.543	0.000396 ***
tn6	-0.44161	0.28455	-1.552	0.120672
tn30	-0.43698	0.31917	-1.369	0.170971
tn34	0.16351	0.30306	0.540	0.589526
tn8	0.09556	0.22970	0.416	0.677394

```

tn48      0.03403    0.22367    0.152 0.879076
tn50      0.31727    0.36141    0.878 0.380020
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

(Dispersion parameter for binomial family taken to be 1)

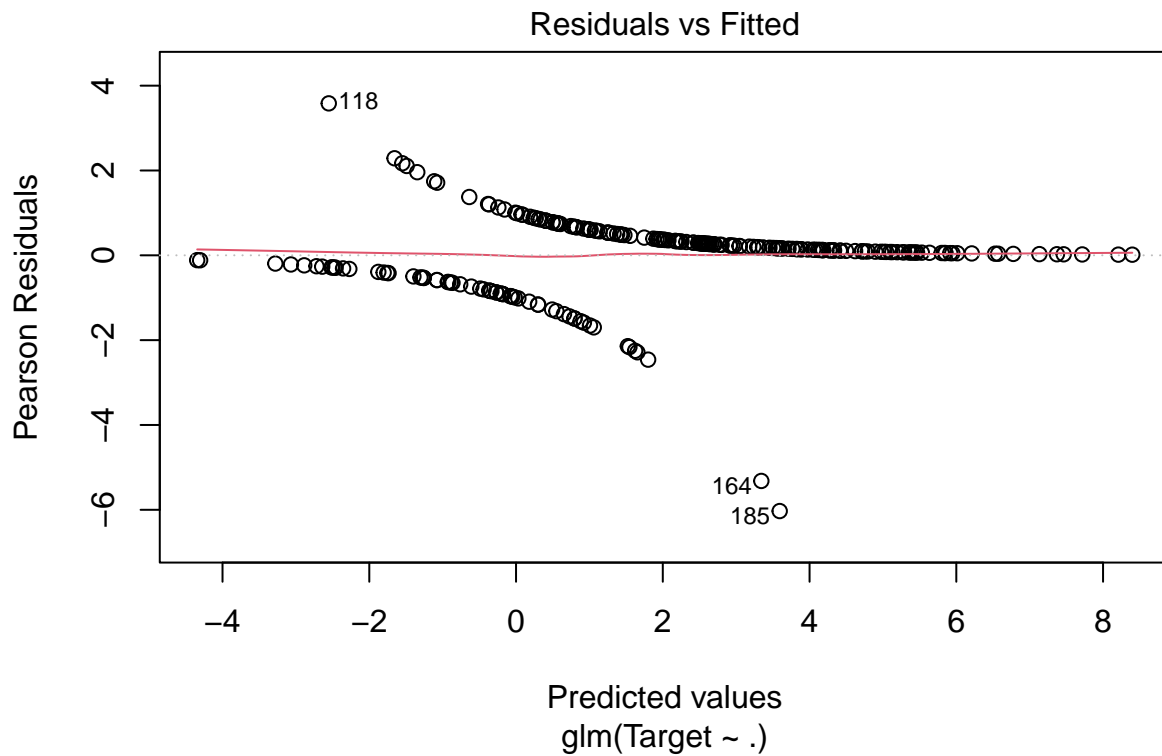
```

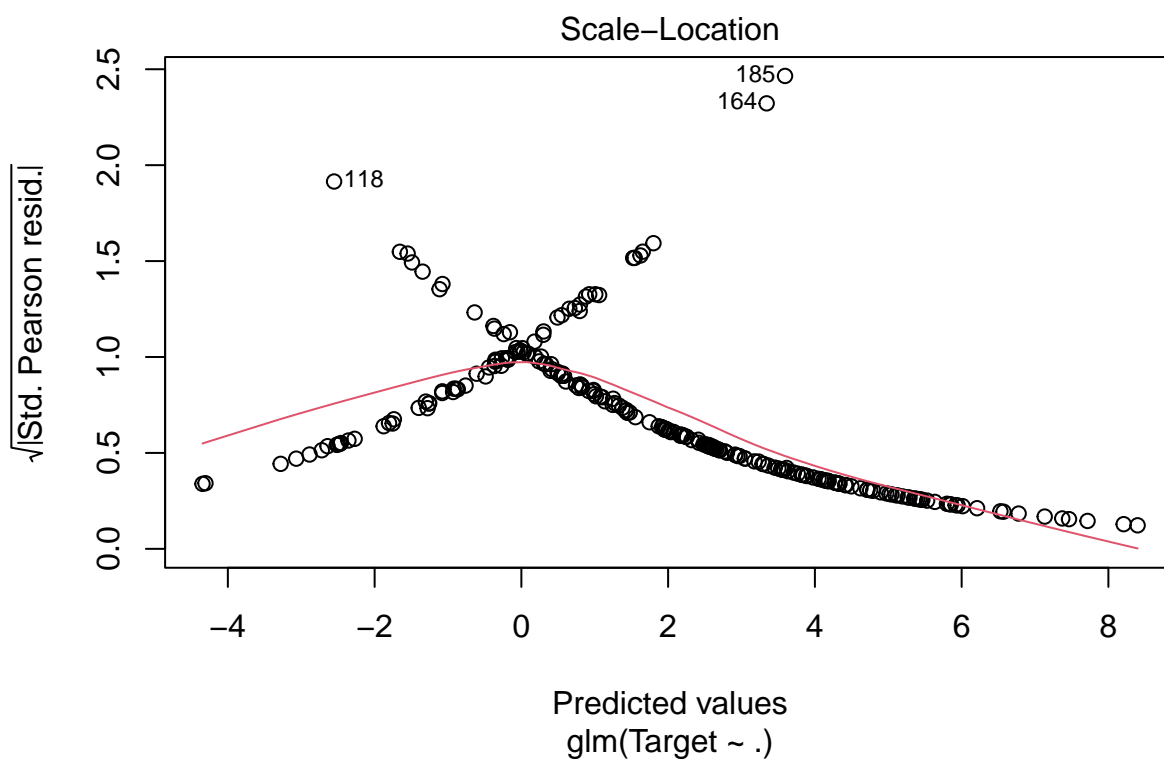
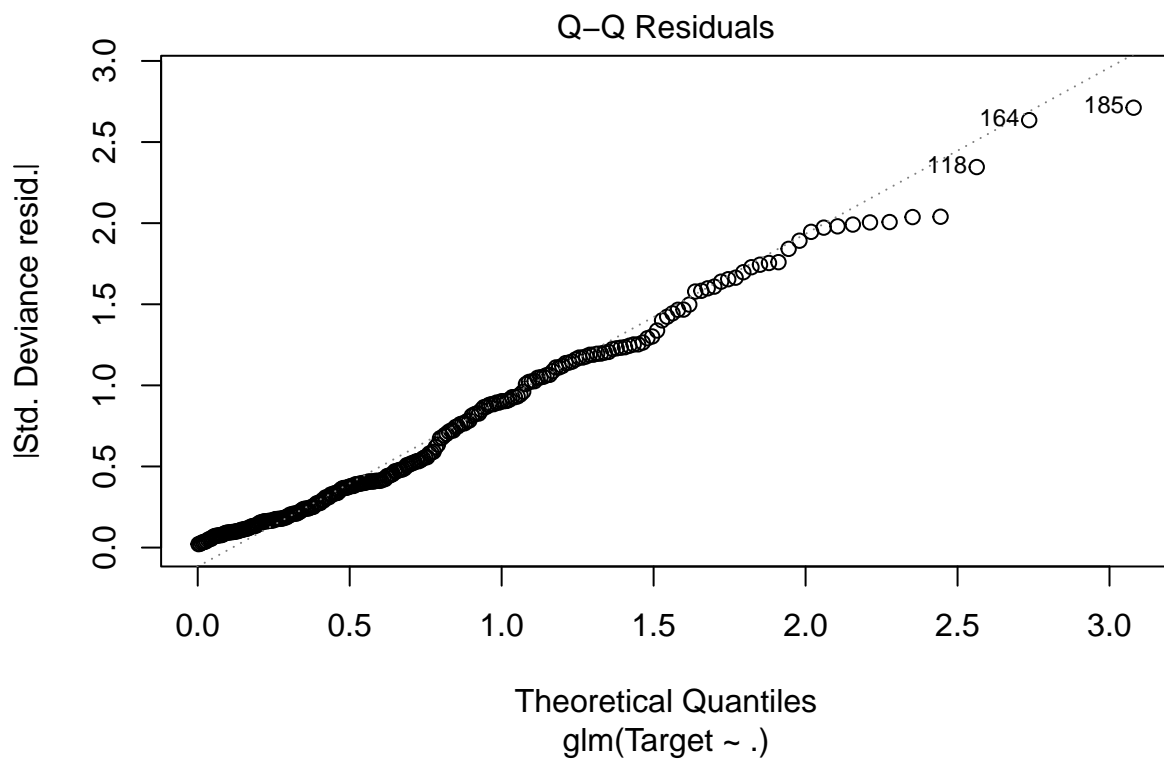
Null deviance: 284.89  on 240  degrees of freedom
Residual deviance: 168.21  on 221  degrees of freedom
AIC: 208.21

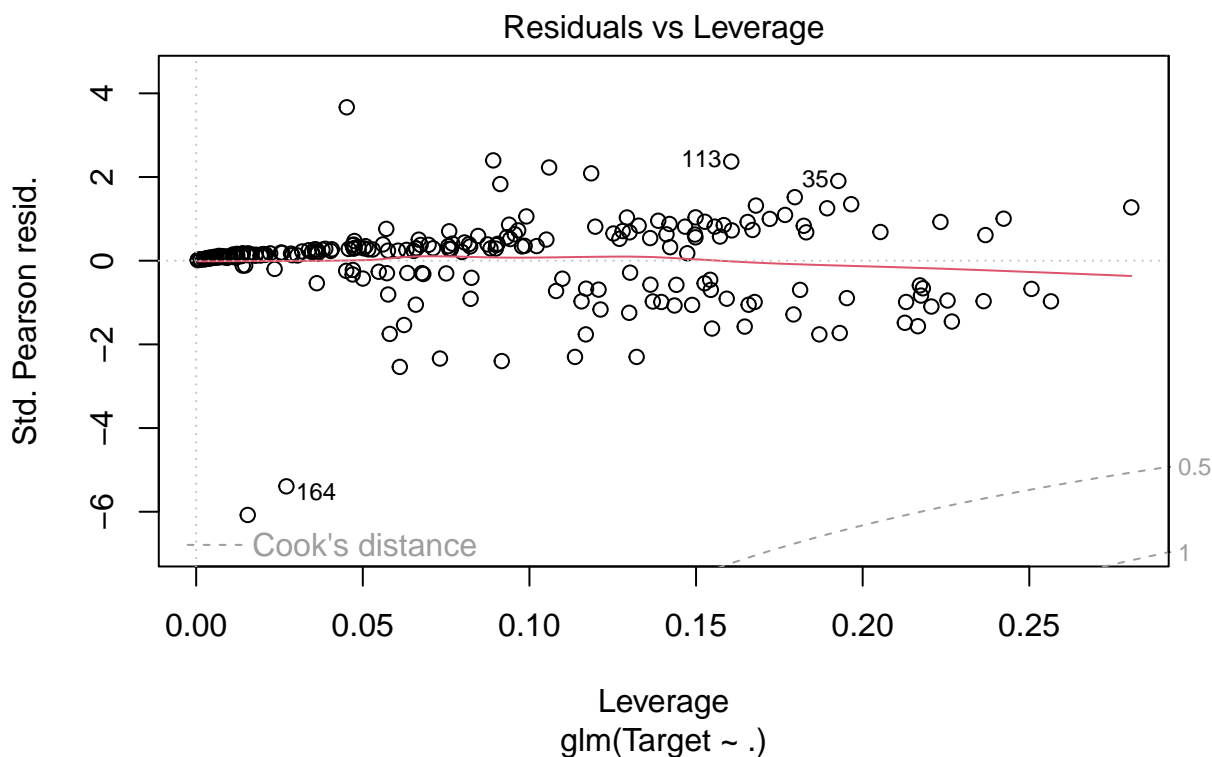
```

Number of Fisher Scoring iterations: 6

```
plot(logistic_model)
```







```
lda_model <- lda(Target ~ ., data = numeric_vars_log)
print(lda_model)
```

Call:

```
lda(Target ~ ., data = numeric_vars_log)
```

Prior probabilities of groups:

```
0      1
0.2780083 0.7219917
```

Group means:

	ag	ptg19	tn46	tn52	tn36	tn38	tn40
0	47.47761	25.81204	0.2259733	-0.06904551	0.1190203	0.2911807	0.2333448
1	49.19483	28.36726	-0.2281609	0.05804598	-0.3883333	-0.3063218	-0.4293103

	tn42	tn22	tn44	tn14	tn24	tn12	tn6
0	-0.1538321	0.6035883	0.01932217	0.9523573	0.07277495	0.98681344	0.5961114
1	-0.9448276	0.3247126	-0.50919540	0.2040230	-0.70000000	-0.02241379	-0.4871264

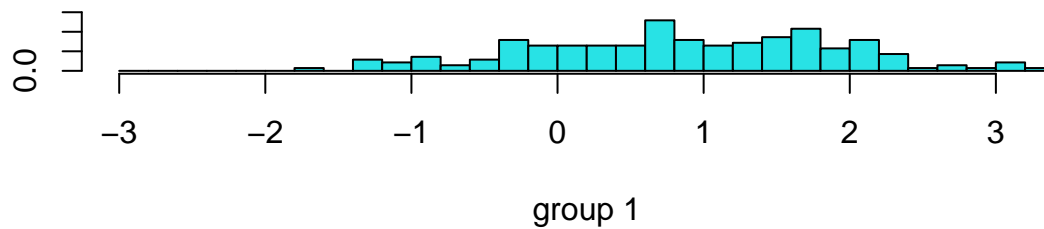
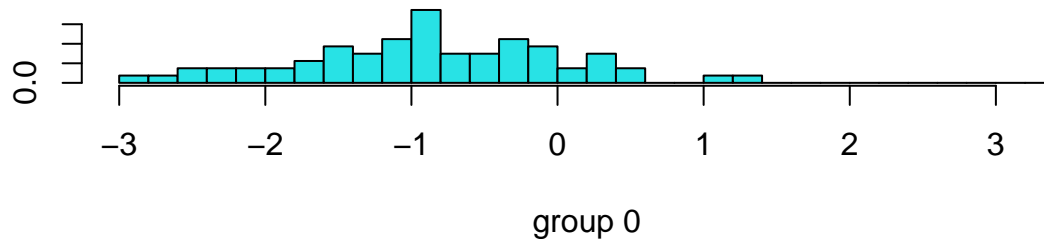
	tn30	tn34	tn8	tn48	tn50
0	0.6772284	-0.1888864	-0.09170517	1.837205	0.4618182
1	-0.2007471	-0.4195402	-0.12701149	1.535632	0.1425287

Coefficients of linear discriminants:

```
LD1
ag      0.04098263
ptg19   0.02624177
tn46   -0.25344260
tn52    0.27158291
tn36    0.40766393
tn38   -0.16422852
tn40   -0.29297120
```

```
tn42 -0.18203216
tn22 -0.06666090
tn44 -0.05936998
tn14 -0.18897618
tn24 -0.08990566
tn12 -0.61817323
tn6 -0.23181184
tn30 -0.18537675
tn34 0.16492408
tn8 0.09127756
tn48 0.08539564
tn50 0.04940253
```

```
plot(lda_model)
```



```
# Realizar el test de Kruskal-Wallis para cada variable
kruskal_results <- sapply(colnames(numeric_vars_log)[-which(colnames(numeric_vars_log) == "Target")], function(var) {
  kruskal.test(as.formula(paste(var, "~ Target")), data = numeric_vars_log)$p.value
})

# Convertir los resultados a un data.frame
kruskal_results_df <- data.frame(
  Variable = names(kruskal_results),
  P_Value = kruskal_results
)

# Ordenar por valor p
kruskal_results_df <- kruskal_results_df %>% arrange(P_Value)

# Mostrar los resultados
print(kruskal_results_df)
```

	Variable	P_Value
tn12	tn12	1.346705e-15

```

tn14      tn14 7.077679e-11
tn6       tn6 5.015610e-10
tn30      tn30 9.667375e-10
tn42      tn42 2.663421e-08
tn40      tn40 1.865985e-07
tn24      tn24 3.572774e-07
tn38      tn38 2.263308e-06
tn50      tn50 8.166312e-04
tn36      tn36 8.963381e-04
tn44      tn44 1.180907e-03
tn46      tn46 2.347338e-03
tn48      tn48 4.823635e-03
ptg19     ptg19 3.622734e-02
tn22      tn22 3.688904e-02
tn34      tn34 6.026432e-02
tn52      tn52 2.036706e-01
ag        ag 2.757166e-01
tn8       tn8 7.211850e-01

```

```

# Opcional: Filtrar variables significativas (p < 0.05)
significant_vars_krus <- kruskal_results_df %>%
  filter(P_Value < 0.05)

```

Generamos los modelos

```

# Suponiendo que `df_def_clean` ya contiene las características y la columna `Target`
# Dividir el dataset en entrenamiento (80%) y prueba (20%)
set.seed(2443) # Para reproducibilidad
#semillas interesantes: (0.8) 1234, 1237, 1238, 1339
#semillas interesantes: (0.7) 1339, 1441 por el recall
#semillas interesantes (0.75) 1234
#Tras mejoras (que es evitar el rol del desbalance y que la clase positiva sea 1), es decir,
#identificar los casos de long covid y 0.75: 1236, 1237, #1239
# tras mejoras y 0,8: 2435, 2429, 2433, 2435

trainIndex <- createDataPartition(df_def_clean$Target, p = 0.8, list = FALSE)
train_data <- df_def_clean[trainIndex, ]
test_data <- df_def_clean[-trainIndex, ]

# Definir la columna objetivo para entrenamiento y prueba
train_target <- train_data$Target
test_target <- test_data$Target

# Eliminar la columna objetivo de los datos de entrenamiento y prueba
train_data <- train_data %>% dplyr::select(-Target)
test_data <- test_data %>% dplyr::select(-Target)

# Convertir los datasets a matrices para XGBoost
train_matrix <- as.matrix(train_data)
test_matrix <- as.matrix(test_data)

# Crear DMatrix para XGBoost
dtrain <- xgb.DMatrix(data = train_matrix, label = train_target)
dtest <- xgb.DMatrix(data = test_matrix, label = test_target)

```

```

# Calcular el peso para la clase positiva
num_negativos <- sum(train_target == 0) # Casos de la clase 0
num_positivos <- sum(train_target == 1) # Casos de la clase 1

scale_pos_weight <- num_negativos / num_positivos

params <- list(
  objective = "binary:logistic", # Configuración para clasificación binaria
  eval_metric = "logloss",      # Métrica de evaluación para clasificación binaria
  scale_pos_weight = scale_pos_weight # Peso para manejar el desbalance de clases
)

# Entrenar el modelo de clasificación binaria con XGBoost
modelo_xgb <- xgboost(data = dtrain, params = params, nrounds = 500)

[1] train-logloss:0.563283
[2] train-logloss:0.481515
[3] train-logloss:0.419843
[4] train-logloss:0.372299
[5] train-logloss:0.333977
[6] train-logloss:0.302680
[7] train-logloss:0.273890
[8] train-logloss:0.254324
[9] train-logloss:0.236549
[10] train-logloss:0.217855
[11] train-logloss:0.204182
[12] train-logloss:0.190731
[13] train-logloss:0.180605
[14] train-logloss:0.169654
[15] train-logloss:0.158723
[16] train-logloss:0.150307
[17] train-logloss:0.142247
[18] train-logloss:0.135132
[19] train-logloss:0.128481
[20] train-logloss:0.123429
[21] train-logloss:0.119813
[22] train-logloss:0.115978
[23] train-logloss:0.112281
[24] train-logloss:0.109127
[25] train-logloss:0.106008
[26] train-logloss:0.102319
[27] train-logloss:0.098278
[28] train-logloss:0.096241
[29] train-logloss:0.093938
[30] train-logloss:0.090129
[31] train-logloss:0.088019
[32] train-logloss:0.086471
[33] train-logloss:0.084705
[34] train-logloss:0.083406
[35] train-logloss:0.081934
[36] train-logloss:0.080555
[37] train-logloss:0.079598
[38] train-logloss:0.077407

```



[39] train-logloss:0.076015  
[40] train-logloss:0.074463  
[41] train-logloss:0.073296  
[42] train-logloss:0.071826  
[43] train-logloss:0.070887  
[44] train-logloss:0.069892  
[45] train-logloss:0.068948  
[46] train-logloss:0.067955  
[47] train-logloss:0.067070  
[48] train-logloss:0.065803  
[49] train-logloss:0.065132  
[50] train-logloss:0.064404  
[51] train-logloss:0.063838  
[52] train-logloss:0.063135  
[53] train-logloss:0.062080  
[54] train-logloss:0.060893  
[55] train-logloss:0.060218  
[56] train-logloss:0.059754  
[57] train-logloss:0.059384  
[58] train-logloss:0.058700  
[59] train-logloss:0.058181  
[60] train-logloss:0.057617  
[61] train-logloss:0.057199  
[62] train-logloss:0.056499  
[63] train-logloss:0.056234  
[64] train-logloss:0.055910  
[65] train-logloss:0.055626  
[66] train-logloss:0.055248  
[67] train-logloss:0.054935  
[68] train-logloss:0.054247  
[69] train-logloss:0.053829  
[70] train-logloss:0.053391  
[71] train-logloss:0.053000  
[72] train-logloss:0.052614  
[73] train-logloss:0.052252  
[74] train-logloss:0.051871  
[75] train-logloss:0.051650  
[76] train-logloss:0.051369  
[77] train-logloss:0.050974  
[78] train-logloss:0.050654  
[79] train-logloss:0.050366  
[80] train-logloss:0.050112  
[81] train-logloss:0.049789  
[82] train-logloss:0.049541  
[83] train-logloss:0.049309  
[84] train-logloss:0.048993  
[85] train-logloss:0.048754  
[86] train-logloss:0.048456  
[87] train-logloss:0.048270  
[88] train-logloss:0.047914  
[89] train-logloss:0.047611  
[90] train-logloss:0.047347  
[91] train-logloss:0.047108  
[92] train-logloss:0.046852

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```

```

# Generar predicciones en el conjunto de prueba
predicciones <- predict(modelo_xgb, test_matrix)

# Convertir las probabilidades en clases (0 o 1) usando un umbral de 0.5
predicciones_clase <- ifelse(predicciones >= 0.5, 1, 0)

# Crear y mostrar la matriz de confusión comparando predicciones con etiquetas reales
matriz_confusion <- table(Predicho = predicciones_clase, Real = test_target)
print(matriz_confusion)

```

```

      Real
Predicho 0  1
      0 10  4
      1  6 28

```

```

# Utilizar caret para una matriz de confusión con métricas adicionales
confusion <- confusionMatrix(factor(predicciones_clase),
                              factor(test_target),
                              mode="everything",
                              positive = "1")

print(confusion)

```

Confusion Matrix and Statistics

	Reference	
Prediction	0	1
0	10	4
1	6	28

Accuracy : 0.7917  
 95% CI : (0.6501, 0.8953)  
 No Information Rate : 0.6667  
 P-Value [Acc > NIR] : 0.04225

Kappa : 0.5161

McNemar's Test P-Value : 0.75183

Sensitivity : 0.8750  
 Specificity : 0.6250  
 Pos Pred Value : 0.8235  
 Neg Pred Value : 0.7143  
 Precision : 0.8235  
 Recall : 0.8750  
 F1 : 0.8485  
 Prevalence : 0.6667  
 Detection Rate : 0.5833  
 Detection Prevalence : 0.7083  
 Balanced Accuracy : 0.7500

'Positive' Class : 1

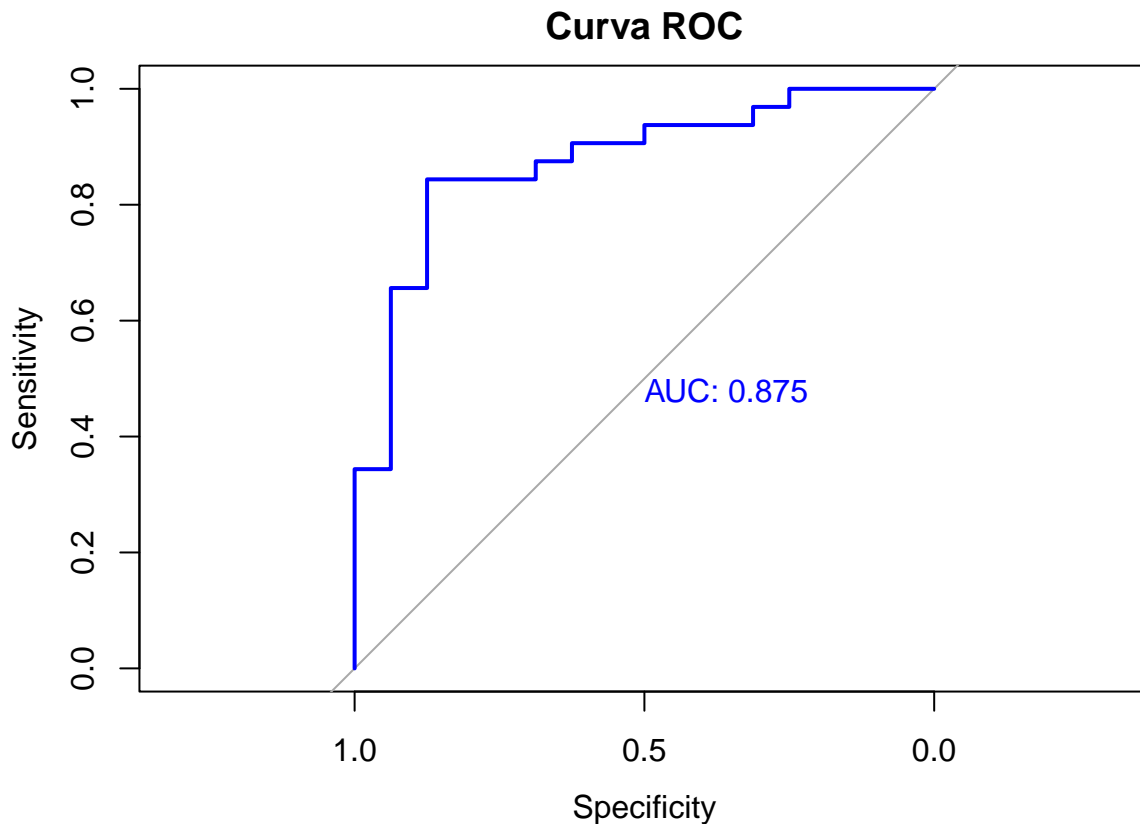
```

#predicciones <- predict(modelo_xgb, test_matrix)

# Crear la curva ROC
roc_obj <- roc(test_target, predicciones)

# Graficar la curva ROC
plot(roc_obj, col = "blue", main = "Curva ROC", print.auc = TRUE)

```



```
# Calcular el AUC
auc_valor <- auc(roc_obj)
print(paste("El AUC es:", auc_valor))

[1] "El AUC es: 0.875"

# Obtener la importancia de las variables
importancia <- xgb.importance(model = modelo_xgb, feature_names = colnames(train_matrix))

# Convertir la importancia en un data.frame
importancia_df <- as.data.frame(importancia)

# Opcional: ordenar las variables por importancia de mayor a menor
importancia_df <- importancia_df[order(-importancia_df$Gain), ]

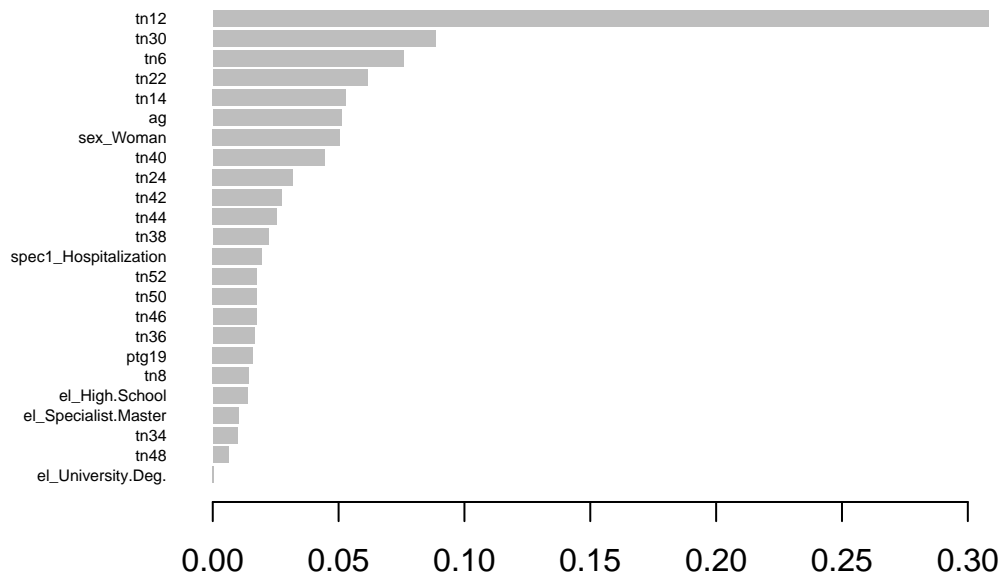
# Mostrar la importancia de las variables
print(importancia)
```

	Feature <char>	Gain <num>	Cover <num>	Frequency <num>
1:	tn12	0.3081574275	0.080903069	0.051162791
2:	tn30	0.0884876880	0.104197717	0.108139535
3:	tn6	0.0756919635	0.111375236	0.111627907
4:	tn22	0.0615674602	0.080054197	0.075581395
5:	tn14	0.0529809393	0.026760600	0.015116279
6:	ag	0.0512532636	0.064178246	0.072093023
7:	sex_Woman	0.0506338962	0.009351063	0.004651163
8:	tn40	0.0444053365	0.058155601	0.051162791

9:	tn24	0.0319256084	0.016915259	0.016279070
10:	tn42	0.0275703809	0.113162988	0.137209302
11:	tn44	0.0255161959	0.038014467	0.039534884
12:	tn38	0.0221177198	0.036401584	0.041860465
13:	spec1_Hospitalization	0.0196221524	0.012998049	0.003488372
14:	tn52	0.0176664100	0.028920006	0.024418605
15:	tn50	0.0174935906	0.012536896	0.011627907
16:	tn46	0.0174647243	0.022108332	0.029069767
17:	tn36	0.0165454101	0.036818706	0.038372093
18:	ptg19	0.0159604116	0.077150720	0.102325581
19:	tn8	0.0144680297	0.018145517	0.020930233
20:	el_High.School	0.0137545420	0.022421510	0.015116279
21:	el_Specialist.Master	0.0101955625	0.005362287	0.003488372
22:	tn34	0.0097462746	0.012136458	0.013953488
23:	tn48	0.0062923955	0.009682384	0.010465116
24:	el_University.Deg.	0.0004826168	0.002249108	0.002325581
	Feature	Gain	Cover	Frequency

*# Graficar la importancia de las variables*

```
xgb.plot.importance(importance_matrix = importancia)
```



Para darle mayor explicabilidad al modelo vamos a calcular los valores SHAP de observaciones individuales

*# Definir la función de predicción personalizada para el modelo xgboost*

```
pred_fun <- function(object, newdata) {
  predict(object, newdata = xgb.DMatrix(data = as.matrix(newdata)))
}
```

*# Convertir test\_data a data.frame y seleccionar una observación para explicar*

```
test_df <- as.data.frame(as.matrix(test_data))
new_observation <- test_df[2, , drop = FALSE] # Seleccionar la primera observación
```

*# Calcular los valores SHAP para la observación seleccionada*

```
set.seed(123) # Para reproducibilidad
shap_values_fast <- fastshap::explain(
  object = modelo_xgb, # El modelo de xgboost
  feature_names = colnames(train_data), # Los nombres de las variables
```

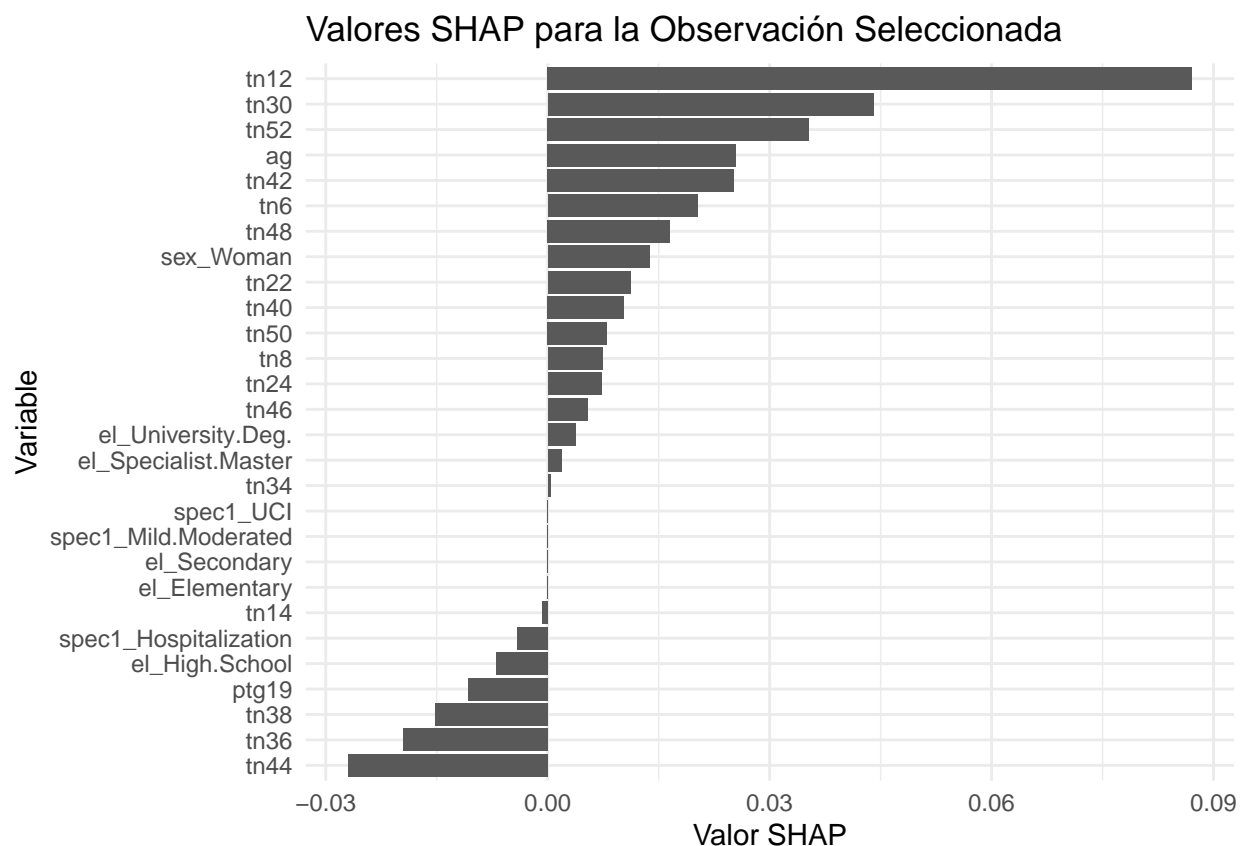
```

newdata = new_observation,      # La observación para la cual queremos valores SHAP
pred_wrapper = pred_fun,       # La función de predicción personalizada
X = train_data,                # El conjunto de entrenamiento completo
nsim = 100                     # Número de simulaciones
)

# Convertir los valores SHAP a un data.frame para graficar
shap_df <- data.frame(
  Variable = colnames(new_observation),
  SHAP = shap_values_fast[1, ] # Solo la primera fila (si tienes más observaciones)
)

# Graficar los valores SHAP
ggplot(shap_df, aes(x = reorder(Variable, SHAP), y = SHAP)) +
  geom_bar(stat = "identity") +
  coord_flip() +
  labs(title = "Valores SHAP para la Observación Seleccionada",
       x = "Variable",
       y = "Valor SHAP") +
  theme_minimal()

```



##### VARIOS VALORES #####

```

# Convertir test_data a data.frame y seleccionar múltiples observaciones para explicar (por ejemplo, la
test_df <- as.data.frame(as.matrix(test_data))
new_observations <- test_df[1:48, , drop = FALSE] # Seleccionar las primeras 10 observaciones

```

```

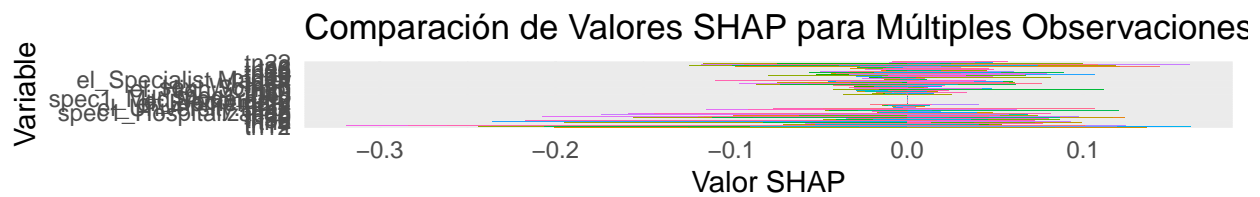
# Calcular los valores SHAP para las observaciones seleccionadas
set.seed(123) # Para reproducibilidad
shap_values_fast_24 <- fastshap::explain(
  object = modelo_xgb,          # El modelo de xgboost
  feature_names = colnames(train_data), # Los nombres de las variables
  newdata = new_observations,    # Las observaciones seleccionadas
  pred_wrapper = pred_fun,      # La función de predicción personalizada
  X = train_data,              # El conjunto de entrenamiento completo
  nsim = 100                    # Número de simulaciones
)

# Convertir shap_values_fast a data.frame y añadir un identificador de observación en el proceso de piv
shap_df_24 <- as.data.frame(shap_values_fast_24)
colnames(shap_df_24) <- colnames(train_data) # Asegurarse de que los nombres de columnas sean los nomb
shap_df_24$Observation <- factor(1:48) # Añadir la identificación de observación como factor

# Convertir a formato largo para visualización
shap_df_long <- shap_df_24 %>%
  pivot_longer(cols = -Observation, names_to = "Variable", values_to = "SHAP")

# Graficar los valores SHAP para múltiples observaciones
ggplot(shap_df_long, aes(x = reorder(Variable, SHAP), y = SHAP, fill = Observation)) +
  geom_bar(stat = "identity", position = "dodge") +
  coord_flip() +
  labs(title = "Comparación de Valores SHAP para Múltiples Observaciones",
       x = "Variable",
       y = "Valor SHAP") +
  theme_minimal() +
  theme(legend.position = "bottom")

```

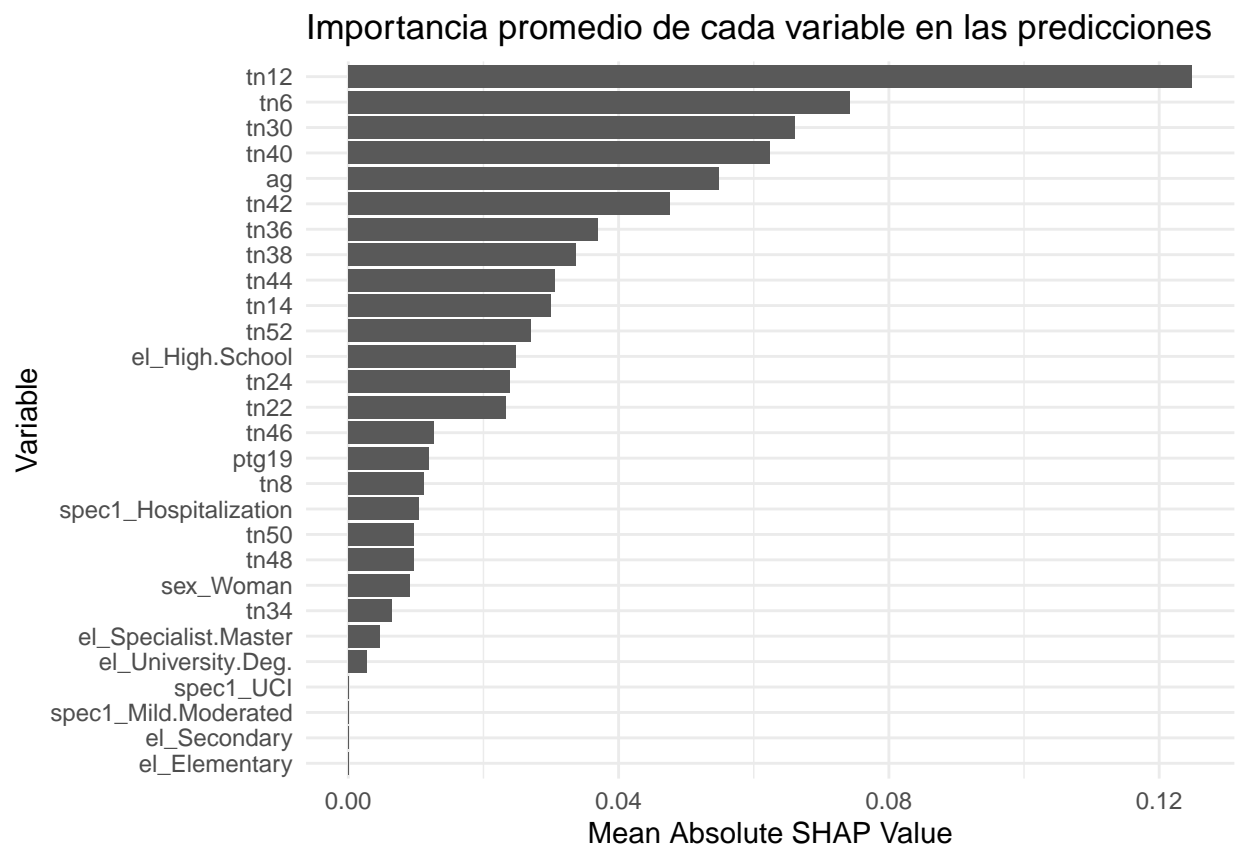


Observation	1	11	21	31	41
	2	12	22	32	42
	3	13	23	33	43
	4	14	24	34	44
	5	15	25	35	45
	6	16	26	36	46
	7	17	27	37	47
	8	18	28	38	48
	9	19	29	39	
	10	20	30	40	

```
##### COMPARAR TODOS LOS VALORES SHAP #####

# Calcular la importancia promedio absoluta de cada variable
importance_summary <- shap_df_24 %>%
  dplyr::select(-Observation) %>% # Excluir la columna de observación
  summarise_all(~ mean(abs(.))) %>% # Calcular la media absoluta para cada variable
  pivot_longer(cols = everything(), names_to = "Variable", values_to = "Mean_Absolute_SHAP") %>%
  arrange(desc(Mean_Absolute_SHAP)) # Ordenar de mayor a menor importancia

# Graficar la importancia promedio de cada variable
ggplot(importance_summary, aes(x = reorder(Variable, Mean_Absolute_SHAP), y = Mean_Absolute_SHAP)) +
  geom_bar(stat = "identity") +
  coord_flip() +
  labs(title = "Importancia promedio de cada variable en las predicciones",
       x = "Variable",
       y = "Mean Absolute SHAP Value") +
  theme_minimal()
```



```
# Mostrar el data.frame con la importancia promedio de cada variable
print(importance_summary)
```

```
# A tibble: 28 x 2
  Variable Mean_Absolute_SHAP
  <chr>      <dbl>
1 tn12      0.125
2 tn6       0.0742
3 tn30      0.0661
4 tn40      0.0623
5 ag        0.0548
6 tn42      0.0475
7 tn36      0.0369
8 tn38      0.0337
9 tn44      0.0306
10 tn14     0.0299
# i 18 more rows
```