

Tratamiento, Limpieza y Selección de variables

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1. TRATAMIENTO

En este primer paso, cargaremos los datos y aplicaremos un tratamiento previo, incluyendo la combinación de variables duplicadas y la renombración de columnas para mejorar su comprensión.

1.1 Carga de librerías

```
library(readxl)
library(dplyr)
```

```
##
## Attaching package: 'dplyr'

## The following objects are masked from 'package:stats':
##
##   filter, lag

## The following objects are masked from 'package:base':
##
##   intersect, setdiff, setequal, union
```

```
library(ggplot2)
library(mice)
```

```
##
## Attaching package: 'mice'

## The following object is masked from 'package:stats':
##
##   filter

## The following objects are masked from 'package:base':
##
##   cbind, rbind
```

```
library(missForest)
```

```
## Warning: package 'missForest' was built under R version 4.3.3
```

1.2 Carga de datos

```
# Reemplaza "datos.xlsx" y "hoja1" con el nombre real de tu archivo y hoja  
datos <- read_excel("/Users/manuelrocamoravalenti/Desktop/TFG/Datos_Crudo/Base_Pembro_1L(febrero_24)_v2
```

```
## New names:  
## * 'Estudios' -> 'Estudios...14'  
## * 'Est_civil' -> 'Est_civil...15'  
## * 'Comp_hogar' -> 'Comp_hogar...16'  
## * 'MOSs' -> 'MOSs...17'  
## * 'Ansiedad' -> 'Ansiedad...18'  
## * 'Depresion' -> 'Depresion...19'  
## * 'MNA' -> 'MNA...20'  
## * 'MNA' -> 'MNA...27'  
## * 'Estudios' -> 'Estudios...29'  
## * 'Est_civil' -> 'Est_civil...31'  
## * 'Comp_hogar' -> 'Comp_hogar...32'  
## * 'MOSs' -> 'MOSs...33'  
## * 'Ansiedad' -> 'Ansiedad...36'  
## * 'Depresion' -> 'Depresion...37'
```

1.3 Cambios de algunos nombres

```
names(datos)[names(datos) == "Joven(0)_Anciano(1)"] <- "Anciano"  
names(datos)[names(datos) == "%_perd_peso"] <- "Porcentaje_perdpeso"  
names(datos)[names(datos) == "PD-L1"] <- "PD_L1"  
names(datos)[names(datos) == "1ª_eval"] <- "primera_eval"  
names(datos)[names(datos) == "1ªeval_num"] <- "primera_eval_num"  
names(datos)[names(datos) == "Toxicidad_si/no"] <- "Toxicidad"  
names(datos)[names(datos) == "Progresión_sí/no"] <- "Progresion"  
names(datos)[names(datos) == "2ªL_sí/no"] <- "segunda_eval"  
names(datos)[names(datos) == "Exitus_sí/no"] <- "Exitus"  
names(datos)[names(datos) == "T"] <- "Tamaño_tumor"  
names(datos)[names(datos) == "N"] <- "Afectacion_ganglionar"  
names(datos)[names(datos) == "M"] <- "Afectacion_metastasica"
```

1.4 Combinacion de variables repetidas

```
datos <- datos %>%  
  mutate(estudios = coalesce(`Estudios...14`, `Estudios...29`)) %>%  
  select(-`Estudios...14`, -`Estudios...29`)
```

```

datos <- datos %>%
  mutate(est_civil = coalesce(`Est_civil...15`, `Est_civil...31`)) %>%
  select(-`Est_civil...15`, -`Est_civil...31`)

datos <- datos %>%
  mutate(hogar = coalesce(`Comp_hogar...16`, `Comp_hogar...32`)) %>%
  select(-`Comp_hogar...16`, -`Comp_hogar...32`)

datos <- datos %>%
  mutate(MOSSs = coalesce(`MOSs...17`, `MOSs...33`)) %>%
  select(-`MOSs...17`, -`MOSs...33`)

datos <- datos %>%
  mutate(ansiedad = coalesce(`Ansiedad...18`, `Ansiedad...36`)) %>%
  select(-`Ansiedad...18`, -`Ansiedad...36`)

datos <- datos %>%
  mutate(depresion = coalesce(`Depresion...19`, `Depresion...37`)) %>%
  select(-`Depresion...19`, -`Depresion...37`)

datos <- datos %>%
  mutate(MNA = coalesce(`MNA...20`, `MNA...27`)) %>%
  select(-`MNA...20`, -`MNA...27`)

```

2. LIMPIEZA

Para garantizar la calidad del análisis, es necesario limpiar el conjunto de datos eliminando aquellas columnas con un alto porcentaje de valores faltantes.

En este caso, estableceremos un **umbral del 20%**, por lo que cualquier columna que supere este porcentaje será eliminada. Esto nos permite reducir el impacto de datos incompletos y mejorar la fiabilidad de los resultados obtenidos.

2.1 Eliminación de columnas previa

```

datos <- datos %>% select(-Num_pac, -Fecha_nac, -Fecha_dx, -Peso, -Talla,
  -G8, -Audicion, -Barthel, -Lawton_Brody, -SPPB,
  -Caida_6m, -Pfeiffer, -Mini_mental, -Social_Gijon,
  -Yesavage, -CIRS, -Charlson, -Polifarmacia, -Sd_geriatr,
  -Clasif_geriatr_Balducci, -Clasif_geriatr_SIOG1,
  -Biopsia_liq, -Tipo_mut_Liq, -Fecha_inicio_pem,
  -primera_eval, -Mejor_resp, -Fecha_exitus, -Fecha_últ_control,
  -Fecha_SLP, -Fecha_SG, -Fecha_progresión, -Observaciones, -'Leuc%', -'Linf%',
  -'CD45+')

```

2.2 Análisis de datos faltantes (Columnas)

Ahora vamos a calcular el porcentaje de valores faltantes por columna, crear un resumen con los resultados y ordenarlos de mayor a menor según su porcentaje.

```

# Calcular el porcentaje de valores faltantes por columna
missing_percent <- colSums(is.na(datos)) / nrow(datos) * 100

# Crear un dataframe con el resumen de valores faltantes
missing_summary <- data.frame(Columna = names(missing_percent), Porcentaje_Faltante = missing_percent)

# Mostrar el resumen
dplyr::arrange(missing_summary, desc(Porcentaje_Faltante))

```

##	Columna	Porcentaje_Faltante
##	CD3_C_Leuc	CD3_C_Leuc 100.000000
##	CD4+_C_CD45	CD4+_C_CD45 100.000000
##	CD45+_C	CD45+_C 100.000000
##	CD127-/lowFoxP3+_%CD4	CD127-/lowFoxP3+_%CD4 88.888889
##	CD25+FoxP3+_%Linf	CD25+FoxP3+_%Linf 88.888889
##	CD39+FoxP3+_%CD4	CD39+FoxP3+_%CD4 88.888889
##	CD4+_%CD45	CD4+_%CD45 88.888889
##	CD45RA+_%Linf	CD45RA+_%Linf 88.888889
##	CD45RA+FoxP3+_%CD4	CD45RA+FoxP3+_%CD4 88.888889
##	Helios+FoxP3+_%CD4	Helios+FoxP3+_%CD4 88.888889
##	CD127-/lowFoxP3+_C_CD4	CD127-/lowFoxP3+_C_CD4 88.888889
##	CD25+FoxP3+_C_Linf	CD25+FoxP3+_C_Linf 88.888889
##	CD39+FoxP3+_C_CD4	CD39+FoxP3+_C_CD4 88.888889
##	CD45RA+_C_Linf	CD45RA+_C_Linf 88.888889
##	CD45RA+FoxP3+_C_CD4	CD45RA+FoxP3+_C_CD4 88.888889
##	Helios+FoxP3+_C_CD4	Helios+FoxP3+_C_CD4 88.888889
##	CD25+CD127low_%Linf	CD25+CD127low_%Linf 72.222222
##	CD25+CD127low_C_Linf	CD25+CD127low_C_Linf 72.222222
##	HLADR+Lin_%Leuc	HLADR+Lin_%Leuc 63.888889
##	mDC_%Leuc	mDC_%Leuc 63.888889
##	pDC_%Leuc	pDC_%Leuc 63.888889
##	CD4_Central_Mem_%Linf	CD4_Central_Mem_%Linf 63.888889
##	CD4_Effector_Mem_%Linf	CD4_Effector_Mem_%Linf 63.888889
##	CD4_Naïve_%Linf	CD4_Naïve_%Linf 63.888889
##	CD4_TEMRA_%Linf	CD4_TEMRA_%Linf 63.888889
##	CD8_Central_Mem_%Linf	CD8_Central_Mem_%Linf 63.888889
##	CD8_Effector_Mem_%Linf	CD8_Effector_Mem_%Linf 63.888889
##	CD8_Naïve_%Linf	CD8_Naïve_%Linf 63.888889
##	CD8_TEMRA_%Linf	CD8_TEMRA_%Linf 63.888889
##	mDC_CD16_%mDC	mDC_CD16_%mDC 63.888889
##	mDC_CD1c_%mDC	mDC_CD1c_%mDC 63.888889
##	mDC_Clec9A_%mDC	mDC_Clec9A_%mDC 63.888889
##	CD3+_%Linf	CD3+_%Linf 63.888889
##	CD27-CD57+CD3+_%Linf	CD27-CD57+CD3+_%Linf 63.888889
##	CD27-CD57+CD4+_%CD3	CD27-CD57+CD4+_%CD3 63.888889
##	CD27-CD57+CD8+_%CD3	CD27-CD57+CD8+_%CD3 63.888889
##	CD3_%Leuc	CD3_%Leuc 63.888889
##	CD3+CD4+_%Linf	CD3+CD4+_%Linf 63.888889
##	CD3+CD57+_%Linf	CD3+CD57+_%Linf 63.888889
##	CD3+CD8+_%Linf	CD3+CD8+_%Linf 63.888889
##	CD45RA+CCR7+CD3+_%Linf	CD45RA+CCR7+CD3+_%Linf 63.888889
##	CD8+_term_efect_%CD3	CD8+_term_efect_%CD3 63.888889
##	CD8_exhausted_%CD3	CD8_exhausted_%CD3 63.888889

## CD4_TCR_ab+_%CD3	CD4_TCR_ab+_%CD3	63.888889
## CD4+_%Linf	CD4+_%Linf	63.888889
## CD8+_TCR_ab+_%CD3	CD8+_TCR_ab+_%CD3	63.888889
## CD8+_%Linf	CD8+_%Linf	63.888889
## CD8+CD4+_%Linf	CD8+CD4+_%Linf	63.888889
## CD8-CD4-_%Linf	CD8-CD4-_%Linf	63.888889
## HLADR+CD3+_%Linf	HLADR+CD3+_%Linf	63.888889
## TCR_ab+_%Linf	TCR_ab+_%Linf	63.888889
## TCR_gd+_%Linf	TCR_gd+_%Linf	63.888889
## gd_VD1+_%CD3	gd_VD1+_%CD3	63.888889
## gd_VD1+VD2+_%CD3	gd_VD1+VD2+_%CD3	63.888889
## gd_VD1-VD2-_%CD3	gd_VD1-VD2-_%CD3	63.888889
## gd_VD2+_%CD3	gd_VD2+_%CD3	63.888889
## LDH_PE	LDH_PE	61.111111
## CD25+CD4+_%Linf	CD25+CD4+_%Linf	61.111111
## HLADR+Lin_C_Leuc	HLADR+Lin_C_Leuc	61.111111
## mDC_C_Leuc	mDC_C_Leuc	61.111111
## pDC_C_Leuc	pDC_C_Leuc	61.111111
## CD4_Central_Mem_C_Linf	CD4_Central_Mem_C_Linf	61.111111
## CD4_Effector_Mem_C_Linf	CD4_Effector_Mem_C_Linf	61.111111
## CD4_Naïve_C_Linf	CD4_Naïve_C_Linf	61.111111
## CD4_TEMRA_C_Linf	CD4_TEMRA_C_Linf	61.111111
## CD8_Central_Mem_C_Linf	CD8_Central_Mem_C_Linf	61.111111
## CD8_Effector_Mem_C_Linf	CD8_Effector_Mem_C_Linf	61.111111
## CD8_Naïve_C_Linf	CD8_Naïve_C_Linf	61.111111
## CD8_TEMRA_C_Linf	CD8_TEMRA_C_Linf	61.111111
## mDC_CD16_C_mDC	mDC_CD16_C_mDC	61.111111
## mDC_CD1c_C_mDC	mDC_CD1c_C_mDC	61.111111
## mDC_Clec9A_C_mDC	mDC_Clec9A_C_mDC	61.111111
## CD3+_C_Linf	CD3+_C_Linf	61.111111
## CD27-CD57+CD3+_C_Linf	CD27-CD57+CD3+_C_Linf	61.111111
## CD27-CD57+CD4+_C_CD3	CD27-CD57+CD4+_C_CD3	61.111111
## CD27-CD57+CD8+_C_CD3	CD27-CD57+CD8+_C_CD3	61.111111
## CD3+CD4+_C_Linf	CD3+CD4+_C_Linf	61.111111
## CD3+CD57+_C_Linf	CD3+CD57+_C_Linf	61.111111
## CD3+CD8+_C_Linf	CD3+CD8+_C_Linf	61.111111
## CD45RA+CCR7+CD3+_C_Linf	CD45RA+CCR7+CD3+_C_Linf	61.111111
## CD8+_term_efect_C_CD3	CD8+_term_efect_C_CD3	61.111111
## CD8_exhausted_C_CD3	CD8_exhausted_C_CD3	61.111111
## CD4_TCR_ab+_C_CD3	CD4_TCR_ab+_C_CD3	61.111111
## CD4+_C_Linf	CD4+_C_Linf	61.111111
## CD8+_TCR_ab+_C_CD3	CD8+_TCR_ab+_C_CD3	61.111111
## CD8+_C_Linf	CD8+_C_Linf	61.111111
## CD8+CD4+_C_Linf	CD8+CD4+_C_Linf	61.111111
## CD8-CD4-_C_Linf	CD8-CD4-_C_Linf	61.111111
## HLADR+CD3+_C_Linf	HLADR+CD3+_C_Linf	61.111111
## TCR_ab+_C_Linf	TCR_ab+_C_Linf	61.111111
## TCR_gd+_C_Linf	TCR_gd+_C_Linf	61.111111
## gd_VD1+_C_CD3	gd_VD1+_C_CD3	61.111111
## gd_VD1+VD2+_C_CD3	gd_VD1+VD2+_C_CD3	61.111111
## gd_VD1-VD2-_C_CD3	gd_VD1-VD2-_C_CD3	61.111111
## gd_VD2+_C_CD3	gd_VD2+_C_CD3	61.111111
## CD25+CD4+_C_Linf	CD25+CD4+_C_Linf	61.111111
## MOOSs	MOOSs	61.111111

## IL-6	IL-6	58.333333
## IgM_CMV	IgM_CMV	55.555556
## IgG_CMV	IgG_CMV	55.555556
## LDH_2C	LDH_2C	50.000000
## Tipo_mut_Tej	Tipo_mut_Tej	47.222222
## Col_LDL	Col_LDL	47.222222
## MNA	MNA	47.222222
## Prot_PE	Prot_PE	44.444444
## ansiedad	ansiedad	44.444444
## depresion	depresion	44.444444
## Col_HDL	Col_HDL	41.666667
## Alb_PE	Alb_PE	41.666667
## Hb_PE	Hb_PE	41.666667
## Leucoc_PE	Leucoc_PE	41.666667
## Neutr_PE	Neutr_PE	41.666667
## Linf_PE	Linf_PE	41.666667
## Plaq_PE	Plaq_PE	41.666667
## NLR_PE	NLR_PE	41.666667
## NLRPE_corte 4	NLRPE_corte 4	41.666667
## NLRPE_corte 5	NLRPE_corte 5	41.666667
## PLR_PE	PLR_PE	41.666667
## PNI_PE	PNI_PE	41.666667
## SII_PE	SII_PE	41.666667
## LDH_1C	LDH_1C	36.111111
## LDH_1eval	LDH_1eval	33.333333
## estudios	estudios	27.777778
## est_civil	est_civil	27.777778
## LinfT_cel	LinfT_cel	25.000000
## LinfT_%	LinfT_%	25.000000
## CD4_cel	CD4_cel	25.000000
## CD4_%	CD4_%	25.000000
## CD8_cel	CD8_cel	25.000000
## CD8_%	CD8_%	25.000000
## CD4:CD8	CD4:CD8	25.000000
## LinfB_cel	LinfB_cel	25.000000
## LinfB_%	LinfB_%	25.000000
## LinfNK_cel	LinfNK_cel	25.000000
## LinfNK_%	LinfNK_%	25.000000
## hogar	hogar	25.000000
## PCR	PCR	22.222222
## segunda_eval	segunda_eval	19.444444
## Estado_mut	Estado_mut	16.666667
## Prot_2C	Prot_2C	16.666667
## Alb_2C	Alb_2C	13.888889
## PNI_2C	PNI_2C	13.888889
## Prot_1eval	Prot_1eval	13.888889
## Porcentaje_perdpeso	Porcentaje_perdpeso	11.111111
## p_peso_no_sí	p_peso_no_sí	11.111111
## Prot_1C	Prot_1C	11.111111
## Hb_2C	Hb_2C	11.111111
## Leucoc_2C	Leucoc_2C	11.111111
## Neutr_2C	Neutr_2C	11.111111
## Linf_2C	Linf_2C	11.111111
## Plaq_2C	Plaq_2C	11.111111

## NLR_2C	NLR_2C	11.111111
## NLR2C_corte4o5	NLR2C_corte4o5	11.111111
## PLR_2C	PLR_2C	11.111111
## SII_2C	SII_2C	11.111111
## Tipo_tox	Tipo_tox	11.111111
## LDH	LDH	8.333333
## Alb_1C	Alb_1C	8.333333
## Hab_tabaq	Hab_tabaq	5.555556
## Exp_tab	Exp_tab	5.555556
## NLR1C_corte5	NLR1C_corte5	5.555556
## PNI_1C	PNI_1C	5.555556
## primera_eval_num	primera_eval_num	5.555556
## Alb_1eval	Alb_1eval	5.555556
## Hb_1eval	Hb_1eval	5.555556
## Leucoc_1eval	Leucoc_1eval	5.555556
## Neutr_1eval	Neutr_1eval	5.555556
## Linf_1eval	Linf_1eval	5.555556
## Plaq_1eval	Plaq_1eval	5.555556
## NLR_1eval	NLR_1eval	5.555556
## PLR_1eval	PLR_1eval	5.555556
## PNI_1eval	PNI_1eval	5.555556
## SII_1eval	SII_1eval	5.555556
## Mejor_resp_num	Mejor_resp_num	5.555556
## Tamaño_tumor	Tamaño_tumor	2.777778
## Afectacion_ganglionar	Afectacion_ganglionar	2.777778
## Afectacion_metastasica	Afectacion_metastasica	2.777778
## Col_total	Col_total	2.777778
## Prot_tot	Prot_tot	2.777778
## Albumina	Albumina	2.777778
## PNI_pre	PNI_pre	2.777778
## ALI_pre	ALI_pre	2.777778
## Hb_1C	Hb_1C	2.777778
## Leucoc_1C	Leucoc_1C	2.777778
## Neutr_1C	Neutr_1C	2.777778
## Linf_1C	Linf_1C	2.777778
## Plaq_1C	Plaq_1C	2.777778
## NLR_1C	NLR_1C	2.777778
## NLR1C_corte4	NLR1C_corte4	2.777778
## PLR_1C	PLR_1C	2.777778
## SII_1C	SII_1C	2.777778
## Idpac	Idpac	0.000000
## Sexo	Sexo	0.000000
## Edad_dx	Edad_dx	0.000000
## Anciano	Anciano	0.000000
## ECOG	ECOG	0.000000
## IMC	IMC	0.000000
## Diabetes	Diabetes	0.000000
## Cardiop	Cardiop	0.000000
## Enf_neurod	Enf_neurod	0.000000
## Histologia	Histologia	0.000000
## Histología_num	Histología_num	0.000000
## Estadio	Estadio	0.000000
## Estadio_num	Estadio_num	0.000000
## PD_L1	PD_L1	0.000000

```
## Estatinas          Estatinas          0.000000
## Hb                  Hb                  0.000000
## Leucoc_tot          Leucoc_tot          0.000000
## Neutrofilos         Neutrofilos         0.000000
## Linf_tot            Linf_tot            0.000000
## Plaquetas           Plaquetas           0.000000
## NLR_pre             NLR_pre             0.000000
## PLR_pre             PLR_pre             0.000000
## SII_pre             SII_pre             0.000000
## N_ciclos            N_ciclos            0.000000
## Toxicidad           Toxicidad           0.000000
## Grado_tox           Grado_tox           0.000000
## Interrupc_tto       Interrupc_tto       0.000000
## Motivo_inter        Motivo_inter        0.000000
## Progresion          Progresion          0.000000
## Exitus              Exitus              0.000000
## SLP                 SLP                 0.000000
## SLP_cens            SLP_cens            0.000000
## SG                  SG                  0.000000
## SG_cens             SG_cens             0.000000
```

```
# Crear un dataframe con los porcentajes de valores faltantes por columna
df_missing_col <- data.frame(Porcentaje_Faltante = missing_percent)
```

```
# Generar el histograma
```

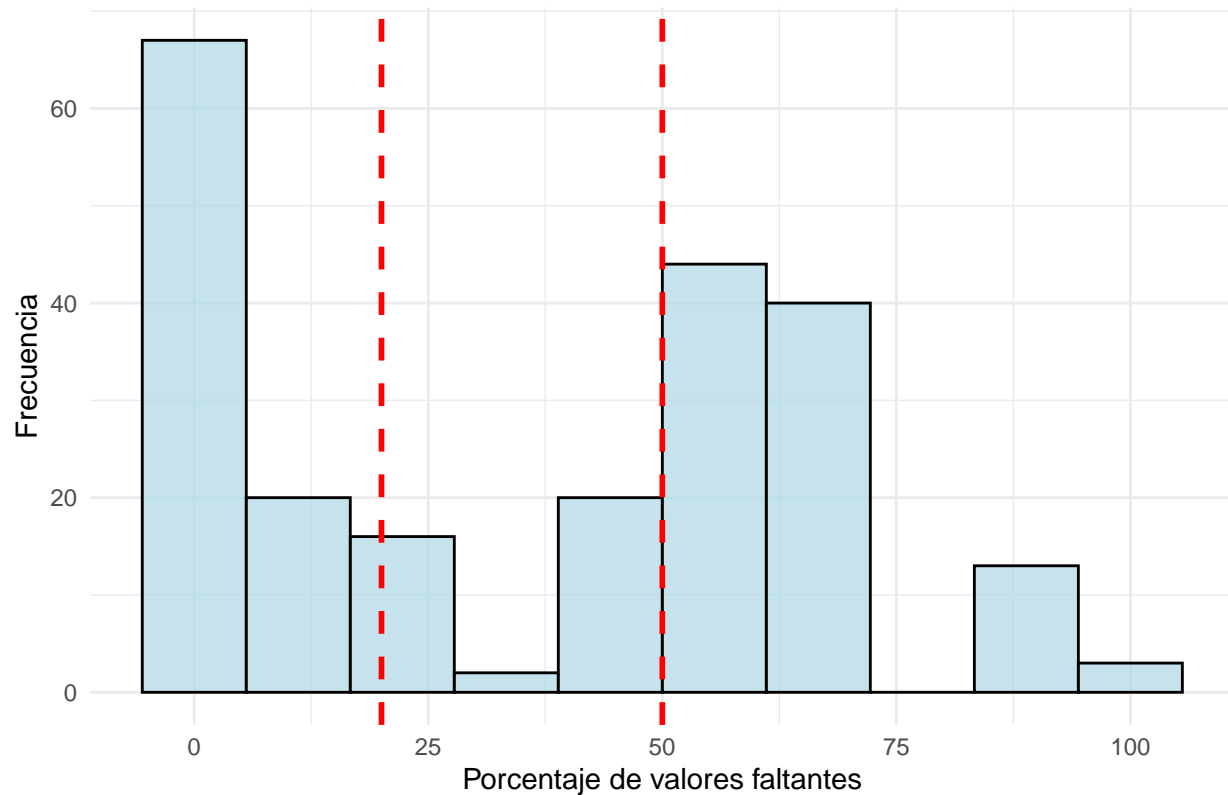
```
ggplot(df_missing_col, aes(x = Porcentaje_Faltante)) +
  geom_histogram(bins = 10, fill = "lightblue", color = "black", alpha = 0.7) +
  geom_vline(xintercept = 20, color = "red", linetype = "dashed", size = 1, label = "Umbral 20%") +
  geom_vline(xintercept = 50, color = "red", linetype = "dashed", size = 1, label = "Umbral 50%") +
  labs(title = "Distribución de valores faltantes por columna",
       x = "Porcentaje de valores faltantes",
       y = "Frecuencia") +
  theme_minimal()
```

```
## Warning: Using 'size' aesthetic for lines was deprecated in ggplot2 3.4.0.
## i Please use 'linewidth' instead.
## This warning is displayed once every 8 hours.
## Call 'lifecycle::last_lifecycle_warnings()' to see where this warning was
## generated.
```

```
## Warning in geom_vline(xintercept = 20, color = "red", linetype = "dashed", :
## Ignoring unknown parameters: 'label'
```

```
## Warning in geom_vline(xintercept = 50, color = "red", linetype = "dashed", :
## Ignoring unknown parameters: 'label'
```


Distribución de valores faltantes por columna



2.2.1 Eliminación de columnas con más del 20% de valores faltantes

```
threshold <- 20 # Umbral de eliminación
datos_limpios <- datos %>% select(which(missing_percent <= threshold))

# Verificar las columnas eliminadas
columnas_eliminadas <- names(missing_percent[missing_percent > threshold])
columnas_eliminadas
```

```
## [1] "Tipo_mut_Tej"      "Col_HDL"
## [3] "Col_LDL"           "PCR"
## [5] "IL-6"              "LinfT_cel"
## [7] "LinfT_%"           "CD4_cel"
## [9] "CD4_%"             "CD8_cel"
## [11] "CD8_%"             "CD4:CD8"
## [13] "LinfB_cel"         "LinfB_%"
## [15] "LinfNK_cel"        "LinfNK_%"
## [17] "IgM_CMV"           "IgG_CMV"
## [19] "LDH_1C"            "LDH_2C"
## [21] "LDH_1eval"         "LDH_PE"
## [23] "Prot_PE"           "Alb_PE"
## [25] "Hb_PE"             "Leucoc_PE"
## [27] "Neutr_PE"          "Linf_PE"
## [29] "Plaq_PE"           "NLR_PE"
```

## [31]	"NLRPE_corte 4"	"NLRPE_corte 5"
## [33]	"PLR_PE"	"PNI_PE"
## [35]	"SII_PE"	"HLADR+Lin_%Leuc"
## [37]	"mDC_%Leuc"	"pDC_%Leuc"
## [39]	"CD4_Central_Mem_%Linf"	"CD4_Effector_Mem_%Linf"
## [41]	"CD4_Naïve_%Linf"	"CD4_TEMRA_%Linf"
## [43]	"CD8_Central_Mem_%Linf"	"CD8_Effector_Mem_%Linf"
## [45]	"CD8_Naïve_%Linf"	"CD8_TEMRA_%Linf"
## [47]	"mDC_CD16_%mDC"	"mDC_CD1c_%mDC"
## [49]	"mDC_Clec9A_%mDC"	"CD3+_%Linf"
## [51]	"CD27-CD57+CD3+_%Linf"	"CD27-CD57+CD4+_%CD3"
## [53]	"CD27-CD57+CD8+_%CD3"	"CD3_%Leuc"
## [55]	"CD3+CD4+_%Linf"	"CD3+CD57+_%Linf"
## [57]	"CD3+CD8+_%Linf"	"CD45RA+CCR7+CD3+_%Linf"
## [59]	"CD8+_term_efect_%CD3"	"CD8_exhausted_%CD3"
## [61]	"CD4_TCR_ab+_%CD3"	"CD4+_%Linf"
## [63]	"CD8+_TCR_ab+_%CD3"	"CD8+_%Linf"
## [65]	"CD8+CD4+_%Linf"	"CD8-CD4-_%Linf"
## [67]	"HLADR+CD3+_%Linf"	"TCR_ab+_%Linf"
## [69]	"TCR_gd+_%Linf"	"gd_VD1+_%CD3"
## [71]	"gd_VD1+VD2+_%CD3"	"gd_VD1-VD2-_%CD3"
## [73]	"gd_VD2+_%CD3"	"CD127-/lowFoxP3+_%CD4"
## [75]	"CD25+CD127low_%Linf"	"CD25+CD4+_%Linf"
## [77]	"CD25+FoxP3+_%Linf"	"CD39+FoxP3+_%CD4"
## [79]	"CD4+_%CD45"	"CD45RA+_%Linf"
## [81]	"CD45RA+FoxP3+_%CD4"	"Helios+FoxP3+_%CD4"
## [83]	"HLADR+Lin_C_Leuc"	"mDC_C_Leuc"
## [85]	"pDC_C_Leuc"	"CD4_Central_Mem_C_Linf"
## [87]	"CD4_Effector_Mem_C_Linf"	"CD4_Naïve_C_Linf"
## [89]	"CD4_TEMRA_C_Linf"	"CD8_Central_Mem_C_Linf"
## [91]	"CD8_Effector_Mem_C_Linf"	"CD8_Naïve_C_Linf"
## [93]	"CD8_TEMRA_C_Linf"	"mDC_CD16_C_mDC"
## [95]	"mDC_CD1c_C_mDC"	"mDC_Clec9A_C_mDC"
## [97]	"CD3+_C_Linf"	"CD27-CD57+CD3+_C_Linf"
## [99]	"CD27-CD57+CD4+_C_CD3"	"CD27-CD57+CD8+_C_CD3"
## [101]	"CD3_C_Leuc"	"CD3+CD4+_C_Linf"
## [103]	"CD3+CD57+_C_Linf"	"CD3+CD8+_C_Linf"
## [105]	"CD45RA+CCR7+CD3+_C_Linf"	"CD8+_term_efect_C_CD3"
## [107]	"CD8_exhausted_C_CD3"	"CD4_TCR_ab+_C_CD3"
## [109]	"CD4+_C_Linf"	"CD8+_TCR_ab+_C_CD3"
## [111]	"CD8+_C_Linf"	"CD8+CD4+_C_Linf"
## [113]	"CD8-CD4-_C_Linf"	"HLADR+CD3+_C_Linf"
## [115]	"TCR_ab+_C_Linf"	"TCR_gd+_C_Linf"
## [117]	"gd_VD1+_C_CD3"	"gd_VD1+VD2+_C_CD3"
## [119]	"gd_VD1-VD2-_C_CD3"	"gd_VD2+_C_CD3"
## [121]	"CD127-/lowFoxP3+_C_CD4"	"CD25+CD127low_C_Linf"
## [123]	"CD25+CD4+_C_Linf"	"CD25+FoxP3+_C_Linf"
## [125]	"CD39+FoxP3+_C_CD4"	"CD4+_C_CD45"
## [127]	"CD45+_C"	"CD45RA+_C_Linf"
## [129]	"CD45RA+FoxP3+_C_CD4"	"Helios+FoxP3+_C_CD4"
## [131]	"estudios"	"est_civil"
## [133]	"hogar"	"MOOSs"
## [135]	"ansiedad"	"depresion"
## [137]	"MNA"	

Por lo que las columnas que quedan **dentro del analisis** son las siguientes:

```
colnames(datos_limpios)
```

```
## [1] "Idpac"          "Sexo"           "Edad_dx"
## [4] "Anciano"        "ECOG"           "IMC"
## [7] "Porcentaje_perdpeso" "p_peso_no_sí"   "Hab_tabaq"
## [10] "Exp_tab"        "Diabetes"       "Cardiop"
## [13] "Enf_neurod"     "Histologia"     "Histología_num"
## [16] "Tamaño_tumor"   "Afectacion_ganglionar" "Afectacion_metastasica"
## [19] "Estadio"        "Estadio_num"    "PD_L1"
## [22] "Estado_mut"     "Estatinas"      "Col_total"
## [25] "LDH"           "Prot_tot"       "Albumina"
## [28] "Hb"            "Leucoc_tot"     "Neutrofilos"
## [31] "Linf_tot"       "Plaquetas"      "NLR_pre"
## [34] "PLR_pre"        "PNI_pre"        "ALI_pre"
## [37] "SII_pre"        "Prot_1C"        "Alb_1C"
## [40] "Hb_1C"          "Leucoc_1C"      "Neutr_1C"
## [43] "Linf_1C"        "Pla_1C"         "NLR_1C"
## [46] "NLR1C_corte4"   "NLR1C_corte5"   "PLR_1C"
## [49] "PNI_1C"         "SII_1C"         "Prot_2C"
## [52] "Alb_2C"         "Hb_2C"          "Leucoc_2C"
## [55] "Neutr_2C"       "Linf_2C"        "Pla_2C"
## [58] "NLR_2C"         "NLR2C_corte4o5" "PLR_2C"
## [61] "PNI_2C"         "SII_2C"         "primera_eval_num"
## [64] "Prot_1eval"     "Alb_1eval"      "Hb_1eval"
## [67] "Leucoc_1eval"   "Neutr_1eval"    "Linf_1eval"
## [70] "Pla_1eval"      "NLR_1eval"      "PLR_1eval"
## [73] "PNI_1eval"      "SII_1eval"      "Mejor_resp_num"
## [76] "N_ciclos"       "Toxicidad"      "Tipo_tox"
## [79] "Grado_tox"      "Interrupc_tto"  "Motivo_inter"
## [82] "Progresion"     "segunda_eval"   "Exitus"
## [85] "SLP"           "SLP_cens"       "SG"
## [88] "SG_cens"
```

2.3 Análisis de datos faltantes (Filas)

Hasta ahora, hemos analizado los datos faltantes por columna, pero ahora es momento de hacerlo por filas, así que procedamos con ello.

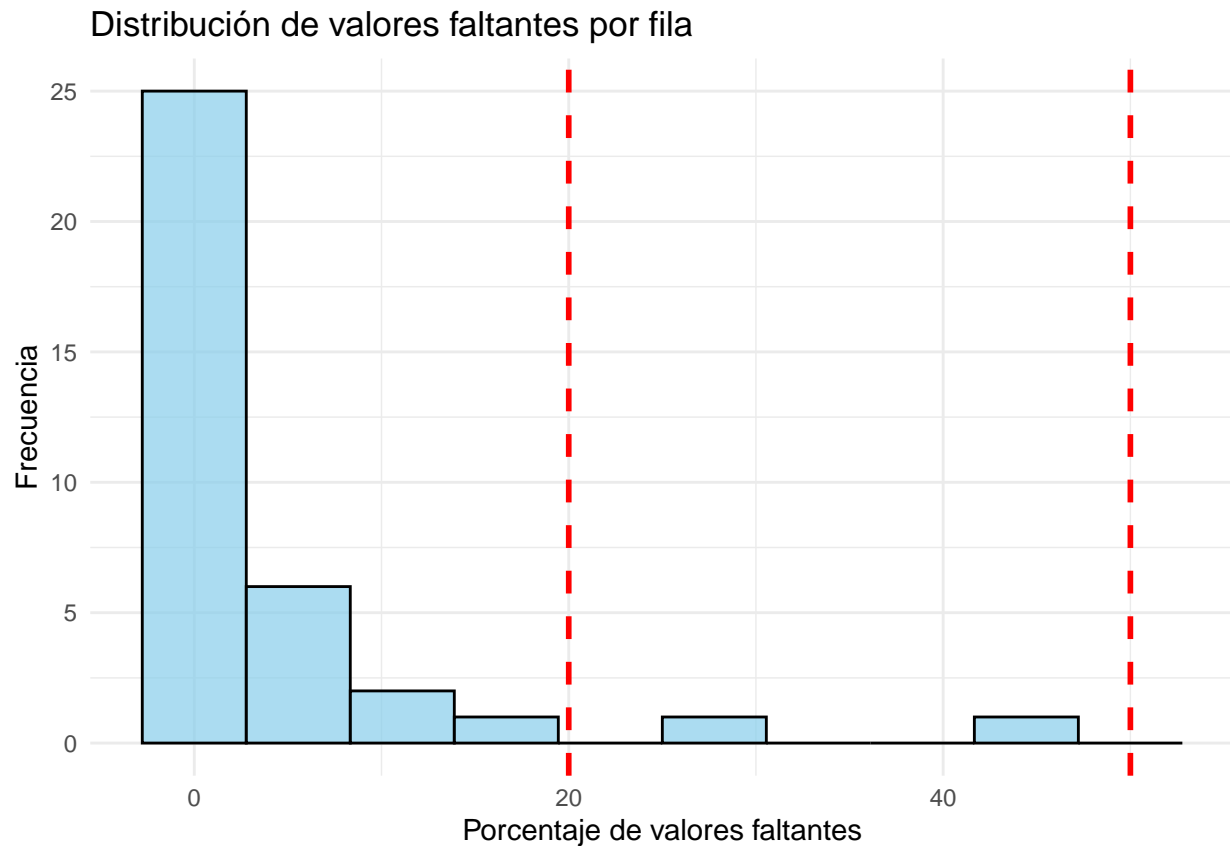
```
datos_limpios$Porcentaje_NA_Fila <- rowSums(is.na(datos_limpios)) / ncol(datos_limpios) * 100

# Crear un dataframe con los porcentajes de valores faltantes
df_missing <- data.frame(Porcentaje_NA_Fila = datos_limpios$Porcentaje_NA_Fila)

ggplot(df_missing, aes(x = Porcentaje_NA_Fila)) +
  geom_histogram(bins = 10, fill = "skyblue", color = "black", alpha = 0.7) +
  geom_vline(xintercept = 20, color = "red", linetype = "dashed", linewidth = 1, label = "Umbral 20%") +
  geom_vline(xintercept = 50, color = "red", linetype = "dashed", linewidth = 1, label = "Umbral 50%") +
  labs(title = "Distribución de valores faltantes por fila",
       x = "Porcentaje de valores faltantes",
       y = "Frecuencia") +
  theme_minimal()
```

```
## Warning in geom_vline(xintercept = 20, color = "red", linetype = "dashed", :  
## Ignoring unknown parameters: 'label'
```

```
## Warning in geom_vline(xintercept = 50, color = "red", linetype = "dashed", :  
## Ignoring unknown parameters: 'label'
```



Dado que solo dos pacientes tienen un porcentaje de valores faltantes del 28.4% y 45.45%, respectivamente, y considerando que la cantidad de datos perdidos en estas filas es significativa, procederemos a eliminarlos para evitar posibles sesgos o problemas en el análisis.

```
# Filtrar las filas donde el porcentaje de valores faltantes es menor al 28%  
datos_limpios <- datos_limpios %>%  
  filter(Porcentaje_NA_Fila < 28)  
  
datos = datos_limpios
```

3. IMPUTACION

En este apartado abordaremos la imputación de datos faltantes, explorando distintos métodos para su tratamiento, como la imputación por la media, la mediana y técnicas más avanzadas según el tipo de variable y el patrón de los datos perdidos.

```
tipos_var <- data.frame(Columna = names(datos))
tipos_var$Tipo <- sapply(datos, function(x) class(x)[1])
tipos_var
```

```
##           Columna      Tipo
## 1           Idpac character
## 2           Sexo  numeric
## 3         Edad_dx  numeric
## 4         Anciano  numeric
## 5           ECOG  numeric
## 6           IMC  numeric
## 7 Porcentaje_perdpeso  numeric
## 8         p_peso_no_sí  numeric
## 9         Hab_tabaq  numeric
## 10          Exp_tab  numeric
## 11          Diabetes  numeric
## 12          Cardiop  numeric
## 13        Enf_neurod  numeric
## 14          Histologia character
## 15      Histología_num  numeric
## 16        Tamaño_tumor character
## 17  Afectacion_ganglionar character
## 18  Afectacion_metastasica character
## 19          Estadio character
## 20        Estadio_num  numeric
## 21          PD_L1  numeric
## 22        Estado_mut character
## 23        Estatinas character
## 24          Col_total  numeric
## 25          LDH  numeric
## 26          Prot_tot  numeric
## 27          Albumina  numeric
## 28          Hb  numeric
## 29        Leucoc_tot  numeric
## 30        Neutrofilos  numeric
## 31          Linf_tot  numeric
## 32          Plaquetas  numeric
## 33          NLR_pre  numeric
## 34          PLR_pre  numeric
## 35          PNI_pre  numeric
## 36          ALI_pre  numeric
## 37          SII_pre  numeric
## 38          Prot_1C  numeric
## 39          Alb_1C  numeric
## 40          Hb_1C  numeric
## 41        Leucoc_1C  numeric
## 42        Neutr_1C  numeric
## 43          Linf_1C  numeric
## 44          Plaq_1C  numeric
## 45          NLR_1C  numeric
## 46        NLR1C_corte4  numeric
## 47        NLR1C_corte5  numeric
## 48          PLR_1C  numeric
```

```

## 49          PNI_1C    numeric
## 50          SII_1C    numeric
## 51          Prot_2C    numeric
## 52          Alb_2C    numeric
## 53          Hb_2C     numeric
## 54          Leucoc_2C  numeric
## 55          Neutr_2C   numeric
## 56          Linf_2C    numeric
## 57          Plaq_2C    numeric
## 58          NLR_2C     numeric
## 59          NLR2C_corte4o5  numeric
## 60          PLR_2C     numeric
## 61          PNI_2C     numeric
## 62          SII_2C     numeric
## 63          primera_eval_num  numeric
## 64          Prot_1eval  numeric
## 65          Alb_1eval  numeric
## 66          Hb_1eval   numeric
## 67          Leucoc_1eval  numeric
## 68          Neutr_1eval  numeric
## 69          Linf_1eval  numeric
## 70          Plaq_1eval  numeric
## 71          NLR_1eval   numeric
## 72          PLR_1eval   numeric
## 73          PNI_1eval   numeric
## 74          SII_1eval   numeric
## 75          Mejor_resp_num  numeric
## 76          N_ciclos    numeric
## 77          Toxicidad   numeric
## 78          Tipo_tox    character
## 79          Grado_tox   character
## 80          Interrupc_tto  numeric
## 81          Motivo_inter  character
## 82          Progresion   numeric
## 83          segunda_eval  numeric
## 84          Exitus       numeric
## 85          SLP          numeric
## 86          SLP_cens     numeric
## 87          SG           numeric
## 88          SG_cens      numeric
## 89          Porcentaje_NA_Fila  numeric

```

Antes de continuar, tenemos que transformar a *factor* aquellas variables que son de naturaleza categorica.

```

datos$Idpac <- as.factor(datos$Idpac)
datos$Sexo <- as.factor(datos$Sexo)
datos$ECOG <- as.factor(datos$ECOG)
datos$Hab_tabaq <- as.factor(datos$Hab_tabaq)
datos$p_peso_no_sí <- as.factor(datos$p_peso_no_sí)
datos$Diabetes <- as.factor(datos$Diabetes)
datos$Cardiop <- as.factor(datos$Cardiop)
datos$Enf_neurod <- as.factor(datos$Enf_neurod)
datos$Histologia <- as.factor(datos$Histologia)
datos$Histología_num <- as.factor(datos$Histología_num)

```

```

datos$Tamaño_tumor <- as.factor(datos$Tamaño_tumor)
datos$Afectacion_ganglionar <- as.factor(datos$Afectacion_ganglionar)
datos$Afectacion_metastasis <- as.factor(datos$Afectacion_metastasis)
datos$Estadio <- as.factor(datos$Estadio)
datos$Estadio_num <- as.factor(datos$Estadio_num)
datos$Estado_mut <- as.factor(datos$Estado_mut)
datos$Estatinas <- as.factor(datos$Estatinas)
datos$Mejor_resp_num <- as.factor(datos$Mejor_resp_num)
datos$NLR1C_corte4 <- as.factor(datos$NLR1C_corte4)
datos$NLR1C_corte5 <- as.factor(datos$NLR1C_corte5)
datos$primera_eval_num <- as.factor(datos$primera_eval_num)
datos$Toxicidad <- as.factor(datos$Toxicidad)
datos$Tipo_tox <- as.factor(datos$Tipo_tox)
datos$Grado_tox <- as.factor(datos$Grado_tox)
datos$Interrupc_tto <- as.factor(datos$Interrupc_tto)
datos$Motivo_inter <- as.factor(datos$Motivo_inter)
datos$Progresion <- as.factor(datos$Progresion)
datos$segunda_eval <- as.factor(datos$segunda_eval)
datos$Exitus <- as.factor(datos$Exitus)

```

Comprobamos que el cambio ha sido correcto

```

tipos_var <- data.frame(Columna = names(datos))
tipos_var$Tipo <- sapply(datos, function(x) class(x)[1])
tipos_var

```

```

##           Columna  Tipo
## 1           Idpac  factor
## 2            Sexo  factor
## 3          Edad_dx numeric
## 4          Anciano numeric
## 5            ECOG  factor
## 6             IMC numeric
## 7 Porcentaje_perdpeso numeric
## 8      p_peso_no_sí  factor
## 9      Hab_tabaq  factor
## 10         Exp_tab numeric
## 11         Diabetes  factor
## 12         Cardiop  factor
## 13        Enf_neurod  factor
## 14        Histologia  factor
## 15   Histología_num  factor
## 16    Tamaño_tumor  factor
## 17 Afectacion_ganglionar  factor
## 18 Afectacion_metastasis  factor
## 19           Estadio  factor
## 20    Estadio_num  factor
## 21          PD_L1 numeric
## 22    Estado_mut  factor
## 23    Estatinas  factor
## 24    Col_total numeric
## 25          LDH numeric
## 26    Prot_tot numeric

```

```

## 27          Albumina numeric
## 28              Hb numeric
## 29          Leucoc_tot numeric
## 30          Neutrofilos numeric
## 31              Linf_tot numeric
## 32          Plaquetas numeric
## 33              NLR_pre numeric
## 34              PLR_pre numeric
## 35              PNI_pre numeric
## 36              ALI_pre numeric
## 37              SII_pre numeric
## 38              Prot_1C numeric
## 39              Alb_1C numeric
## 40              Hb_1C numeric
## 41          Leucoc_1C numeric
## 42          Neutr_1C numeric
## 43              Linf_1C numeric
## 44              Plaq_1C numeric
## 45              NLR_1C numeric
## 46          NLR1C_corte4  factor
## 47          NLR1C_corte5  factor
## 48              PLR_1C numeric
## 49              PNI_1C numeric
## 50              SII_1C numeric
## 51          Prot_2C numeric
## 52          Alb_2C numeric
## 53          Hb_2C numeric
## 54          Leucoc_2C numeric
## 55          Neutr_2C numeric
## 56          Linf_2C numeric
## 57          Plaq_2C numeric
## 58          NLR_2C numeric
## 59          NLR2C_corte4o5 numeric
## 60              PLR_2C numeric
## 61              PNI_2C numeric
## 62              SII_2C numeric
## 63          primera_eval_num  factor
## 64              Prot_1eval numeric
## 65              Alb_1eval numeric
## 66              Hb_1eval numeric
## 67          Leucoc_1eval numeric
## 68          Neutr_1eval numeric
## 69          Linf_1eval numeric
## 70          Plaq_1eval numeric
## 71          NLR_1eval numeric
## 72          PLR_1eval numeric
## 73          PNI_1eval numeric
## 74          SII_1eval numeric
## 75          Mejor_resp_num  factor
## 76              N_ciclos numeric
## 77          Toxicidad  factor
## 78          Tipo_tox  factor
## 79          Grado_tox  factor
## 80          Interrupc_tto  factor

```



```
## 81      Motivo_inter factor
## 82      Progresion factor
## 83      segunda_eval factor
## 84      Exitus factor
## 85      SLP numeric
## 86      SLP_cens numeric
## 87      SG numeric
## 88      SG_cens numeric
## 89      Porcentaje_NA_Fila numeric
```

3.1. Metodo de la media

Una vez localizados los tipos de variables, vamos a imputar las variables numericas haciendo uso de la libreria **mice**

```
# Excluir automáticamente todas las variables que sean factores
imputed_data1 <- mice(datos %>%
  select(which(!sapply(datos, is.factor))),
  method = "mean", print = FALSE)
```

```
## Warning: Number of logged events: 720
```

```
df_imputado1 <- complete(imputed_data1)
df_imputado1
```

##	Edad_dx	Anciano	IMC	Porcentaje_perdpeso	Exp_tab	PD_L1	Col_total
## 1	46.00000	0	28.08899	0.04583871	20.00000	100	217.0000
## 2	68.00000	0	35.37981	0.00000000	0.00000	70	154.0000
## 3	59.00000	0	25.71101	0.00000000	45.00000	50	146.0000
## 4	72.00000	1	28.24859	0.00000000	92.00000	2	149.0000
## 5	50.00000	0	29.05475	0.10000000	55.00000	60	210.0000
## 6	71.00000	1	28.83059	0.00000000	200.00000	90	162.0000
## 7	71.00000	1	29.77778	0.07600000	60.00000	100	142.0000
## 8	79.00000	1	25.84777	0.00000000	40.00000	70	167.0000
## 9	73.14168	1	22.32143	0.04583871	50.00000	90	153.0000
## 10	68.51745	0	23.65618	0.00000000	50.00000	70	132.0000
## 11	56.48734	0	24.09796	0.04583871	36.00000	80	113.0000
## 12	66.27242	0	24.95389	0.22900000	50.00000	50	182.0000
## 13	61.96304	0	26.21631	0.13800000	0.00000	80	242.0000
## 14	64.45996	0	20.28651	0.11000000	50.00000	100	133.0000
## 15	58.70500	0	22.06035	0.00000000	25.00000	95	215.0000
## 16	60.30664	0	24.62296	0.00000000	82.00000	90	202.0000
## 17	63.00000	0	29.77778	0.00000000	60.00000	100	262.0000
## 18	82.00000	1	22.95909	0.00000000	100.00000	90	200.0000
## 19	66.00000	0	24.76756	0.00000000	50.00000	60	199.0000
## 20	50.00000	0	20.70082	0.00000000	51.65625	100	194.0000
## 21	78.00000	1	28.72008	0.06700000	75.00000	70	176.0000
## 22	73.00000	1	27.95976	0.01000000	83.00000	70	160.0000
## 23	60.00000	0	22.38631	0.13000000	40.00000	100	111.0000
## 24	70.00000	1	25.27344	0.03000000	51.65625	90	276.0000
## 25	64.00000	0	18.35937	0.03100000	40.00000	70	155.0000
## 26	57.00000	0	23.24341	0.04000000	30.00000	70	197.0000

## 27	71.00000	1	21.36752		0.05500000	0.00000	80	206.0000
## 28	75.00000	1	24.21875		0.08800000	25.00000	100	208.0000
## 29	68.00000	0	26.57313		0.03800000	40.00000	90	191.0000
## 30	62.00000	0	27.04164		0.00000000	60.00000	90	201.0000
## 31	51.00000	0	18.92494		0.07500000	45.00000	60	142.0000
## 32	80.00000	1	21.23057		0.15900000	40.00000	80	187.4545
## 33	65.00000	0	31.22130		0.04500000	85.00000	95	192.0000
## 34	62.00000	0	35.62902		0.00000000	25.00000	70	397.0000
##	LDH	Prot_tot	Albumina	Hb	Leucoc_tot	Neutrofilos	Linf_tot	Plaquetas
## 1	163.0000	7.000000	4.100000	15.3	10100	8000	1400	273000
## 2	171.0000	6.000000	3.700000	11.8	9800	8200	800	124000
## 3	197.0000	7.200000	4.300000	16.7	9900	7400	1400	349000
## 4	159.0000	7.100000	4.000000	14.6	5600	3000	1800	220000
## 5	198.0000	7.900000	4.400000	15.7	8900	5300	2300	350000
## 6	220.0000	8.000000	4.300000	13.1	8800	6000	1600	279000
## 7	198.0000	6.400000	3.600000	11.5	10000	9000	400	269000
## 8	184.0000	7.000000	3.900000	12.3	7700	5300	1600	201000
## 9	176.0000	6.900000	3.700000	12.9	16200	13200	1900	317000
## 10	199.0000	7.300000	4.000000	13.0	15700	13400	1400	324000
## 11	190.0000	7.300000	3.500000	13.6	10600	8000	1500	489000
## 12	198.0000	6.500000	3.700000	11.4	8700	5000	2500	216000
## 13	184.0000	7.300000	4.100000	13.7	12600	8700	2200	455000
## 14	183.0000	7.000000	4.100000	13.7	11400	7600	2300	274000
## 15	204.0000	7.200000	4.400000	11.9	13400	9400	2900	208000
## 16	276.0000	5.600000	3.200000	15.5	13100	12300	500	112000
## 17	201.0000	5.400000	2.800000	11.6	4100	2500	800	293000
## 18	383.0000	8.900000	3.900000	12.1	11700	7100	2700	358000
## 19	382.0000	6.300000	3.700000	10.6	4000	2600	900	338000
## 20	156.0000	7.300000	4.300000	11.5	10800	9400	500	464000
## 21	175.0000	7.100000	4.300000	11.5	10600	7300	1200	158000
## 22	195.0000	7.300000	4.100000	12.6	7700	4800	1900	248000
## 23	326.0000	6.400000	3.000000	13.5	6800	5400	700	335000
## 24	263.6452	7.000000	3.900000	14.1	6000	3500	2000	143000
## 25	1644.0000	6.900000	3.400000	10.9	13800	8700	3000	709000
## 26	279.0000	6.900000	4.400000	13.0	6300	4300	1100	287000
## 27	164.0000	6.600000	4.000000	14.0	6200	4800	800	295000
## 28	222.0000	7.400000	4.100000	12.4	7200	4900	1500	228000
## 29	267.0000	6.900000	3.500000	14.5	20500	11500	3300	271000
## 30	248.0000	7.300000	4.200000	13.6	14100	8400	3500	461000
## 31	263.6452	7.100000	3.900000	12.8	9600	7100	1300	327000
## 32	263.6452	7.006061	3.912121	12.6	7600	5300	1300	249000
## 33	230.0000	7.700000	4.400000	15.3	8400	5400	1800	306000
## 34	201.0000	7.000000	4.200000	14.2	12500	10700	1300	234000
##	NLR_pre	PLR_pre	PNI_pre	ALI_pre	SII_pre	Prot_1C	Alb_1C	Hb_1C
## 1	5.714286	195.00000	48.00000	20.153851	1560000.0	7.500000	4.60000	15.9
## 2	10.250000	155.00000	41.00000	12.771249	1271000.0	6.200000	3.90000	11.7
## 3	5.285714	249.28571	50.00000	20.916253	1844714.3	7.200000	4.40000	16.1
## 4	1.666667	122.22222	49.00000	67.796610	366666.7	7.048387	3.90625	15.5
## 5	2.304348	152.17391	55.50000	55.478130	806521.7	7.400000	4.50000	15.9
## 6	3.750000	174.37500	51.00000	33.059079	1046250.0	8.100000	4.20000	14.5
## 7	22.500000	672.50000	38.00000	4.764444	6052500.0	6.500000	4.00000	11.6
## 8	3.312500	125.62500	47.00000	30.432089	665812.5	7.700000	4.00000	12.1
## 9	6.947368	166.84211	46.50000	11.887852	2202315.8	7.048387	3.90000	14.5
## 10	9.571429	231.42857	47.00000	9.886163	3101142.9	7.400000	4.00000	13.5

## 11	5.333333	326.00000	42.50000	15.814286	2608000.0	7.000000	3.40000	11.4
## 12	2.000000	86.40000	49.50000	46.164703	432000.0	7.100000	4.10000	11.6
## 13	3.954545	206.81818	52.00000	27.180583	1799318.2	7.300000	4.00000	14.3
## 14	3.304348	119.13043	52.50000	25.171285	905391.3	7.000000	3.70000	10.8
## 15	3.241379	71.72414	58.50000	29.945757	674206.9	6.900000	3.70000	11.3
## 16	24.600000	224.00000	34.50000	3.202987	2755200.0	7.800000	4.20000	13.7
## 17	3.125000	366.25000	32.00000	26.680889	915625.0	4.600000	3.10000	8.4
## 18	2.629630	132.59259	52.50000	34.050591	941407.4	9.100000	4.00000	11.9
## 19	2.888889	375.55556	41.50000	31.721533	976444.4	6.000000	3.10000	11.1
## 20	18.800000	928.00000	45.50000	4.734761	8723200.0	8.000000	4.50000	11.9
## 21	6.083333	131.66667	49.00000	20.300771	961166.7	7.200000	4.20000	10.2
## 22	2.526316	130.52632	50.50000	45.376360	626526.3	7.300000	3.80000	12.5
## 23	7.714286	478.57143	33.50000	8.705789	2584285.7	6.800000	3.60000	16.0
## 24	1.750000	71.50000	49.00000	56.323661	250250.0	7.400000	3.90000	12.9
## 25	2.900000	236.33333	49.00000	21.524784	2056100.0	7.300000	3.60000	11.1
## 26	3.909091	260.90909	49.50000	26.162348	1121909.1	6.400000	3.70000	11.7
## 27	6.000000	368.75000	44.00000	14.245014	1770000.0	6.900000	4.10000	13.6
## 28	3.266667	152.00000	48.50000	30.397003	744800.0	7.100000	4.00000	11.7
## 29	3.484848	82.12121	51.50000	26.688665	944393.9	6.800000	3.30000	14.5
## 30	2.400000	131.71429	59.50000	47.322877	1106400.0	6.300000	3.80000	11.6
## 31	5.461538	251.53846	45.50000	13.514007	1785923.1	6.500000	3.30000	10.5
## 32	4.076923	191.53846	47.42424	26.858982	1015153.8	7.100000	4.30000	13.2
## 33	3.000000	170.00000	53.00000	45.791246	918000.0	7.048387	3.90625	15.6
## 34	8.230769	180.00000	48.50000	18.180788	1926000.0	6.600000	4.10000	15.4
##	Leucoc_1C	Neutr_1C	Linf_1C	Pla_1C	NLR_1C	PLR_1C	PNI_1C	SII_1C
## 1	8100	6000	1600	369000	3.750000	230.62500	54.00000	1383750.0
## 2	8500	7000	800	112000	8.750000	140.00000	43.00000	980000.0
## 3	12000	8000	2500	382000	3.200000	152.80000	56.50000	1222400.0
## 4	7300	4400	2000	266000	2.200000	133.00000	10.00000	585200.0
## 5	9200	4900	3100	364000	1.580645	117.41935	60.50000	575354.8
## 6	8000	5300	1600	293000	3.312500	183.12500	50.00000	970562.5
## 7	4500	3200	900	218000	3.555556	242.22222	44.50000	775111.1
## 8	3600	1900	900	131000	2.111111	145.55556	44.50000	276555.6
## 9	16100	12100	2900	380000	4.172414	131.03448	53.50000	1585517.2
## 10	12800	9800	1900	214000	5.157895	112.63158	49.50000	1103789.5
## 11	12500	10400	1000	463000	10.400000	463.00000	39.00000	4815200.0
## 12	7600	3700	2900	242000	1.275862	83.44828	55.50000	308758.6
## 13	11300	6200	3600	452000	1.722222	125.55556	58.00000	778444.4
## 14	15400	11300	2500	444000	4.520000	177.60000	49.50000	2006880.0
## 15	10400	8400	1000	272000	8.400000	272.00000	42.00000	2284800.0
## 16	14100	9700	3100	494000	3.129032	159.35484	57.50000	1545741.9
## 17	4800	3300	700	226000	4.714286	322.85714	34.50000	1065428.6
## 18	11100	6000	3000	380000	2.000000	126.66667	55.00000	760000.0
## 19	3600	1700	1100	338000	1.545455	307.27273	36.50000	522363.6
## 20	10900	8900	1200	465000	7.416667	387.50000	51.00000	3448750.0
## 21	13800	9300	1600	205000	5.812500	128.12500	50.00000	1191562.5
## 22	7400	4600	1800	304000	2.555556	168.88889	47.00000	776888.9
## 23	4600	3000	1100	274000	2.727273	249.09091	41.50000	747272.7
## 24	6000	3000	2200	277000	1.363636	125.90909	50.00000	377727.3
## 25	12600	7000	3600	650000	1.944444	180.55556	54.00000	1263888.9
## 26	9000	6300	800	457000	7.875000	571.25000	41.00000	3598875.0
## 27	4900	2700	1600	192000	1.687500	120.00000	49.00000	324000.0
## 28	6900	5900	600	259000	9.833333	431.66667	43.00000	2546833.3
## 29	12800	6800	2800	366000	2.428571	130.71429	47.00000	888857.1

## 30	8700	4600	2200	553000	2.090909	251.36364	49.00000	1156272.7
## 31	9000	6300	1200	513000	5.250000	427.50000	39.00000	2693250.0
## 32	13100	8700	2900	345000	3.000000	118.96552	57.50000	1035000.0
## 33	10000	6800	2000	323000	3.400000	161.50000	47.33333	1098200.0
## 34	10200	7600	1700	208000	4.470588	122.35294	49.50000	929882.4
##	Prot_2C	Alb_2C	Hb_2C	Leucoc_2C	Neutr_2C	Linf_2C	Pla_2C	NLR_2C
## 1	7.900000	4.500000	16.50000	6400.00	3400.000	2300.000	253000.0	1.478261
## 2	6.100000	3.700000	11.40000	8300.00	6200.000	1200.000	126000.0	5.166667
## 3	7.200000	4.400000	16.70000	8400.00	4500.000	3000.000	329000.0	1.500000
## 4	7.200000	4.100000	15.30000	7200.00	4300.000	1900.000	254000.0	2.263158
## 5	8.000000	4.500000	16.30000	9200.00	5100.000	2800.000	313000.0	1.821429
## 6	8.100000	4.500000	15.50000	8000.00	5300.000	1800.000	285.0	2.944444
## 7	7.200000	4.100000	12.40000	8200.00	6400.000	1100.000	265000.0	5.818182
## 8	7.223333	4.012903	13.15312	9106.25	5796.875	2090.625	305883.9	3.348731
## 9	7.223333	3.700000	13.40000	19400.00	16100.000	2300.000	406000.0	7.000000
## 10	7.500000	4.100000	13.40000	8700.00	5500.000	2200.000	226000.0	2.500000
## 11	6.500000	2.900000	8.60000	14500.00	12100.000	1200.000	858000.0	10.083333
## 12	6.800000	3.900000	11.40000	9400.00	4400.000	3100.000	260000.0	1.419355
## 13	7.100000	4.200000	15.00000	10200.00	5100.000	2900.000	319000.0	1.758621
## 14	7.200000	4.000000	10.90000	14700.00	9900.000	3500.000	408000.0	2.828571
## 15	7.200000	3.900000	11.20000	10600.00	7900.000	1400.000	254000.0	5.642857
## 16	7.200000	3.900000	13.30000	9900.00	6100.000	2500.000	297000.0	2.440000
## 17	5.300000	3.000000	10.30000	3800.00	2300.000	800.000	218000.0	2.875000
## 18	8.800000	4.100000	12.90000	11200.00	6100.000	3300.000	272000.0	1.848485
## 19	6.700000	3.700000	11.60000	4000.00	1800.000	1500.000	394000.0	1.200000
## 20	7.900000	4.600000	12.30000	8600.00	6300.000	1200.000	425000.0	5.250000
## 21	6.900000	4.400000	10.40000	12500.00	8500.000	1500.000	217000.0	5.666667
## 22	7.700000	3.900000	13.40000	6400.00	3600.000	1800.000	194000.0	2.000000
## 23	7.800000	3.600000	15.80000	5400.00	3700.000	1000.000	237000.0	3.700000
## 24	7.400000	4.000000	13.20000	5100.00	2400.000	2100.000	240000.0	1.142857
## 25	7.500000	3.800000	11.40000	13900.00	7900.000	3900.000	615000.0	2.025641
## 26	7.223333	4.012903	13.15312	9106.25	5796.875	2090.625	305883.9	3.348731
## 27	6.800000	4.200000	13.50000	5300.00	3300.000	1400.000	235000.0	2.357143
## 28	7.100000	4.000000	11.70000	6900.00	5900.000	600.000	259000.0	9.833333
## 29	7.500000	3.700000	13.80000	12000.00	5700.000	4000.000	256000.0	1.425000
## 30	6.300000	4.100000	13.00000	8300.00	4100.000	2900.000	343000.0	1.413793
## 31	7.223333	4.012903	11.40000	7600.00	4900.000	1300.000	412000.0	3.769231
## 32	7.500000	4.500000	12.90000	8800.00	5800.000	1800.000	337000.0	3.222222
## 33	7.700000	4.300000	15.70000	9600.00	5900.000	2200.000	312000.0	2.681818
## 34	6.600000	4.100000	16.30000	8900.00	5000.000	2400.000	254000.0	2.083333
##	NLR2C_corte4o5	PLR_2C	PNI_2C	SII_2C	Prot_1eval	Alb_1eval		
## 1	0.00	110.0000000	56.50000	374000.0000	7.300000	4.4		
## 2	1.00	105.0000000	43.00000	651000.0000	6.000000	3.9		
## 3	0.00	109.6666667	59.00000	493500.0000	7.100000	4.5		
## 4	0.00	133.6842105	50.50000	574842.1053	7.200000	4.0		
## 5	0.00	111.7857143	59.00000	570107.1429	7.087097	4.3		
## 6	0.00	0.1583333	54.00000	839.1667	7.600000	4.4		
## 7	1.00	240.9090909	46.50000	1541818.1818	6.700000	3.9		
## 8	0.25	177.5906414	50.70968	1134689.6890	6.400000	4.2		
## 9	1.00	176.5217391	48.50000	2842000.0000	6.900000	3.7		
## 10	0.00	102.7272727	52.00000	565000.0000	7.700000	4.5		
## 11	1.00	715.0000000	35.00000	8651500.0000	7.000000	3.1		
## 12	0.00	83.8709677	54.50000	369032.2581	7.200000	4.0		
## 13	0.00	110.0000000	56.50000	561000.0000	7.100000	4.2		

## 14	0.00	116.5714286	57.50000	1154057.1429	8.000000	4.5	
## 15	1.00	181.4285714	46.00000	1433285.7143	7.087097	3.8	
## 16	0.00	118.8000000	51.50000	724680.0000	7.000000	4.1	
## 17	0.00	272.5000000	34.00000	626750.0000	5.500000	3.6	
## 18	0.00	82.4242424	57.50000	502787.8788	8.500000	4.1	
## 19	0.00	262.6666667	44.50000	472800.0000	7.100000	3.9	
## 20	1.00	354.1666667	52.00000	2231250.0000	7.300000	3.7	
## 21	1.00	144.6666667	51.50000	1229666.6667	6.400000	3.9	
## 22	0.00	107.7777778	48.00000	388000.0000	7.700000	3.9	
## 23	0.00	237.0000000	41.00000	876900.0000	7.087097	3.1	
## 24	0.00	114.2857143	50.50000	274285.7143	7.700000	4.1	
## 25	0.00	157.6923077	57.50000	1245769.2308	7.800000	3.9	
## 26	0.25	177.5906414	50.70968	1134689.6890	6.400000	3.7	
## 27	0.00	167.8571429	49.00000	553928.5714	7.000000	4.2	
## 28	1.00	431.6666667	43.00000	2546833.3333	7.100000	4.0	
## 29	0.00	64.0000000	57.00000	364800.0000	6.900000	3.9	
## 30	0.00	118.2758621	55.50000	484931.0345	6.300000	4.1	
## 31	0.00	316.9230769	50.70968	1552923.0769	7.000000	4.0	
## 32	0.00	187.2222222	54.00000	1085888.8889	7.700000	4.6	
## 33	0.00	141.8181818	54.00000	836727.2727	7.500000	4.3	
## 34	0.00	105.8333333	53.00000	529166.6667	6.600000	4.1	
##	Hb_1eval	Leucoc_1eval	Neutr_1eval	Linf_1eval	Plaq_1eval	NLR_1eval	PLR_1eval
## 1	16.9	8000	5300	2100	262000	2.523810	124.76190
## 2	11.2	7500	5500	1300	96000	4.230769	73.84615
## 3	16.6	8700	4900	2900	315000	1.689655	108.62069
## 4	15.7	7800	4400	2500	259000	1.760000	103.60000
## 5	16.5	8700	4600	2800	285000	1.642857	101.78571
## 6	15.4	8000	5900	1400	263000	4.214286	187.85714
## 7	11.1	7600	6000	1000	229000	6.000000	229.00000
## 8	11.6	8600	6300	1400	124000	4.500000	88.57143
## 9	12.9	16200	13200	1900	317000	6.947368	166.84211
## 10	14.6	7400	4700	1900	219000	2.473684	115.26316
## 11	8.8	10200	9100	500	527000	18.200000	1054.00000
## 12	10.8	11100	7200	2600	227000	2.769231	87.30769
## 13	15.1	9100	5000	2900	308000	1.724138	106.20690
## 14	12.9	18400	13600	3200	328000	4.250000	102.50000
## 15	11.3	9400	7200	900	309000	8.000000	343.33333
## 16	15.8	9800	6000	2600	205000	2.307692	78.84615
## 17	13.7	4600	2600	1100	220000	2.363636	200.00000
## 18	12.7	10200	5400	3100	282000	1.741935	90.96774
## 19	12.2	3100	1400	1300	290000	1.076923	223.07692
## 20	12.5	9200	7100	1100	400000	6.454545	363.63636
## 21	9.9	10000	7500	800	216000	9.375000	270.00000
## 22	13.4	6400	3600	1800	194000	2.000000	107.77778
## 23	14.9	4500	2900	1000	220000	2.900000	220.00000
## 24	13.8	4300	2200	1600	234000	1.375000	146.25000
## 25	11.8	13900	7700	4000	647000	1.925000	161.75000
## 26	11.7	9000	6300	800	457000	7.875000	571.25000
## 27	13.8	4600	3100	1100	257000	2.818182	233.63636
## 28	11.7	6900	5900	600	259000	9.833333	431.66667
## 29	14.5	9100	4200	3500	199000	1.200000	56.85714
## 30	13.0	8300	4100	2900	343000	1.413793	118.27586
## 31	11.3	7300	4300	1800	353000	2.388889	196.11111
## 32	12.9	9700	6500	2100	308000	3.095238	146.66667

## 33	15.9	8700	6000	1600	320000	3.750000	200.00000
## 34	16.3	8900	5000	2400	254000	2.083333	105.83333
##	PNI_1eval	SII_1eval	N_ciclos	SLP	SLP_cens	SG	SG_cens
## 1	54.5	661238.1	35	29.8644764	1	29.864476	1
## 2	45.5	406153.8	35	60.3860370	1	60.386037	1
## 3	59.5	532241.4	7	35.1868583	0	57.626283	1
## 4	52.5	455840.0	11	7.3921971	0	33.741273	0
## 5	57.0	468214.3	7	4.5010267	0	18.825462	0
## 6	51.0	1108357.1	4	41.4948665	1	41.494867	1
## 7	44.0	1374000.0	19	24.0164271	0	30.225873	0
## 8	49.0	558000.0	1	7.8850103	1	7.885010	1
## 9	46.5	2202315.8	3	1.8069815	0	1.806982	0
## 10	54.5	541736.8	26	19.9425051	1	19.942505	0
## 11	33.5	9591400.0	3	1.8726899	0	17.478439	0
## 12	53.0	628615.4	5	40.7392197	1	40.739220	0
## 13	56.5	531034.5	11	7.4579055	0	41.002053	0
## 14	61.0	1394000.0	6	5.6509240	0	8.837782	0
## 15	42.5	2472000.0	6	3.6796715	0	6.570842	0
## 16	54.0	473076.9	32	22.3737166	0	28.747433	0
## 17	41.5	520000.0	25	21.9794661	1	21.979466	0
## 18	56.5	491225.8	35	46.3244353	1	46.324435	1
## 19	45.5	312307.7	16	35.3182752	0	35.318275	1
## 20	42.5	2581818.2	25	45.6344969	1	45.634497	1
## 21	43.0	2025000.0	3	1.9055441	0	5.749487	0
## 22	48.0	388000.0	5	7.2607803	0	22.078029	0
## 23	36.0	638000.0	8	9.3305955	1	9.330595	0
## 24	49.0	321750.0	8	7.1293634	0	16.131417	0
## 25	59.0	1245475.0	8	5.6180698	0	10.611910	0
## 26	41.0	3598875.0	1	0.7556468	0	41.166324	0
## 27	47.5	724272.7	35	31.1457906	1	31.145791	1
## 28	43.0	2546833.3	2	1.3798768	0	1.839836	0
## 29	56.5	238800.0	27	17.8069815	0	26.579055	0
## 30	55.5	484931.0	13	11.1704312	0	22.702259	0
## 31	49.0	843277.8	4	1.8069815	0	16.000000	0
## 32	56.5	953333.3	35	40.2135524	1	40.213552	1
## 33	51.0	1200000.0	23	16.1642710	0	35.778234	0
## 34	53.0	529166.7	35	39.4579055	1	39.457906	1
##	Porcentaje_NA_Fila						
## 1	3.409091						
## 2	0.000000						
## 3	0.000000						
## 4	3.409091						
## 5	2.272727						
## 6	1.136364						
## 7	1.136364						
## 8	14.772727						
## 9	5.681818						
## 10	0.000000						
## 11	2.272727						
## 12	0.000000						
## 13	0.000000						
## 14	0.000000						
## 15	1.136364						
## 16	0.000000						

```
## 17          1.136364
## 18          1.136364
## 19          0.000000
## 20          2.272727
## 21          0.000000
## 22          1.136364
## 23          2.272727
## 24          6.818182
## 25          0.000000
## 26         13.636364
## 27          0.000000
## 28          1.136364
## 29          0.000000
## 30          0.000000
## 31          5.681818
## 32         10.227273
## 33          4.545455
## 34          1.136364
```

```
df_imputado1$Edad_dx <- round(df_imputado1$Edad_dx, 0)
df_imputado1$LDH <- round(df_imputado1$LDH, 0)
df_imputado1$LDH <- round(df_imputado1$LDH, 0)
df_imputado1$Exp_tab <- round(df_imputado1$Exp_tab, 0)
df_imputado1$Col_total <- round(df_imputado1$Col_total, 0)
df_imputado1$Leucoc_1C <- round(df_imputado1$Leucoc_1C, 0)
df_imputado1$Neutr_1C <- round(df_imputado1$Neutr_1C, 0)
df_imputado1$Linf_1C <- round(df_imputado1$Linf_1C, 0)
df_imputado1$Pla_1C <- round(df_imputado1$Pla_1C, 0)
df_imputado1$Prot_1C <- round(df_imputado1$Prot_1C, 1)
df_imputado1$Alb_1C <- round(df_imputado1$Alb_1C, 1)
df_imputado1$Hb_1C <- round(df_imputado1$Hb_1C, 1)
df_imputado1$Prot_tot <- round(df_imputado1$Prot_tot, 1)
df_imputado1$Albumina <- round(df_imputado1$Albumina, 1)
df_imputado1$Porcentaje_perdpeso <- round(df_imputado1$Porcentaje_perdpeso, 4)
df_imputado1$NLR_pre <- round(df_imputado1$NLR_pre, 2)
df_imputado1$PLR_pre <- round(df_imputado1$PLR_pre, 2)
df_imputado1$PNI_pre <- round(df_imputado1$PNI_pre, 2)
df_imputado1$ALI_pre <- round(df_imputado1$ALI_pre, 2)
df_imputado1$SII_pre <- round(df_imputado1$SII_pre, 2)
df_imputado1$NLR_1C <- round(df_imputado1$NLR_1C, 2)
df_imputado1$PLR_2C <- round(df_imputado1$PLR_2C, 2)
df_imputado1$SII_2C <- round(df_imputado1$SII_2C, 2)
df_imputado1$IMC <- round(df_imputado1$IMC , 2)
df_imputado1$PLR_1C <- round(df_imputado1$PLR_1C, 2)
df_imputado1$PNI_1C <- round(df_imputado1$PNI_1C, 2)
df_imputado1$Prot_2C <- round(df_imputado1$Prot_2C, 2)
df_imputado1$Alb_2C <- round(df_imputado1$Alb_2C, 2)
df_imputado1$Hb_2C <- round(df_imputado1$Hb_2C, 2)
df_imputado1$Leucoc_2C <- round(df_imputado1$Leucoc_2C, 2)
df_imputado1$Neutr_2C <- round(df_imputado1$Neutr_2C, 2)
df_imputado1$Linf_2C <- round(df_imputado1$Linf_2C, 2)
df_imputado1$PNI_2C <- round(df_imputado1$PNI_2C, 2)
df_imputado1$Prot_1eval <- round(df_imputado1$Prot_1eval, 2)
df_imputado1$NLR_1eval <- round(df_imputado1$NLR_1eval, 2)
```

```
df_imputado1$PLR_1eval <- round(df_imputado1$PLR_1eval, 2)
df_imputado1$SLP <- round(df_imputado1$SLP, 2)
df_imputado1$SG <- round(df_imputado1$SG, 2)
```

```
df_imputado1
```

##	Edad_dx	Anciano	IMC	Porcentaje_perdpeso	Exp_tab	PD_L1	Col_total	LDH
## 1	46	0	28.09	0.0458	20	100	217	163
## 2	68	0	35.38	0.0000	0	70	154	171
## 3	59	0	25.71	0.0000	45	50	146	197
## 4	72	1	28.25	0.0000	92	2	149	159
## 5	50	0	29.05	0.1000	55	60	210	198
## 6	71	1	28.83	0.0000	200	90	162	220
## 7	71	1	29.78	0.0760	60	100	142	198
## 8	79	1	25.85	0.0000	40	70	167	184
## 9	73	1	22.32	0.0458	50	90	153	176
## 10	69	0	23.66	0.0000	50	70	132	199
## 11	56	0	24.10	0.0458	36	80	113	190
## 12	66	0	24.95	0.2290	50	50	182	198
## 13	62	0	26.22	0.1380	0	80	242	184
## 14	64	0	20.29	0.1100	50	100	133	183
## 15	59	0	22.06	0.0000	25	95	215	204
## 16	60	0	24.62	0.0000	82	90	202	276
## 17	63	0	29.78	0.0000	60	100	262	201
## 18	82	1	22.96	0.0000	100	90	200	383
## 19	66	0	24.77	0.0000	50	60	199	382
## 20	50	0	20.70	0.0000	52	100	194	156
## 21	78	1	28.72	0.0670	75	70	176	175
## 22	73	1	27.96	0.0100	83	70	160	195
## 23	60	0	22.39	0.1300	40	100	111	326
## 24	70	1	25.27	0.0300	52	90	276	264
## 25	64	0	18.36	0.0310	40	70	155	1644
## 26	57	0	23.24	0.0400	30	70	197	279
## 27	71	1	21.37	0.0550	0	80	206	164
## 28	75	1	24.22	0.0880	25	100	208	222
## 29	68	0	26.57	0.0380	40	90	191	267
## 30	62	0	27.04	0.0000	60	90	201	248
## 31	51	0	18.92	0.0750	45	60	142	264
## 32	80	1	21.23	0.1590	40	80	187	264
## 33	65	0	31.22	0.0450	85	95	192	230
## 34	62	0	35.63	0.0000	25	70	397	201
##	Prot_tot	Albumina	Hb	Leucoc_tot	Neutrofilos	Linf_tot	Plaquetas	NLR_pre
## 1	7.0	4.1	15.3	10100	8000	1400	273000	5.71
## 2	6.0	3.7	11.8	9800	8200	800	124000	10.25
## 3	7.2	4.3	16.7	9900	7400	1400	349000	5.29
## 4	7.1	4.0	14.6	5600	3000	1800	220000	1.67
## 5	7.9	4.4	15.7	8900	5300	2300	350000	2.30
## 6	8.0	4.3	13.1	8800	6000	1600	279000	3.75
## 7	6.4	3.6	11.5	10000	9000	400	269000	22.50
## 8	7.0	3.9	12.3	7700	5300	1600	201000	3.31
## 9	6.9	3.7	12.9	16200	13200	1900	317000	6.95
## 10	7.3	4.0	13.0	15700	13400	1400	324000	9.57
## 11	7.3	3.5	13.6	10600	8000	1500	489000	5.33

## 12	6.5	3.7	11.4	8700	5000	2500	216000	2.00	
## 13	7.3	4.1	13.7	12600	8700	2200	455000	3.95	
## 14	7.0	4.1	13.7	11400	7600	2300	274000	3.30	
## 15	7.2	4.4	11.9	13400	9400	2900	208000	3.24	
## 16	5.6	3.2	15.5	13100	12300	500	112000	24.60	
## 17	5.4	2.8	11.6	4100	2500	800	293000	3.12	
## 18	8.9	3.9	12.1	11700	7100	2700	358000	2.63	
## 19	6.3	3.7	10.6	4000	2600	900	338000	2.89	
## 20	7.3	4.3	11.5	10800	9400	500	464000	18.80	
## 21	7.1	4.3	11.5	10600	7300	1200	158000	6.08	
## 22	7.3	4.1	12.6	7700	4800	1900	248000	2.53	
## 23	6.4	3.0	13.5	6800	5400	700	335000	7.71	
## 24	7.0	3.9	14.1	6000	3500	2000	143000	1.75	
## 25	6.9	3.4	10.9	13800	8700	3000	709000	2.90	
## 26	6.9	4.4	13.0	6300	4300	1100	287000	3.91	
## 27	6.6	4.0	14.0	6200	4800	800	295000	6.00	
## 28	7.4	4.1	12.4	7200	4900	1500	228000	3.27	
## 29	6.9	3.5	14.5	20500	11500	3300	271000	3.48	
## 30	7.3	4.2	13.6	14100	8400	3500	461000	2.40	
## 31	7.1	3.9	12.8	9600	7100	1300	327000	5.46	
## 32	7.0	3.9	12.6	7600	5300	1300	249000	4.08	
## 33	7.7	4.4	15.3	8400	5400	1800	306000	3.00	
## 34	7.0	4.2	14.2	12500	10700	1300	234000	8.23	
##	PLR_pre	PNI_pre	ALI_pre	SII_pre	Prot_1C	Alb_1C	Hb_1C	Leucoc_1C	Neutr_1C
## 1	195.00	48.00	20.15	1560000.0	7.5	4.6	15.9	8100	6000
## 2	155.00	41.00	12.77	1271000.0	6.2	3.9	11.7	8500	7000
## 3	249.29	50.00	20.92	1844714.3	7.2	4.4	16.1	12000	8000
## 4	122.22	49.00	67.80	366666.7	7.0	3.9	15.5	7300	4400
## 5	152.17	55.50	55.48	806521.7	7.4	4.5	15.9	9200	4900
## 6	174.38	51.00	33.06	1046250.0	8.1	4.2	14.5	8000	5300
## 7	672.50	38.00	4.76	6052500.0	6.5	4.0	11.6	4500	3200
## 8	125.62	47.00	30.43	665812.5	7.7	4.0	12.1	3600	1900
## 9	166.84	46.50	11.89	2202315.8	7.0	3.9	14.5	16100	12100
## 10	231.43	47.00	9.89	3101142.9	7.4	4.0	13.5	12800	9800
## 11	326.00	42.50	15.81	2608000.0	7.0	3.4	11.4	12500	10400
## 12	86.40	49.50	46.16	432000.0	7.1	4.1	11.6	7600	3700
## 13	206.82	52.00	27.18	1799318.2	7.3	4.0	14.3	11300	6200
## 14	119.13	52.50	25.17	905391.3	7.0	3.7	10.8	15400	11300
## 15	71.72	58.50	29.95	674206.9	6.9	3.7	11.3	10400	8400
## 16	224.00	34.50	3.20	2755200.0	7.8	4.2	13.7	14100	9700
## 17	366.25	32.00	26.68	915625.0	4.6	3.1	8.4	4800	3300
## 18	132.59	52.50	34.05	941407.4	9.1	4.0	11.9	11100	6000
## 19	375.56	41.50	31.72	976444.4	6.0	3.1	11.1	3600	1700
## 20	928.00	45.50	4.73	8723200.0	8.0	4.5	11.9	10900	8900
## 21	131.67	49.00	20.30	961166.7	7.2	4.2	10.2	13800	9300
## 22	130.53	50.50	45.38	626526.3	7.3	3.8	12.5	7400	4600
## 23	478.57	33.50	8.71	2584285.7	6.8	3.6	16.0	4600	3000
## 24	71.50	49.00	56.32	250250.0	7.4	3.9	12.9	6000	3000
## 25	236.33	49.00	21.52	2056100.0	7.3	3.6	11.1	12600	7000
## 26	260.91	49.50	26.16	1121909.1	6.4	3.7	11.7	9000	6300
## 27	368.75	44.00	14.25	1770000.0	6.9	4.1	13.6	4900	2700
## 28	152.00	48.50	30.40	744800.0	7.1	4.0	11.7	6900	5900
## 29	82.12	51.50	26.69	944393.9	6.8	3.3	14.5	12800	6800
## 30	131.71	59.50	47.32	1106400.0	6.3	3.8	11.6	8700	4600

##	31	251.54	45.50	13.51	1785923.1	6.5	3.3	10.5	9000	6300
##	32	191.54	47.42	26.86	1015153.8	7.1	4.3	13.2	13100	8700
##	33	170.00	53.00	45.79	918000.0	7.0	3.9	15.6	10000	6800
##	34	180.00	48.50	18.18	1926000.0	6.6	4.1	15.4	10200	7600
##		Linf_1C	Plaq_1C	NLR_1C	PLR_1C	PNI_1C	SII_1C	Prot_2C	Alb_2C	Hb_2C
##	1	1600	369000	3.75	230.62	54.00	1383750.0	7.90	4.50	16.50
##	2	800	112000	8.75	140.00	43.00	980000.0	6.10	3.70	11.40
##	3	2500	382000	3.20	152.80	56.50	1222400.0	7.20	4.40	16.70
##	4	2000	266000	2.20	133.00	10.00	585200.0	7.20	4.10	15.30
##	5	3100	364000	1.58	117.42	60.50	575354.8	8.00	4.50	16.30
##	6	1600	293000	3.31	183.12	50.00	970562.5	8.10	4.50	15.50
##	7	900	218000	3.56	242.22	44.50	775111.1	7.20	4.10	12.40
##	8	900	131000	2.11	145.56	44.50	276555.6	7.22	4.01	13.15
##	9	2900	380000	4.17	131.03	53.50	1585517.2	7.22	3.70	13.40
##	10	1900	214000	5.16	112.63	49.50	1103789.5	7.50	4.10	13.40
##	11	1000	463000	10.40	463.00	39.00	4815200.0	6.50	2.90	8.60
##	12	2900	242000	1.28	83.45	55.50	308758.6	6.80	3.90	11.40
##	13	3600	452000	1.72	125.56	58.00	778444.4	7.10	4.20	15.00
##	14	2500	444000	4.52	177.60	49.50	2006880.0	7.20	4.00	10.90
##	15	1000	272000	8.40	272.00	42.00	2284800.0	7.20	3.90	11.20
##	16	3100	494000	3.13	159.35	57.50	1545741.9	7.20	3.90	13.30
##	17	700	226000	4.71	322.86	34.50	1065428.6	5.30	3.00	10.30
##	18	3000	380000	2.00	126.67	55.00	760000.0	8.80	4.10	12.90
##	19	1100	338000	1.55	307.27	36.50	522363.6	6.70	3.70	11.60
##	20	1200	465000	7.42	387.50	51.00	3448750.0	7.90	4.60	12.30
##	21	1600	205000	5.81	128.12	50.00	1191562.5	6.90	4.40	10.40
##	22	1800	304000	2.56	168.89	47.00	776888.9	7.70	3.90	13.40
##	23	1100	274000	2.73	249.09	41.50	747272.7	7.80	3.60	15.80
##	24	2200	277000	1.36	125.91	50.00	377727.3	7.40	4.00	13.20
##	25	3600	650000	1.94	180.56	54.00	1263888.9	7.50	3.80	11.40
##	26	800	457000	7.88	571.25	41.00	3598875.0	7.22	4.01	13.15
##	27	1600	192000	1.69	120.00	49.00	324000.0	6.80	4.20	13.50
##	28	600	259000	9.83	431.67	43.00	2546833.3	7.10	4.00	11.70
##	29	2800	366000	2.43	130.71	47.00	888857.1	7.50	3.70	13.80
##	30	2200	553000	2.09	251.36	49.00	1156272.7	6.30	4.10	13.00
##	31	1200	513000	5.25	427.50	39.00	2693250.0	7.22	4.01	11.40
##	32	2900	345000	3.00	118.97	57.50	1035000.0	7.50	4.50	12.90
##	33	2000	323000	3.40	161.50	47.33	1098200.0	7.70	4.30	15.70
##	34	1700	208000	4.47	122.35	49.50	929882.4	6.60	4.10	16.30
##		Leucoc_2C	Neutr_2C	Linf_2C	Plaq_2C	NLR_2C	NLR2C_corte4o5	PLR_2C	PNI_2C	
##	1	6400.00	3400.00	2300.00	253000.0	1.478261		0.00	110.00	56.50
##	2	8300.00	6200.00	1200.00	126000.0	5.166667		1.00	105.00	43.00
##	3	8400.00	4500.00	3000.00	329000.0	1.500000		0.00	109.67	59.00
##	4	7200.00	4300.00	1900.00	254000.0	2.263158		0.00	133.68	50.50
##	5	9200.00	5100.00	2800.00	313000.0	1.821429		0.00	111.79	59.00
##	6	8000.00	5300.00	1800.00	285.0	2.944444		0.00	0.16	54.00
##	7	8200.00	6400.00	1100.00	265000.0	5.818182		1.00	240.91	46.50
##	8	9106.25	5796.88	2090.62	305883.9	3.348731		0.25	177.59	50.71
##	9	19400.00	16100.00	2300.00	406000.0	7.000000		1.00	176.52	48.50
##	10	8700.00	5500.00	2200.00	226000.0	2.500000		0.00	102.73	52.00
##	11	14500.00	12100.00	1200.00	858000.0	10.083333		1.00	715.00	35.00
##	12	9400.00	4400.00	3100.00	260000.0	1.419355		0.00	83.87	54.50
##	13	10200.00	5100.00	2900.00	319000.0	1.758621		0.00	110.00	56.50
##	14	14700.00	9900.00	3500.00	408000.0	2.828571		0.00	116.57	57.50

## 15	10600.00	7900.00	1400.00	254000.0	5.642857	1.00	181.43	46.00
## 16	9900.00	6100.00	2500.00	297000.0	2.440000	0.00	118.80	51.50
## 17	3800.00	2300.00	800.00	218000.0	2.875000	0.00	272.50	34.00
## 18	11200.00	6100.00	3300.00	272000.0	1.848485	0.00	82.42	57.50
## 19	4000.00	1800.00	1500.00	394000.0	1.200000	0.00	262.67	44.50
## 20	8600.00	6300.00	1200.00	425000.0	5.250000	1.00	354.17	52.00
## 21	12500.00	8500.00	1500.00	217000.0	5.666667	1.00	144.67	51.50
## 22	6400.00	3600.00	1800.00	194000.0	2.000000	0.00	107.78	48.00
## 23	5400.00	3700.00	1000.00	237000.0	3.700000	0.00	237.00	41.00
## 24	5100.00	2400.00	2100.00	240000.0	1.142857	0.00	114.29	50.50
## 25	13900.00	7900.00	3900.00	615000.0	2.025641	0.00	157.69	57.50
## 26	9106.25	5796.88	2090.62	305883.9	3.348731	0.25	177.59	50.71
## 27	5300.00	3300.00	1400.00	235000.0	2.357143	0.00	167.86	49.00
## 28	6900.00	5900.00	600.00	259000.0	9.833333	1.00	431.67	43.00
## 29	12000.00	5700.00	4000.00	256000.0	1.425000	0.00	64.00	57.00
## 30	8300.00	4100.00	2900.00	343000.0	1.413793	0.00	118.28	55.50
## 31	7600.00	4900.00	1300.00	412000.0	3.769231	0.00	316.92	50.71
## 32	8800.00	5800.00	1800.00	337000.0	3.222222	0.00	187.22	54.00
## 33	9600.00	5900.00	2200.00	312000.0	2.681818	0.00	141.82	54.00
## 34	8900.00	5000.00	2400.00	254000.0	2.083333	0.00	105.83	53.00
##	SII_2C	Prot_1eval	Alb_1eval	Hb_1eval	Leucoc_1eval	Neutr_1eval	Linf_1eval	
## 1	374000.00	7.30	4.4	16.9	8000	5300	2100	
## 2	651000.00	6.00	3.9	11.2	7500	5500	1300	
## 3	493500.00	7.10	4.5	16.6	8700	4900	2900	
## 4	574842.11	7.20	4.0	15.7	7800	4400	2500	
## 5	570107.14	7.09	4.3	16.5	8700	4600	2800	
## 6	839.17	7.60	4.4	15.4	8000	5900	1400	
## 7	1541818.18	6.70	3.9	11.1	7600	6000	1000	
## 8	1134689.69	6.40	4.2	11.6	8600	6300	1400	
## 9	2842000.00	6.90	3.7	12.9	16200	13200	1900	
## 10	565000.00	7.70	4.5	14.6	7400	4700	1900	
## 11	8651500.00	7.00	3.1	8.8	10200	9100	500	
## 12	369032.26	7.20	4.0	10.8	11100	7200	2600	
## 13	561000.00	7.10	4.2	15.1	9100	5000	2900	
## 14	1154057.14	8.00	4.5	12.9	18400	13600	3200	
## 15	1433285.71	7.09	3.8	11.3	9400	7200	900	
## 16	724680.00	7.00	4.1	15.8	9800	6000	2600	
## 17	626750.00	5.50	3.6	13.7	4600	2600	1100	
## 18	502787.88	8.50	4.1	12.7	10200	5400	3100	
## 19	472800.00	7.10	3.9	12.2	3100	1400	1300	
## 20	2231250.00	7.30	3.7	12.5	9200	7100	1100	
## 21	1229666.67	6.40	3.9	9.9	10000	7500	800	
## 22	388000.00	7.70	3.9	13.4	6400	3600	1800	
## 23	876900.00	7.09	3.1	14.9	4500	2900	1000	
## 24	274285.71	7.70	4.1	13.8	4300	2200	1600	
## 25	1245769.23	7.80	3.9	11.8	13900	7700	4000	
## 26	1134689.69	6.40	3.7	11.7	9000	6300	800	
## 27	553928.57	7.00	4.2	13.8	4600	3100	1100	
## 28	2546833.33	7.10	4.0	11.7	6900	5900	600	
## 29	364800.00	6.90	3.9	14.5	9100	4200	3500	
## 30	484931.03	6.30	4.1	13.0	8300	4100	2900	
## 31	1552923.08	7.00	4.0	11.3	7300	4300	1800	
## 32	1085888.89	7.70	4.6	12.9	9700	6500	2100	
## 33	836727.27	7.50	4.3	15.9	8700	6000	1600	

## 34	529166.67	6.60	4.1	16.3	8900	5000	2400
##	Plaq_1eval	NLR_1eval	PLR_1eval	PNI_1eval	SII_1eval	N_ciclos	SLP SLP_cens
## 1	262000	2.52	124.76	54.5	661238.1	35	29.86 1
## 2	96000	4.23	73.85	45.5	406153.8	35	60.39 1
## 3	315000	1.69	108.62	59.5	532241.4	7	35.19 0
## 4	259000	1.76	103.60	52.5	455840.0	11	7.39 0
## 5	285000	1.64	101.79	57.0	468214.3	7	4.50 0
## 6	263000	4.21	187.86	51.0	1108357.1	4	41.49 1
## 7	229000	6.00	229.00	44.0	1374000.0	19	24.02 0
## 8	124000	4.50	88.57	49.0	558000.0	1	7.89 1
## 9	317000	6.95	166.84	46.5	2202315.8	3	1.81 0
## 10	219000	2.47	115.26	54.5	541736.8	26	19.94 1
## 11	527000	18.20	1054.00	33.5	9591400.0	3	1.87 0
## 12	227000	2.77	87.31	53.0	628615.4	5	40.74 1
## 13	308000	1.72	106.21	56.5	531034.5	11	7.46 0
## 14	328000	4.25	102.50	61.0	1394000.0	6	5.65 0
## 15	309000	8.00	343.33	42.5	2472000.0	6	3.68 0
## 16	205000	2.31	78.85	54.0	473076.9	32	22.37 0
## 17	220000	2.36	200.00	41.5	520000.0	25	21.98 1
## 18	282000	1.74	90.97	56.5	491225.8	35	46.32 1
## 19	290000	1.08	223.08	45.5	312307.7	16	35.32 0
## 20	400000	6.45	363.64	42.5	2581818.2	25	45.63 1
## 21	216000	9.38	270.00	43.0	2025000.0	3	1.91 0
## 22	194000	2.00	107.78	48.0	388000.0	5	7.26 0
## 23	220000	2.90	220.00	36.0	638000.0	8	9.33 1
## 24	234000	1.38	146.25	49.0	321750.0	8	7.13 0
## 25	647000	1.93	161.75	59.0	1245475.0	8	5.62 0
## 26	457000	7.88	571.25	41.0	3598875.0	1	0.76 0
## 27	257000	2.82	233.64	47.5	724272.7	35	31.15 1
## 28	259000	9.83	431.67	43.0	2546833.3	2	1.38 0
## 29	199000	1.20	56.86	56.5	238800.0	27	17.81 0
## 30	343000	1.41	118.28	55.5	484931.0	13	11.17 0
## 31	353000	2.39	196.11	49.0	843277.8	4	1.81 0
## 32	308000	3.10	146.67	56.5	953333.3	35	40.21 1
## 33	320000	3.75	200.00	51.0	1200000.0	23	16.16 0
## 34	254000	2.08	105.83	53.0	529166.7	35	39.46 1
##	SG SG_cens	Porcentaje_NA_Fila					
## 1	29.86	1	3.409091				
## 2	60.39	1	0.000000				
## 3	57.63	1	0.000000				
## 4	33.74	0	3.409091				
## 5	18.83	0	2.272727				
## 6	41.49	1	1.136364				
## 7	30.23	0	1.136364				
## 8	7.89	1	14.772727				
## 9	1.81	0	5.681818				
## 10	19.94	0	0.000000				
## 11	17.48	0	2.272727				
## 12	40.74	0	0.000000				
## 13	41.00	0	0.000000				
## 14	8.84	0	0.000000				
## 15	6.57	0	1.136364				
## 16	28.75	0	0.000000				
## 17	21.98	0	1.136364				

```
## 18 46.32      1      1.136364
## 19 35.32      1      0.000000
## 20 45.63      1      2.272727
## 21  5.75      0      0.000000
## 22 22.08      0      1.136364
## 23  9.33      0      2.272727
## 24 16.13      0      6.818182
## 25 10.61      0      0.000000
## 26 41.17      0     13.636364
## 27 31.15      1      0.000000
## 28  1.84      0      1.136364
## 29 26.58      0      0.000000
## 30 22.70      0      0.000000
## 31 16.00      0      5.681818
## 32 40.21      1     10.227273
## 33 35.78      0      4.545455
## 34 39.46      1      1.136364
```

3.2. Metodo Cart

```
imputed_data2 <- mice(datos %>%
  select(which(sapply(datos, is.factor))),
  method = "cart", print = FALSE)
```

```
## Warning: Number of logged events: 270
```

```
df_imputado2 <- complete(imputed_data2)
df_imputado2
```

```
##      Idpac Sexo ECOG p_peso_no_sí Hab_tabaq Diabetes Cardiop Enf_neurod
## 1  P_01     2     1      0          1      0      0      0
## 2  P_02     1     2      0          0      0      0      0
## 3  P_03     2     0      0          2      0      0      0
## 4  P_04     2     0      0          2      0      1      0
## 5  P_05     2     1      1          1      0      0      0
## 6  P_06     2     1      0          2      0      0      0
## 7  P_07     1     1      1          2      1      0      0
## 8  P_08     2     2      0          2      0      1      1
## 9  P_09     2     1      0          2      0      1      1
## 10 P_10     2     1      0          1      0      0      0
## 11 P_11     2     2      1          1      0      0      0
## 12 P_12     2     1      1          1      0      1      0
## 13 P_13     1     1      1          0      1      0      0
## 14 P_14     2     1      1          1      0      0      0
## 15 P_15     1     2      0          1      0      0      0
## 16 P_16     2     1      0          1      0      0      0
## 17 P_18     1     0      0          1      0      0      0
## 18 P_19     2     1      0          2      0      0      0
## 19 P_20     1     1      0          2      0      0      0
## 20 P_22     2     0      0          2      0      0      0
## 21 P_23     2     2      1          2      1      0      1
```

##	22	P_24	2	1	0	1	0	0	0
##	23	P_25	2	1	1	1	0	0	0
##	24	P_26	1	1	0	1	0	0	0
##	25	P_27	2	1	0	2	0	0	0
##	26	P_28	1	1	0	2	0	0	0
##	27	P_29	1	1	1	0	0	0	0
##	28	P_30	2	1	1	2	0	0	0
##	29	P_31	2	1	0	2	0	0	0
##	30	P_32	2	1	0	1	0	0	0
##	31	P_33	2	1	1	1	0	0	0
##	32	P_34	2	0	1	2	1	0	0
##	33	P_35	1	2	0	1	0	0	0
##	34	P_36	2	1	0	1	0	0	0
##		Histologia	Histología_num	Tamaño_tumor	Afectacion_ganglionar				
##	1	Adenocarcinoma	0	2b	2				
##	2	Otros	2	X	2				
##	3	Adenocarcinoma	0	2b	1				
##	4	Escamoso	1	4	2				
##	5	Escamoso	1	2a	2				
##	6	Otros	2	x	3				
##	7	Escamoso	1	3	2				
##	8	Escamoso	1	1c	2				
##	9	Escamoso	1	4	2				
##	10	Adenocarcinoma	0	3	2				
##	11	Adenocarcinoma	0	4	3				
##	12	Adenocarcinoma	0	2b	1				
##	13	Adenocarcinoma	0	4	3				
##	14	Otros	2	x	0				
##	15	Adenocarcinoma	0	4	0				
##	16	Adenocarcinoma	0	x	2				
##	17	Adenocarcinoma	0	2b	2				
##	18	Adenocarcinoma	0	4	3				
##	19	Ca. indiferenciado	2	4	2				
##	20	Adenocarcinoma	0	x	x				
##	21	Adenocarcinoma	0	3	0				
##	22	Escamoso	1	3	2				
##	23	Adenocarcinoma	0	3	3				
##	24	Adenocarcinoma	0	4	3				
##	25	Adenocarcinoma	0	4	3				
##	26	Adenocarcinoma	0	4	3				
##	27	Otros	2	4	3				
##	28	Adenocarcinoma	0	4	3				
##	29	Adenocarcinoma	0	4	2				
##	30	Adenocarcinoma	0	4	3				
##	31	Adenocarcinoma	0	4	2				
##	32	Otros	2	3	3				
##	33	Adenocarcinoma	0	4	2				
##	34	Adenocarcinoma	0	4	0				
##		Afectacion_metastasica	Estadio	Estadio_num	Estado_mut	Estatinas			
##	1	1c	IVB	4	0	0			
##	2	1c	IVB	4	0	0			
##	3	1c	IVB	4	0	1			
##	4	0	IIIB	3	0	1			
##	5	1a	IVA	4	0	0			

## 6	1c	IVB	4	0	0
## 7	1c	IVB	4	0	1
## 8	0	IIIA	3	0	0
## 9	0	IIIB	3	0	1
## 10	1c	IVB	4	0	0
## 11	1c	IVB	4	0	0
## 12	1b	IVA	4	0	1
## 13	1a	IVA	4	0	0
## 14	1b	IVA	4	0	0
## 15	1c	IVB	4	0	0
## 16	1b	IVB	4	0	0
## 17	1c	IVB	4	0	0
## 18	1c	IVB	4	0	0
## 19	1c	IVB	4	0	0
## 20	1a	IVA	4	0	0
## 21	1c	IVB	4	0	1
## 22	1c	IVB	4	0	0
## 23	1c	IVB	4	0	0
## 24	1b	IVB	4	EGFR	0
## 25	1c	IVB	4	0 0 (fibrato)	
## 26	1c	IVB	4	0	1
## 27	1b	IVB	4	0	0
## 28	1b	IVB	4	0	0
## 29	1a	IVA	4	0	0
## 30	1c	IVB	4	0	0
## 31	1b	IVB	4	0	0
## 32	0	IIIC	3	0	1
## 33	1c	IVB	4	0	1
## 34	1c	IVB	4	0	0
##	NLR1C_corte4	NLR1C_corte5	primera_eval_num	Mejor_resp_num	Toxicidad
## 1	0	0	2	0	1
## 2	1	1	1	0	1
## 3	0	0	1	1	1
## 4	0	0	2	2	0
## 5	0	0	2	2	0
## 6	0	0	1	0	1
## 7	0	0	2	1	1
## 8	0	0	1	1	1
## 9	1	0	2	2	0
## 10	1	1	1	1	1
## 11	1	1	3	3	0
## 12	0	0	2	1	0
## 13	0	0	1	1	1
## 14	1	0	2	2	0
## 15	1	1	2	2	1
## 16	0	0	1	1	1
## 17	1	0	2	1	0
## 18	0	0	1	1	1
## 19	0	0	1	1	1
## 20	1	1	1	0	1
## 21	1	1	3	3	0
## 22	0	0	2	1	1
## 23	0	0	1	1	0
## 24	0	0	2	2	0

## 25	0	0	1	1	0
## 26	1	1	3	3	0
## 27	0	0	1	0	1
## 28	1	1	3	3	0
## 29	0	0	3	2	1
## 30	0	0	2	1	1
## 31	0	0	3	3	0
## 32	0	0	1	1	0
## 33	0	0	1	1	0
## 34	1	0	2	1	1
##	Tipo_tox	Grado_tox	Interrupc_tto	Motivo_inter	
## 1	Miocarditis	3	1	Fin del tratamiento previsto	
## 2	Dermatitis	1	1	Fin del tratamiento previsto	
## 3	Hepatitis	3	1	Toxicidad	
## 4	0	0	1	Progresión	
## 5	0	0	1	Progresión	
## 6	Uveítis	2	1	Toxicidad	
## 7	Neumonitis	3	1	Toxicidad	
## 8	Hepatitis	3	1	Toxicidad	
## 9	0	0	1	Exitus	
## 10	Uveítis	1	1	Exitus (otra causa)	
## 11	0	0	1	Progresión	
## 12	Neumonitis	3	1	Toxicidad	
## 13	Queratitis/Dermatitis	1	1	Progresión	
## 14	0	0	1	Progresión	
## 15	Dermatitis	1	1	Progresión	
## 16	Artritis	1	1	Progresión	
## 17	0	0	1	Exitus (otra causa)	
## 18	Tiroiditis	1	1	Fin del tratamiento previsto	
## 19	Encefalitis	3	1	Toxicidad	
## 20	Neumonitis	2	1	Toxicidad	
## 21	0	0	1	Progresión	
## 22	Neumonitis	3	1	Toxicidad	
## 23	0	0	1	2º tumor/ hepatocarcinoma	
## 24	0	0	1	Progresión	
## 25	0	0	1	Progresión	
## 26	0	0	1	Progresión	
## 27	Tiroiditis	2	1	Fin del tratamiento previsto	
## 28	0	0	1	Progresión	
## 29	Dermatitis	3	1	Toxicidad	
## 30	Dermatitis	1	1	Progresión	
## 31	0	0	1	Progresión	
## 32	Dermatitis	0	1	Fin del tratamiento previsto	
## 33	Miocarditis	0	0	Progresión	
## 34	Artritis	G2	1	Fin del tratamiento previsto	
##	Progresion	segunda_eval	Exitus		
## 1	0	0	1		
## 2	0	0	0		
## 3	1	0	0		
## 4	1	1	1		
## 5	1	1	1		
## 6	0	0	0		
## 7	1	1	1		
## 8	0	0	1		


```
## 9      1      0      1
## 10     0      0      1
## 11     1      1      1
## 12     0      0      1
## 13     1      1      1
## 14     1      0      1
## 15     1      1      1
## 16     1      1      1
## 17     0      0      1
## 18     0      0      0
## 19     1      0      0
## 20     0      0      0
## 21     1      0      1
## 22     1      1      1
## 23     0      0      1
## 24     1      1      1
## 25     1      1      0
## 26     1      1      1
## 27     0      0      0
## 28     1      0      1
## 29     1      0      1
## 30     1      1      1
## 31     1      1      1
## 32     0      0      0
## 33     1      1      1
## 34     0      0      0
```

```
df_completo <- cbind(df_imputado1, df_imputado2)
df_completo
```

```
##      Edad_dx Anciano      IMC Porcentaje_perdpeso Exp_tab PD_L1 Col_total LDH
## 1         46         0 28.09                0.0458      20   100      217  163
## 2         68         0 35.38                0.0000        0    70      154  171
## 3         59         0 25.71                0.0000      45    50      146  197
## 4         72         1 28.25                0.0000     92     2      149  159
## 5         50         0 29.05                0.1000     55    60      210  198
## 6         71         1 28.83                0.0000    200    90      162  220
## 7         71         1 29.78                0.0760     60   100      142  198
## 8         79         1 25.85                0.0000     40    70      167  184
## 9         73         1 22.32                0.0458     50    90      153  176
## 10        69         0 23.66                0.0000     50    70      132  199
## 11        56         0 24.10                0.0458     36    80      113  190
## 12        66         0 24.95                0.2290     50    50      182  198
## 13        62         0 26.22                0.1380        0    80      242  184
## 14        64         0 20.29                0.1100     50   100      133  183
## 15        59         0 22.06                0.0000     25    95      215  204
## 16        60         0 24.62                0.0000     82    90      202  276
## 17        63         0 29.78                0.0000     60   100      262  201
## 18        82         1 22.96                0.0000    100    90      200  383
## 19        66         0 24.77                0.0000     50    60      199  382
## 20        50         0 20.70                0.0000     52   100      194  156
## 21        78         1 28.72                0.0670     75    70      176  175
## 22        73         1 27.96                0.0100     83    70      160  195
## 23        60         0 22.39                0.1300     40   100      111  326
```

## 24	70	1	25.27		0.0300	52	90	276	264
## 25	64	0	18.36		0.0310	40	70	155	1644
## 26	57	0	23.24		0.0400	30	70	197	279
## 27	71	1	21.37		0.0550	0	80	206	164
## 28	75	1	24.22		0.0880	25	100	208	222
## 29	68	0	26.57		0.0380	40	90	191	267
## 30	62	0	27.04		0.0000	60	90	201	248
## 31	51	0	18.92		0.0750	45	60	142	264
## 32	80	1	21.23		0.1590	40	80	187	264
## 33	65	0	31.22		0.0450	85	95	192	230
## 34	62	0	35.63		0.0000	25	70	397	201
##	Prot_tot	Albumina	Hb	Leucoc_tot	Neutrofilos	Linf_tot	Plaquetas	NLR_pre	
## 1	7.0	4.1	15.3	10100	8000	1400	273000	5.71	
## 2	6.0	3.7	11.8	9800	8200	800	124000	10.25	
## 3	7.2	4.3	16.7	9900	7400	1400	349000	5.29	
## 4	7.1	4.0	14.6	5600	3000	1800	220000	1.67	
## 5	7.9	4.4	15.7	8900	5300	2300	350000	2.30	
## 6	8.0	4.3	13.1	8800	6000	1600	279000	3.75	
## 7	6.4	3.6	11.5	10000	9000	400	269000	22.50	
## 8	7.0	3.9	12.3	7700	5300	1600	201000	3.31	
## 9	6.9	3.7	12.9	16200	13200	1900	317000	6.95	
## 10	7.3	4.0	13.0	15700	13400	1400	324000	9.57	
## 11	7.3	3.5	13.6	10600	8000	1500	489000	5.33	
## 12	6.5	3.7	11.4	8700	5000	2500	216000	2.00	
## 13	7.3	4.1	13.7	12600	8700	2200	455000	3.95	
## 14	7.0	4.1	13.7	11400	7600	2300	274000	3.30	
## 15	7.2	4.4	11.9	13400	9400	2900	208000	3.24	
## 16	5.6	3.2	15.5	13100	12300	500	112000	24.60	
## 17	5.4	2.8	11.6	4100	2500	800	293000	3.12	
## 18	8.9	3.9	12.1	11700	7100	2700	358000	2.63	
## 19	6.3	3.7	10.6	4000	2600	900	338000	2.89	
## 20	7.3	4.3	11.5	10800	9400	500	464000	18.80	
## 21	7.1	4.3	11.5	10600	7300	1200	158000	6.08	
## 22	7.3	4.1	12.6	7700	4800	1900	248000	2.53	
## 23	6.4	3.0	13.5	6800	5400	700	335000	7.71	
## 24	7.0	3.9	14.1	6000	3500	2000	143000	1.75	
## 25	6.9	3.4	10.9	13800	8700	3000	709000	2.90	
## 26	6.9	4.4	13.0	6300	4300	1100	287000	3.91	
## 27	6.6	4.0	14.0	6200	4800	800	295000	6.00	
## 28	7.4	4.1	12.4	7200	4900	1500	228000	3.27	
## 29	6.9	3.5	14.5	20500	11500	3300	271000	3.48	
## 30	7.3	4.2	13.6	14100	8400	3500	461000	2.40	
## 31	7.1	3.9	12.8	9600	7100	1300	327000	5.46	
## 32	7.0	3.9	12.6	7600	5300	1300	249000	4.08	
## 33	7.7	4.4	15.3	8400	5400	1800	306000	3.00	
## 34	7.0	4.2	14.2	12500	10700	1300	234000	8.23	
##	PLR_pre	PNI_pre	ALI_pre	SII_pre	Prot_1C	Alb_1C	Hb_1C	Leucoc_1C	Neutr_1C
## 1	195.00	48.00	20.15	1560000.0	7.5	4.6	15.9	8100	6000
## 2	155.00	41.00	12.77	1271000.0	6.2	3.9	11.7	8500	7000
## 3	249.29	50.00	20.92	1844714.3	7.2	4.4	16.1	12000	8000
## 4	122.22	49.00	67.80	366666.7	7.0	3.9	15.5	7300	4400
## 5	152.17	55.50	55.48	806521.7	7.4	4.5	15.9	9200	4900
## 6	174.38	51.00	33.06	1046250.0	8.1	4.2	14.5	8000	5300
## 7	672.50	38.00	4.76	6052500.0	6.5	4.0	11.6	4500	3200

## 8	125.62	47.00	30.43	665812.5	7.7	4.0	12.1	3600	1900
## 9	166.84	46.50	11.89	2202315.8	7.0	3.9	14.5	16100	12100
## 10	231.43	47.00	9.89	3101142.9	7.4	4.0	13.5	12800	9800
## 11	326.00	42.50	15.81	2608000.0	7.0	3.4	11.4	12500	10400
## 12	86.40	49.50	46.16	432000.0	7.1	4.1	11.6	7600	3700
## 13	206.82	52.00	27.18	1799318.2	7.3	4.0	14.3	11300	6200
## 14	119.13	52.50	25.17	905391.3	7.0	3.7	10.8	15400	11300
## 15	71.72	58.50	29.95	674206.9	6.9	3.7	11.3	10400	8400
## 16	224.00	34.50	3.20	2755200.0	7.8	4.2	13.7	14100	9700
## 17	366.25	32.00	26.68	915625.0	4.6	3.1	8.4	4800	3300
## 18	132.59	52.50	34.05	941407.4	9.1	4.0	11.9	11100	6000
## 19	375.56	41.50	31.72	976444.4	6.0	3.1	11.1	3600	1700
## 20	928.00	45.50	4.73	8723200.0	8.0	4.5	11.9	10900	8900
## 21	131.67	49.00	20.30	961166.7	7.2	4.2	10.2	13800	9300
## 22	130.53	50.50	45.38	626526.3	7.3	3.8	12.5	7400	4600
## 23	478.57	33.50	8.71	2584285.7	6.8	3.6	16.0	4600	3000
## 24	71.50	49.00	56.32	250250.0	7.4	3.9	12.9	6000	3000
## 25	236.33	49.00	21.52	2056100.0	7.3	3.6	11.1	12600	7000
## 26	260.91	49.50	26.16	1121909.1	6.4	3.7	11.7	9000	6300
## 27	368.75	44.00	14.25	1770000.0	6.9	4.1	13.6	4900	2700
## 28	152.00	48.50	30.40	744800.0	7.1	4.0	11.7	6900	5900
## 29	82.12	51.50	26.69	944393.9	6.8	3.3	14.5	12800	6800
## 30	131.71	59.50	47.32	1106400.0	6.3	3.8	11.6	8700	4600
## 31	251.54	45.50	13.51	1785923.1	6.5	3.3	10.5	9000	6300
## 32	191.54	47.42	26.86	1015153.8	7.1	4.3	13.2	13100	8700
## 33	170.00	53.00	45.79	918000.0	7.0	3.9	15.6	10000	6800
## 34	180.00	48.50	18.18	1926000.0	6.6	4.1	15.4	10200	7600
##	Linf_1C	Plaq_1C	NLR_1C	PLR_1C	PNI_1C	SII_1C	Prot_2C	Alb_2C	Hb_2C
## 1	1600	369000	3.75	230.62	54.00	1383750.0	7.90	4.50	16.50
## 2	800	112000	8.75	140.00	43.00	980000.0	6.10	3.70	11.40
## 3	2500	382000	3.20	152.80	56.50	1222400.0	7.20	4.40	16.70
## 4	2000	266000	2.20	133.00	10.00	585200.0	7.20	4.10	15.30
## 5	3100	364000	1.58	117.42	60.50	575354.8	8.00	4.50	16.30
## 6	1600	293000	3.31	183.12	50.00	970562.5	8.10	4.50	15.50
## 7	900	218000	3.56	242.22	44.50	775111.1	7.20	4.10	12.40
## 8	900	131000	2.11	145.56	44.50	276555.6	7.22	4.01	13.15
## 9	2900	380000	4.17	131.03	53.50	1585517.2	7.22	3.70	13.40
## 10	1900	214000	5.16	112.63	49.50	1103789.5	7.50	4.10	13.40
## 11	1000	463000	10.40	463.00	39.00	4815200.0	6.50	2.90	8.60
## 12	2900	242000	1.28	83.45	55.50	308758.6	6.80	3.90	11.40
## 13	3600	452000	1.72	125.56	58.00	778444.4	7.10	4.20	15.00
## 14	2500	444000	4.52	177.60	49.50	2006880.0	7.20	4.00	10.90
## 15	1000	272000	8.40	272.00	42.00	2284800.0	7.20	3.90	11.20
## 16	3100	494000	3.13	159.35	57.50	1545741.9	7.20	3.90	13.30
## 17	700	226000	4.71	322.86	34.50	1065428.6	5.30	3.00	10.30
## 18	3000	380000	2.00	126.67	55.00	760000.0	8.80	4.10	12.90
## 19	1100	338000	1.55	307.27	36.50	522363.6	6.70	3.70	11.60
## 20	1200	465000	7.42	387.50	51.00	3448750.0	7.90	4.60	12.30
## 21	1600	205000	5.81	128.12	50.00	1191562.5	6.90	4.40	10.40
## 22	1800	304000	2.56	168.89	47.00	776888.9	7.70	3.90	13.40
## 23	1100	274000	2.73	249.09	41.50	747272.7	7.80	3.60	15.80
## 24	2200	277000	1.36	125.91	50.00	377727.3	7.40	4.00	13.20
## 25	3600	650000	1.94	180.56	54.00	1263888.9	7.50	3.80	11.40
## 26	800	457000	7.88	571.25	41.00	3598875.0	7.22	4.01	13.15

## 27	1600	192000	1.69	120.00	49.00	324000.0	6.80	4.20	13.50
## 28	600	259000	9.83	431.67	43.00	2546833.3	7.10	4.00	11.70
## 29	2800	366000	2.43	130.71	47.00	888857.1	7.50	3.70	13.80
## 30	2200	553000	2.09	251.36	49.00	1156272.7	6.30	4.10	13.00
## 31	1200	513000	5.25	427.50	39.00	2693250.0	7.22	4.01	11.40
## 32	2900	345000	3.00	118.97	57.50	1035000.0	7.50	4.50	12.90
## 33	2000	323000	3.40	161.50	47.33	1098200.0	7.70	4.30	15.70
## 34	1700	208000	4.47	122.35	49.50	929882.4	6.60	4.10	16.30
##	Leucoc_2C	Neutr_2C	Linf_2C	Plaq_2C	NLR_2C	NLR2C_corte4o5	PLR_2C	PNI_2C	
## 1	6400.00	3400.00	2300.00	253000.0	1.478261		0.00	110.00	56.50
## 2	8300.00	6200.00	1200.00	126000.0	5.166667		1.00	105.00	43.00
## 3	8400.00	4500.00	3000.00	329000.0	1.500000		0.00	109.67	59.00
## 4	7200.00	4300.00	1900.00	254000.0	2.263158		0.00	133.68	50.50
## 5	9200.00	5100.00	2800.00	313000.0	1.821429		0.00	111.79	59.00
## 6	8000.00	5300.00	1800.00	285.0	2.944444		0.00	0.16	54.00
## 7	8200.00	6400.00	1100.00	265000.0	5.818182		1.00	240.91	46.50
## 8	9106.25	5796.88	2090.62	305883.9	3.348731		0.25	177.59	50.71
## 9	19400.00	16100.00	2300.00	406000.0	7.000000		1.00	176.52	48.50
## 10	8700.00	5500.00	2200.00	226000.0	2.500000		0.00	102.73	52.00
## 11	14500.00	12100.00	1200.00	858000.0	10.083333		1.00	715.00	35.00
## 12	9400.00	4400.00	3100.00	260000.0	1.419355		0.00	83.87	54.50
## 13	10200.00	5100.00	2900.00	319000.0	1.758621		0.00	110.00	56.50
## 14	14700.00	9900.00	3500.00	408000.0	2.828571		0.00	116.57	57.50
## 15	10600.00	7900.00	1400.00	254000.0	5.642857		1.00	181.43	46.00
## 16	9900.00	6100.00	2500.00	297000.0	2.440000		0.00	118.80	51.50
## 17	3800.00	2300.00	800.00	218000.0	2.875000		0.00	272.50	34.00
## 18	11200.00	6100.00	3300.00	272000.0	1.848485		0.00	82.42	57.50
## 19	4000.00	1800.00	1500.00	394000.0	1.200000		0.00	262.67	44.50
## 20	8600.00	6300.00	1200.00	425000.0	5.250000		1.00	354.17	52.00
## 21	12500.00	8500.00	1500.00	217000.0	5.666667		1.00	144.67	51.50
## 22	6400.00	3600.00	1800.00	194000.0	2.000000		0.00	107.78	48.00
## 23	5400.00	3700.00	1000.00	237000.0	3.700000		0.00	237.00	41.00
## 24	5100.00	2400.00	2100.00	240000.0	1.142857		0.00	114.29	50.50
## 25	13900.00	7900.00	3900.00	615000.0	2.025641		0.00	157.69	57.50
## 26	9106.25	5796.88	2090.62	305883.9	3.348731		0.25	177.59	50.71
## 27	5300.00	3300.00	1400.00	235000.0	2.357143		0.00	167.86	49.00
## 28	6900.00	5900.00	600.00	259000.0	9.833333		1.00	431.67	43.00
## 29	12000.00	5700.00	4000.00	256000.0	1.425000		0.00	64.00	57.00
## 30	8300.00	4100.00	2900.00	343000.0	1.413793		0.00	118.28	55.50
## 31	7600.00	4900.00	1300.00	412000.0	3.769231		0.00	316.92	50.71
## 32	8800.00	5800.00	1800.00	337000.0	3.222222		0.00	187.22	54.00
## 33	9600.00	5900.00	2200.00	312000.0	2.681818		0.00	141.82	54.00
## 34	8900.00	5000.00	2400.00	254000.0	2.083333		0.00	105.83	53.00
##	SII_2C	Prot_1eval	Alb_1eval	Hb_1eval	Leucoc_1eval	Neutr_1eval	Linf_1eval		
## 1	374000.00	7.30	4.4	16.9	8000	5300	2100		
## 2	651000.00	6.00	3.9	11.2	7500	5500	1300		
## 3	493500.00	7.10	4.5	16.6	8700	4900	2900		
## 4	574842.11	7.20	4.0	15.7	7800	4400	2500		
## 5	570107.14	7.09	4.3	16.5	8700	4600	2800		
## 6	839.17	7.60	4.4	15.4	8000	5900	1400		
## 7	1541818.18	6.70	3.9	11.1	7600	6000	1000		
## 8	1134689.69	6.40	4.2	11.6	8600	6300	1400		
## 9	2842000.00	6.90	3.7	12.9	16200	13200	1900		
## 10	565000.00	7.70	4.5	14.6	7400	4700	1900		

## 11	8651500.00	7.00	3.1	8.8	10200	9100	500
## 12	369032.26	7.20	4.0	10.8	11100	7200	2600
## 13	561000.00	7.10	4.2	15.1	9100	5000	2900
## 14	1154057.14	8.00	4.5	12.9	18400	13600	3200
## 15	1433285.71	7.09	3.8	11.3	9400	7200	900
## 16	724680.00	7.00	4.1	15.8	9800	6000	2600
## 17	626750.00	5.50	3.6	13.7	4600	2600	1100
## 18	502787.88	8.50	4.1	12.7	10200	5400	3100
## 19	472800.00	7.10	3.9	12.2	3100	1400	1300
## 20	2231250.00	7.30	3.7	12.5	9200	7100	1100
## 21	1229666.67	6.40	3.9	9.9	10000	7500	800
## 22	388000.00	7.70	3.9	13.4	6400	3600	1800
## 23	876900.00	7.09	3.1	14.9	4500	2900	1000
## 24	274285.71	7.70	4.1	13.8	4300	2200	1600
## 25	1245769.23	7.80	3.9	11.8	13900	7700	4000
## 26	1134689.69	6.40	3.7	11.7	9000	6300	800
## 27	553928.57	7.00	4.2	13.8	4600	3100	1100
## 28	2546833.33	7.10	4.0	11.7	6900	5900	600
## 29	364800.00	6.90	3.9	14.5	9100	4200	3500
## 30	484931.03	6.30	4.1	13.0	8300	4100	2900
## 31	1552923.08	7.00	4.0	11.3	7300	4300	1800
## 32	1085888.89	7.70	4.6	12.9	9700	6500	2100
## 33	836727.27	7.50	4.3	15.9	8700	6000	1600
## 34	529166.67	6.60	4.1	16.3	8900	5000	2400

##	Plaq_1eval	NLR_1eval	PLR_1eval	PNI_1eval	SII_1eval	N_ciclos	SLP	SLP_cens
## 1	262000	2.52	124.76	54.5	661238.1	35	29.86	1
## 2	96000	4.23	73.85	45.5	406153.8	35	60.39	1
## 3	315000	1.69	108.62	59.5	532241.4	7	35.19	0
## 4	259000	1.76	103.60	52.5	455840.0	11	7.39	0
## 5	285000	1.64	101.79	57.0	468214.3	7	4.50	0
## 6	263000	4.21	187.86	51.0	1108357.1	4	41.49	1
## 7	229000	6.00	229.00	44.0	1374000.0	19	24.02	0
## 8	124000	4.50	88.57	49.0	558000.0	1	7.89	1
## 9	317000	6.95	166.84	46.5	2202315.8	3	1.81	0
## 10	219000	2.47	115.26	54.5	541736.8	26	19.94	1
## 11	527000	18.20	1054.00	33.5	9591400.0	3	1.87	0
## 12	227000	2.77	87.31	53.0	628615.4	5	40.74	1
## 13	308000	1.72	106.21	56.5	531034.5	11	7.46	0
## 14	328000	4.25	102.50	61.0	1394000.0	6	5.65	0
## 15	309000	8.00	343.33	42.5	2472000.0	6	3.68	0
## 16	205000	2.31	78.85	54.0	473076.9	32	22.37	0
## 17	220000	2.36	200.00	41.5	520000.0	25	21.98	1
## 18	282000	1.74	90.97	56.5	491225.8	35	46.32	1
## 19	290000	1.08	223.08	45.5	312307.7	16	35.32	0
## 20	400000	6.45	363.64	42.5	2581818.2	25	45.63	1
## 21	216000	9.38	270.00	43.0	2025000.0	3	1.91	0
## 22	194000	2.00	107.78	48.0	388000.0	5	7.26	0
## 23	220000	2.90	220.00	36.0	638000.0	8	9.33	1
## 24	234000	1.38	146.25	49.0	321750.0	8	7.13	0
## 25	647000	1.93	161.75	59.0	1245475.0	8	5.62	0
## 26	457000	7.88	571.25	41.0	3598875.0	1	0.76	0
## 27	257000	2.82	233.64	47.5	724272.7	35	31.15	1
## 28	259000	9.83	431.67	43.0	2546833.3	2	1.38	0
## 29	199000	1.20	56.86	56.5	238800.0	27	17.81	0

## 30	343000	1.41	118.28	55.5	484931.0	13	11.17	0
## 31	353000	2.39	196.11	49.0	843277.8	4	1.81	0
## 32	308000	3.10	146.67	56.5	953333.3	35	40.21	1
## 33	320000	3.75	200.00	51.0	1200000.0	23	16.16	0
## 34	254000	2.08	105.83	53.0	529166.7	35	39.46	1
##	SG	SG_cens	Porcentaje_NA_Fila	Idpac	Sexo	ECOG	p_peso_no_si	Hab_tabaq
## 1	29.86	1	3.409091	P_01	2	1	0	1
## 2	60.39	1	0.000000	P_02	1	2	0	0
## 3	57.63	1	0.000000	P_03	2	0	0	2
## 4	33.74	0	3.409091	P_04	2	0	0	2
## 5	18.83	0	2.272727	P_05	2	1	1	1
## 6	41.49	1	1.136364	P_06	2	1	0	2
## 7	30.23	0	1.136364	P_07	1	1	1	2
## 8	7.89	1	14.772727	P_08	2	2	0	2
## 9	1.81	0	5.681818	P_09	2	1	0	2
## 10	19.94	0	0.000000	P_10	2	1	0	1
## 11	17.48	0	2.272727	P_11	2	2	1	1
## 12	40.74	0	0.000000	P_12	2	1	1	1
## 13	41.00	0	0.000000	P_13	1	1	1	0
## 14	8.84	0	0.000000	P_14	2	1	1	1
## 15	6.57	0	1.136364	P_15	1	2	0	1
## 16	28.75	0	0.000000	P_16	2	1	0	1
## 17	21.98	0	1.136364	P_18	1	0	0	1
## 18	46.32	1	1.136364	P_19	2	1	0	2
## 19	35.32	1	0.000000	P_20	1	1	0	2
## 20	45.63	1	2.272727	P_22	2	0	0	2
## 21	5.75	0	0.000000	P_23	2	2	1	2
## 22	22.08	0	1.136364	P_24	2	1	0	1
## 23	9.33	0	2.272727	P_25	2	1	1	1
## 24	16.13	0	6.818182	P_26	1	1	0	1
## 25	10.61	0	0.000000	P_27	2	1	0	2
## 26	41.17	0	13.636364	P_28	1	1	0	2
## 27	31.15	1	0.000000	P_29	1	1	1	0
## 28	1.84	0	1.136364	P_30	2	1	1	2
## 29	26.58	0	0.000000	P_31	2	1	0	2
## 30	22.70	0	0.000000	P_32	2	1	0	1
## 31	16.00	0	5.681818	P_33	2	1	1	1
## 32	40.21	1	10.227273	P_34	2	0	1	2
## 33	35.78	0	4.545455	P_35	1	2	0	1
## 34	39.46	1	1.136364	P_36	2	1	0	1
##	Diabetes	Cardiop	Enf_neurod	Histologia	Histología_num	Tamaño_tumor		
## 1	0	0	0	Adenocarcinoma	0	2b		
## 2	0	0	0	Otros	2	X		
## 3	0	0	0	Adenocarcinoma	0	2b		
## 4	0	1	0	Escamoso	1	4		
## 5	0	0	0	Escamoso	1	2a		
## 6	0	0	0	Otros	2	x		
## 7	1	0	0	Escamoso	1	3		
## 8	0	1	1	Escamoso	1	1c		
## 9	0	1	1	Escamoso	1	4		
## 10	0	0	0	Adenocarcinoma	0	3		
## 11	0	0	0	Adenocarcinoma	0	4		
## 12	0	1	0	Adenocarcinoma	0	2b		
## 13	1	0	0	Adenocarcinoma	0	4		

## 14	0	0	0	Otros	2	x
## 15	0	0	0	Adenocarcinoma	0	4
## 16	0	0	0	Adenocarcinoma	0	x
## 17	0	0	0	Adenocarcinoma	0	2b
## 18	0	0	0	Adenocarcinoma	0	4
## 19	0	0	0	Ca. indiferenciado	2	4
## 20	0	0	0	Adenocarcinoma	0	x
## 21	1	0	1	Adenocarcinoma	0	3
## 22	0	0	0	Escamoso	1	3
## 23	0	0	0	Adenocarcinoma	0	3
## 24	0	0	0	Adenocarcinoma	0	4
## 25	0	0	0	Adenocarcinoma	0	4
## 26	0	0	0	Adenocarcinoma	0	4
## 27	0	0	0	Otros	2	4
## 28	0	0	0	Adenocarcinoma	0	4
## 29	0	0	0	Adenocarcinoma	0	4
## 30	0	0	0	Adenocarcinoma	0	4
## 31	0	0	0	Adenocarcinoma	0	4
## 32	1	0	0	Otros	2	3
## 33	0	0	0	Adenocarcinoma	0	4
## 34	0	0	0	Adenocarcinoma	0	4
##	Afectacion_ganglionar	Afectacion_metastasica	Estadio	Estadio_num	Estado_mut	
## 1		2	1c	IVB	4	0
## 2		2	1c	IVB	4	0
## 3		1	1c	IVB	4	0
## 4		2	0	IIIB	3	0
## 5		2	1a	IVA	4	0
## 6		3	1c	IVB	4	0
## 7		2	1c	IVB	4	0
## 8		2	0	IIIA	3	0
## 9		2	0	IIIB	3	0
## 10		2	1c	IVB	4	0
## 11		3	1c	IVB	4	0
## 12		1	1b	IVA	4	0
## 13		3	1a	IVA	4	0
## 14		0	1b	IVA	4	0
## 15		0	1c	IVB	4	0
## 16		2	1b	IVB	4	0
## 17		2	1c	IVB	4	0
## 18		3	1c	IVB	4	0
## 19		2	1c	IVB	4	0
## 20		x	1a	IVA	4	0
## 21		0	1c	IVB	4	0
## 22		2	1c	IVB	4	0
## 23		3	1c	IVB	4	0
## 24		3	1b	IVB	4	EGFR
## 25		3	1c	IVB	4	0
## 26		3	1c	IVB	4	0
## 27		3	1b	IVB	4	0
## 28		3	1b	IVB	4	0
## 29		2	1a	IVA	4	0
## 30		3	1c	IVB	4	0
## 31		2	1b	IVB	4	0
## 32		3	0	IIIC	3	0

## 33		2		1c	IVB	4	0
## 34		0		1c	IVB	4	0
##	Estatinas	NLR1C_corte4	NLR1C_corte5	primera_eval_num	Mejor_resp_num		
## 1	0	0	0	2	0		
## 2	0	1	1	1	0		
## 3	1	0	0	1	1		
## 4	1	0	0	2	2		
## 5	0	0	0	2	2		
## 6	0	0	0	1	0		
## 7	1	0	0	2	1		
## 8	0	0	0	1	1		
## 9	1	1	0	2	2		
## 10	0	1	1	1	1		
## 11	0	1	1	3	3		
## 12	1	0	0	2	1		
## 13	0	0	0	1	1		
## 14	0	1	0	2	2		
## 15	0	1	1	2	2		
## 16	0	0	0	1	1		
## 17	0	1	0	2	1		
## 18	0	0	0	1	1		
## 19	0	0	0	1	1		
## 20	0	1	1	1	0		
## 21	1	1	1	3	3		
## 22	0	0	0	2	1		
## 23	0	0	0	1	1		
## 24	0	0	0	2	2		
## 25	0 (fibrato)	0	0	1	1		
## 26	1	1	1	3	3		
## 27	0	0	0	1	0		
## 28	0	1	1	3	3		
## 29	0	0	0	3	2		
## 30	0	0	0	2	1		
## 31	0	0	0	3	3		
## 32	1	0	0	1	1		
## 33	1	0	0	1	1		
## 34	0	1	0	2	1		
##	Toxicidad	Tipo_tox	Grado_tox	Interrupc_tto			
## 1	1	Miocarditis	3	1			
## 2	1	Dermatitis	1	1			
## 3	1	Hepatitis	3	1			
## 4	0	0	0	1			
## 5	0	0	0	1			
## 6	1	Uveítis	2	1			
## 7	1	Neumonitis	3	1			
## 8	1	Hepatitis	3	1			
## 9	0	0	0	1			
## 10	1	Uveítis	1	1			
## 11	0	0	0	1			
## 12	0	Neumonitis	3	1			
## 13	1	Queratitis/Dermatitis	1	1			
## 14	0	0	0	1			
## 15	1	Dermatitis	1	1			
## 16	1	Artritis	1	1			

## 17	0	0	0	1
## 18	1	Tiroiditis	1	1
## 19	1	Encefalitis	3	1
## 20	1	Neumonitis	2	1
## 21	0	0	0	1
## 22	1	Neumonitis	3	1
## 23	0	0	0	1
## 24	0	0	0	1
## 25	0	0	0	1
## 26	0	0	0	1
## 27	1	Tiroiditis	2	1
## 28	0	0	0	1
## 29	1	Dermatitis	3	1
## 30	1	Dermatitis	1	1
## 31	0	0	0	1
## 32	0	Dermatitis	0	1
## 33	0	Miocarditis	0	0
## 34	1	Artritis	G2	1
##		Motivo_inter Progresion	segunda_eval	Exitus
## 1	Fin del tratamiento previsto	0	0	1
## 2	Fin del tratamiento previsto	0	0	0
## 3	Toxicidad	1	0	0
## 4	Progresión	1	1	1
## 5	Progresión	1	1	1
## 6	Toxicidad	0	0	0
## 7	Toxicidad	1	1	1
## 8	Toxicidad	0	0	1
## 9	Exitus	1	0	1
## 10	Exitus (otra causa)	0	0	1
## 11	Progresión	1	1	1
## 12	Toxicidad	0	0	1
## 13	Progresión	1	1	1
## 14	Progresión	1	0	1
## 15	Progresión	1	1	1
## 16	Progresión	1	1	1
## 17	Exitus (otra causa)	0	0	1
## 18	Fin del tratamiento previsto	0	0	0
## 19	Toxicidad	1	0	0
## 20	Toxicidad	0	0	0
## 21	Progresión	1	0	1
## 22	Toxicidad	1	1	1
## 23	2º tumor/ hepatocarcinoma	0	0	1
## 24	Progresión	1	1	1
## 25	Progresión	1	1	0
## 26	Progresión	1	1	1
## 27	Fin del tratamiento previsto	0	0	0
## 28	Progresión	1	0	1
## 29	Toxicidad	1	0	1
## 30	Progresión	1	1	1
## 31	Progresión	1	1	1
## 32	Fin del tratamiento previsto	0	0	0
## 33	Progresión	1	1	1
## 34	Fin del tratamiento previsto	0	0	0

3.3. missForest

Por último vamos a usar el método missForest, esta técnica de imputación basada en **bosques aleatorios** (Random Forest) permite estimar valores faltantes en datasets de manera no paramétrica y eficiente. Su funcionamiento se basa en entrenar un modelo de Random Forest para cada variable con valores ausentes, utilizando el resto de las variables como predictores.

La idea es imputar con missForest los valores numéricos y los categóricos con el método anterior (Cart), crear dos conjuntos diferentes es por probar si existen diferencias entre los dos métodos de imputación y usar el que mejor resultado genere.

```
datos_num <- datos %>% select(where(is.numeric))
datos_num <- as.data.frame(datos_num)

imputed_num <- missForest(
  xmis = datos_num, # Dataset con valores faltantes
  maxiter = 20,     # Máximo número de iteraciones
  ntree = 100,      # Número de árboles en cada Random Forest
  mtry = floor(ncol(datos_num) / 3), # Número de variables usadas en cada árbol
  replace = TRUE,   # Si se reemplazan las muestras en cada árbol
  decreasing = TRUE, # Imputar primero las variables con menos NA
  verbose = TRUE,   # Mostrar el progreso de la imputación
  variablewise = FALSE, # Imputación por bloque o variable por variable
)
```

```
## missForest iteration 1 in progress...
```

```
## Warning in randomForest.default(x = obsX, y = obsY, ntree = ntree, mtry = mtry,
## : The response has five or fewer unique values. Are you sure you want to do
## regression?
```

```
## done!
## estimated error(s): 0.3070776
## difference(s): 0.001737426
## time: 0.37 seconds
##
## missForest iteration 2 in progress...
```

```
## Warning in randomForest.default(x = obsX, y = obsY, ntree = ntree, mtry = mtry,
## : The response has five or fewer unique values. Are you sure you want to do
## regression?
```

```
## done!
## estimated error(s): 0.3156882
## difference(s): 0.0004710156
## time: 0.379 seconds
##
## missForest iteration 3 in progress...
```

```
## Warning in randomForest.default(x = obsX, y = obsY, ntree = ntree, mtry = mtry,
## : The response has five or fewer unique values. Are you sure you want to do
## regression?
```

```
## done!
##      estimated error(s): 0.3103979
##      difference(s): 3.731954e-05
##      time: 0.365 seconds
##
## missForest iteration 4 in progress...

## Warning in randomForest.default(x = obsX, y = obsY, ntree = ntree, mtry = mtry,
## : The response has five or fewer unique values. Are you sure you want to do
## regression?

## done!
##      estimated error(s): 0.3100008
##      difference(s): 0.0004069164
##      time: 0.367 seconds
```

```
imputed_num$OOBerror
```

```
##      NRMSE
## 0.3103979
```

```
df_completo_missForest <- cbind(df_imputado2, imputed_num$ximp)
```

El error **NRMSE** se interpreta como:

NRMSE	Calidad de Imputación
0 - 0.2	Excelente, la imputación es casi perfecta.
0.2 - 0.4	Buena, la imputación introduce un pequeño sesgo.
0.4 - 0.6	Moderada, puede haber errores significativos en la imputación.
0.6 - 1	Mala, la imputación se aleja demasiado de los valores reales.
< 1	Inaceptable, los datos imputados no son confiables.

Exportar los datos completos

```
# Cargar la librería
library(openxlsx)
```

```
## Warning: package 'openxlsx' was built under R version 4.3.3
```

```
# Guardar en formato Excel
write.xlsx(datos_limpios, "datos_limpios.xlsx", rowNames = FALSE)
write.xlsx(df_completo_missForest, "df_completo_missForest.xlsx", rowNames = FALSE)
```