

PEC 1

1. Seleccionar un dataset de metabolómica

En mi caso, he seleccionado y descargado el dataset Cachexia procedente de <https://github.com/nutrimetabolomics/metaboData/>. Para trabajar con estos datos en R, cargamos el archivo “human_cachexia.csv” que tiene los datos con los que trabajaremos.

```
# Indicar la ruta del archivo
ruta_archivo <- "D:/Máster bioinformática/Análisis de datos ómicos/PEC1/human_cachexia.csv"

# Cargar el archivo
cachexia <- read.csv(ruta_archivo)
```

2. Crear un contenedor del tipo SummarizedExperiment

Para crear el contenedor, primero comprobaremos las columnas o variables de nuestro dataset, con el fin de identificar cuales corresponden a datos y cuales a metadatos, y así poder asignarlos adecuadamente.

```
# Obtener el nombre de las columnas
colnames(cachexia)

## [1] "Patient.ID" "Muscle.loss"
## [3] "X1.6.Anhydro.beta.D.glucose" "X1.Methylnicotinamide"
## [5] "X2.Aminobutyrate" "X2.Hydroxyisobutyrate"
## [7] "X2.Oxoglutarate" "X3.Aminoisobutyrate"
## [9] "X3.Hydroxybutyrate" "X3.Hydroxyisovalerate"
## [11] "X3.Indoxylsulfate" "X4.Hydroxyphenylacetate"
## [13] "Acetate" "Acetone"
## [15] "Adipate" "Alanine"
## [17] "Asparagine" "Betaine"
## [19] "Carnitine" "Citrate"
## [21] "Creatine" "Creatinine"
## [23] "Dimethylamine" "Ethanolamine"
## [25] "Formate" "Fucose"
## [27] "Fumarate" "Glucose"
## [29] "Glutamine" "Glycine"
## [31] "Glycolate" "Guanidoacetate"
## [33] "Hippurate" "Histidine"
## [35] "Hypoxanthine" "Isoleucine"
## [37] "Lactate" "Leucine"
## [39] "Lysine" "Methylamine"
## [41] "Methylguanidine" "N.N.Dimethylglycine"
## [43] "O.Acetylcarnitine" "Pantothenate"
## [45] "Pyroglutamate" "Pyruvate"
## [47] "Quinolate" "Serine"
## [49] "Succinate" "Sucrose"
## [51] "Tartrate" "Taurine"
## [53] "Threonine" "Trigonelline"
```

```
## [55] "Trimethylamine.N.oxide"      "Tryptophan"
## [57] "Tyrosine"                    "Uracil"
## [59] "Valine"                      "Xylose"
## [61] "cis.Aconitate"               "myo.Inositol"
## [63] "trans.Aconitate"             "pi.Methylhistidine"
## [65] "tau.Methylhistidine"
```

Podemos observar que las dos primeras columnas corresponden a metadatos: Patient.ID (es el identificador de cada individuo) y Muscle.loss (indica a qué grupo pertenece cada individuo). El resto de columnas son variables que corresponden a metabolitos.

Teniendo esto en cuenta podremos crear nuestro contenedor SummarizedExperiment

```
# Cargar la biblioteca SummarizedExperiment
library(SummarizedExperiment)
```

```
## Cargando paquete requerido: MatrixGenerics
## Cargando paquete requerido: matrixStats
## Warning: package 'matrixStats' was built under R version 4.4.1
##
## Adjuntando el paquete: 'MatrixGenerics'
## The following objects are masked from 'package:matrixStats':
##
##      colAlls, colAnyNAs, colAnys, colAvgPerRowSet, colCollapse,
##      colCounts, colCummaxs, colCummins, colCumprods, colCumsums,
##      colDiffs, colIQRDiffs, colIQRs, colLogSumExps, colMadDiffs,
##      colMads, colMaxs, colMeans2, colMedians, colMins, colOrderStats,
##      colProds, colQuantiles, colRanges, colRanks, colSdDiffs, colSds,
##      colSums2, colTabulates, colVarDiffs, colVars, colWeightedMads,
##      colWeightedMeans, colWeightedMedians, colWeightedSds,
##      colWeightedVars, rowAlls, rowAnyNAs, rowAnys, rowAvgPerColSet,
##      rowCollapse, rowCounts, rowCummaxs, rowCummins, rowCumprods,
##      rowCumsums, rowDiffs, rowIQRDiffs, rowIQRs, rowLogSumExps,
##      rowMadDiffs, rowMads, rowMaxs, rowMeans2, rowMedians, rowMins,
##      rowOrderStats, rowProds, rowQuantiles, rowRanges, rowRanks,
##      rowSdDiffs, rowSds, rowSums2, rowTabulates, rowVarDiffs, rowVars,
##      rowWeightedMads, rowWeightedMeans, rowWeightedMedians,
##      rowWeightedSds, rowWeightedVars
## Cargando paquete requerido: GenomicRanges
## Cargando paquete requerido: stats4
## Cargando paquete requerido: BiocGenerics
##
## Adjuntando el paquete: 'BiocGenerics'
## The following objects are masked from 'package:stats':
##
##      IQR, mad, sd, var, xtabs
## The following objects are masked from 'package:base':
##
##      anyDuplicated, aperm, append, as.data.frame, basename, cbind,
##      colnames, dirname, do.call, duplicated, eval, evalq, Filter, Find,
##      get, grep, grepl, intersect, is.unsorted, lapply, Map, mapply,
```

```

##      match, mget, order, paste, pmax, pmax.int, pmin, pmin.int,
##      Position, rank, rbind, Reduce, rownames, sapply, setdiff, table,
##      tapply, union, unique, unsplit, which.max, which.min

## Cargando paquete requerido: S4Vectors

## Warning: package 'S4Vectors' was built under R version 4.4.1

##
## Adjuntando el paquete: 'S4Vectors'

## The following object is masked from 'package:utils':
##
##      findMatches

## The following objects are masked from 'package:base':
##
##      expand.grid, I, unname

## Cargando paquete requerido: IRanges

## Warning: package 'IRanges' was built under R version 4.4.1

##
## Adjuntando el paquete: 'IRanges'

## The following object is masked from 'package:grDevices':
##
##      windows

## Cargando paquete requerido: GenomeInfoDb

## Cargando paquete requerido: Biobase

## Welcome to Bioconductor
##
##      Vignettes contain introductory material; view with
##      'browseVignettes()'. To cite Bioconductor, see
##      'citation("Biobase")', and for packages 'citation("pkgname")'.

##
## Adjuntando el paquete: 'Biobase'

## The following object is masked from 'package:MatrixGenerics':
##
##      rowMedians

## The following objects are masked from 'package:matrixStats':
##
##      anyMissing, rowMedians

# Definir los metadatos ("Patient.ID" y "Muscle.loss") como las columnas
metadatos_cachexia <- cachexia[, 1:2]
colnames(metadatos_cachexia) <- c("Patient.ID", "Muscle.loss")

# Definir los datos de expresión (columnas correspondientes a los metabolitos)
datos_expresion <- as.matrix(cachexia[, 3:ncol(cachexia)])
datos_expresion <- t(datos_expresion)

# Crear un data frame para indicar en rowData que las variables corresponden a metabolitos
row_data <- data.frame(Metabolitos = rownames(datos_expresion))

```

```

# Crear el objeto SummarizedExperiment
contenedor_cachexia <- SummarizedExperiment(
  assays = list(counts = datos_expresion),
  colData = metadatos_cachexia,
  rowData = row_data
)

# Visualizar el objeto SummarizedExperiment que hemos creado
contenedor_cachexia

## class: SummarizedExperiment
## dim: 63 77
## metadata(0):
## assays(1): counts
## rownames(63): X1.6.Anhydro.beta.D.glucose X1.Methylnicotinamide ...
##   pi.Methylhistidine tau.Methylhistidine
## rowData names(1): Metabolitos
## colnames: NULL
## colData names(2): Patient.ID Muscle.loss

Luego, guardamos el contenedor en formato .Rda.
# Guardar el contenedor en formato .Rda
save(contenedor_cachexia, file = "D:/Máster bioinformática/Análisis de datos ómicos/PEC1/contenedor_cachexia.Rda")

```

3. Exploración del dataset

Análisis convencional del dataset

En primer lugar, podemos visualizar el tamaño y estructura del dataset, así como obtener un resumen estadístico para cada variable.

```

# Mostrar las dimensiones del dataset
dim(cachexia)

## [1] 77 65

# Mostrar los nombres de las columnas
colnames(cachexia)

## [1] "Patient.ID" "Muscle.loss"
## [3] "X1.6.Anhydro.beta.D.glucose" "X1.Methylnicotinamide"
## [5] "X2.Aminobutyrate" "X2.Hydroxyisobutyrate"
## [7] "X2.Oxoglutarate" "X3.Aminoisobutyrate"
## [9] "X3.Hydroxybutyrate" "X3.Hydroxyisovalerate"
## [11] "X3.Indoxylsulfate" "X4.Hydroxyphenylacetate"
## [13] "Acetate" "Acetone"
## [15] "Adipate" "Alanine"
## [17] "Asparagine" "Betaine"
## [19] "Carnitine" "Citrate"
## [21] "Creatine" "Creatinine"
## [23] "Dimethylamine" "Ethanolamine"
## [25] "Formate" "Fucose"
## [27] "Fumarate" "Glucose"
## [29] "Glutamine" "Glycine"
## [31] "Glycolate" "Guanidoacetate"

```

```
## [33] "Hippurate"           "Histidine"
## [35] "Hypoxanthine"        "Isoleucine"
## [37] "Lactate"             "Leucine"
## [39] "Lysine"              "Methylamine"
## [41] "Methylguanidine"     "N.N.Dimethylglycine"
## [43] "O.Acetylcarnitine"   "Pantothenate"
## [45] "Pyroglutamate"       "Pyruvate"
## [47] "Quinolate"           "Serine"
## [49] "Succinate"           "Sucrose"
## [51] "Tartrate"            "Taurine"
## [53] "Threonine"           "Trigonelline"
## [55] "Trimethylamine.N.oxide" "Tryptophan"
## [57] "Tyrosine"            "Uracil"
## [59] "Valine"              "Xylose"
## [61] "cis.Aconitate"        "myo.Inositol"
## [63] "trans.Aconitate"     "pi.Methylhistidine"
## [65] "tau.Methylhistidine"
```

```
# Mostrar los tipos de datos que tiene cada variable
str(cachexia)
```

```
## 'data.frame':    77 obs. of  65 variables:
## $ Patient.ID      : chr  "PIF_178" "PIF_087" "PIF_090" "NETL_005_V1" ...
## $ Muscle.loss      : chr  "cachexic" "cachexic" "cachexic" "cachexic" ...
## $ X1.6.Anhydro.beta.D.glucose: num  40.9 62.2 270.4 154.5 22.2 ...
## $ X1.Methylnicotinamide : num  65.4 340.4 64.7 53 73.7 ...
## $ X2.Aminobutyrate   : num  18.7 24.3 12.2 172.4 15.6 ...
## $ X2.Hydroxyisobutyrate : num  26.1 41.7 65.4 74.4 83.9 ...
## $ X2.Oxoglutarate    : num  71.5 67.4 23.8 1199.9 33.1 ...
## $ X3.Aminoisobutyrate : num  1480.3 116.8 14.3 555.6 29.7 ...
## $ X3.Hydroxybutyrate  : num  56.83 43.82 5.64 175.91 76.71 ...
## $ X3.Hydroxyisovalerate : num  10.1 79.8 23.3 25 69.4 ...
## $ X3.Indoxylsulfate   : num  567 369 665 412 166 ...
## $ X4.Hydroxyphenylacetate : num  120.3 432.7 292.9 214.9 97.5 ...
## $ Acetate            : num  126.5 212.7 314.2 37.3 407.5 ...
## $ Acetone            : num  9.49 11.82 4.44 206.44 44.26 ...
## $ Adipate            : num  38.1 327 131.6 144 15 ...
## $ Alanine            : num  314 871 464 590 1119 ...
## $ Asparagine         : num  159.2 157.6 89.1 273.1 42.5 ...
## $ Betaine            : num  110 245 117 279 392 ...
## $ Carnitine          : num  265.1 120.3 25 200.3 84.8 ...
## $ Citrate            : num  3714 2618 863 13630 854 ...
## $ Creatine           : num  196.4 212.7 221.4 85.6 105.6 ...
## $ Creatinine         : num  16482 15835 24588 20952 6768 ...
## $ Dimethylamine      : num  633 608 735 1064 242 ...
## $ Ethanolamine       : num  645 488 407 821 365 ...
## $ Formate            : num  441 252 250 469 114 ...
## $ Fucose             : num  337 198.3 186.8 407.5 26.1 ...
## $ Fumarate           : num  7.69 18.92 7.1 96.54 19.69 ...
## $ Glucose            : num  395 8691 1353 863 6836 ...
## $ Glutamine          : num  871 602 302 1686 433 ...
## $ Glycine            : num  2039 1108 620 5064 395 ...
## $ Glycolate          : num  685.4 652 141.2 70.8 26.6 ...
## $ Guanidoacetate     : num  154 110 183 103 53 ...
## $ Hippurate          : num  4582 1737 4316 757 1153 ...
```

```
## $ Histidine : num 925 846 284 1043 327 ...
## $ Hypoxanthine : num 97.5 82.3 114.4 223.6 66.7 ...
## $ Isoleucine : num 5.58 8.17 9.3 37.71 40.04 ...
## $ Lactate : num 107 369 750 369 3641 ...
## $ Leucine : num 42.1 77.5 31.5 103.5 101.5 ...
## $ Lysine : num 146.9 284.3 97.5 290 122.7 ...
## $ Methylamine : num 52.5 23.6 18.7 48.9 27.9 ...
## $ Methylguanidine : num 9.97 7.69 4.66 141.17 5.31 ...
## $ N.N.Dimethylglycine : num 23.3 87.4 24.5 40 46.1 ...
## $ O.Acetylcarnitine : num 52.98 50.4 5.58 254.68 45.6 ...
## $ Pantothenate : num 25.8 186.8 145.5 42.5 74.4 ...
## $ Pyroglutamate : num 437 437 713 567 185 ...
## $ Pyruvate : num 21.1 37 29.4 64.1 12.3 ...
## $ Quinolate : num 165.7 73 192.5 86.5 38.1 ...
## $ Serine : num 284 392 296 1249 206 ...
## $ Succinate : num 154.5 244.7 142.6 144 68.7 ...
## $ Sucrose : num 45.1 459.4 160.8 111 75.2 ...
## $ Tartrate : num 97.51 32.79 16.28 837.15 4.53 ...
## $ Taurine : num 1920 1261 4273 1525 469 ...
## $ Threonine : num 184.9 198.3 110 376.1 64.1 ...
## $ Trigonelline : num 943.9 208.5 192.5 992.3 86.5 ...
## $ Trimethylamine.N.oxide : num 2122 639 1153 1451 172 ...
## $ Tryptophan : num 259.8 83.1 82.3 235.1 103.5 ...
## $ Tyrosine : num 290 167.3 60.3 323.8 142.6 ...
## $ Uracil : num 111 47 31.5 30.6 44.3 ...
## $ Valine : num 86.5 110 59.1 102.5 160.8 ...
## $ Xylose : num 72.2 192.5 2164.6 125.2 186.8 ...
## $ cis.Aconitate : num 237 334 330 1863 101 ...
## $ myo.Inositol : num 135.6 376.1 86.5 247.2 750 ...
## $ trans.Aconitate : num 51.9 217 58.6 75.9 98.5 ...
## $ pi.Methylhistidine : num 157.6 308 145.5 249.6 84.8 ...
## $ tau.Methylhistidine : num 160.8 130.3 83.9 254.7 79.8 ...
```

Podemos observar que tenemos 77 muestras con 65 variables, siendo todas numéricas excepto Patient.ID y Muscle.loss, las cuales son de tipo character.

También podemos visualizar las primeras filas del dataset y obtener un resumen estadístico.

```
# Mostrar las primeras filas
head(cachexia)
```

```
## Patient.ID Muscle.loss X1.6.Anhydro.beta.D.glucose X1.Methylnicotinamide
## 1 PIF_178 cachexic 40.85 65.37
## 2 PIF_087 cachexic 62.18 340.36
## 3 PIF_090 cachexic 270.43 64.72
## 4 NETL_005_V1 cachexic 154.47 52.98
## 5 PIF_115 cachexic 22.20 73.70
## 6 PIF_110 cachexic 212.72 31.82
## X2.Aminobutyrate X2.Hydroxyisobutyrate X2.Oxoglutarate X3.Aminoisobutyrate
## 1 18.73 26.05 71.52 1480.30
## 2 24.29 41.68 67.36 116.75
## 3 12.18 65.37 23.81 14.30
## 4 172.43 74.44 1199.91 555.57
## 5 15.64 83.93 33.12 29.67
## 6 18.36 80.64 47.94 17.46
## X3.Hydroxybutyrate X3.Hydroxyisovalerate X3.Indoxylsulfate
```

## 1	56.83		10.07		566.80			
## 2	43.82		79.84		368.71			
## 3	5.64		23.34		665.14			
## 4	175.91		25.03		411.58			
## 5	76.71		69.41		165.67			
## 6	31.82		35.16		183.09			
##	X4.Hydroxyphenylacetate	Acetate	Acetone	Adipate	Alanine	Asparagine	Betaine	
## 1	120.30	126.47	9.49	38.09	314.19	159.17	109.95	
## 2	432.68	212.72	11.82	327.01	871.31	157.59	244.69	
## 3	292.95	314.19	4.44	131.63	464.05	89.12	116.75	
## 4	214.86	37.34	206.44	144.03	589.93	273.14	278.66	
## 5	97.51	407.48	44.26	15.03	1118.79	42.52	391.51	
## 6	132.95	81.45	14.44	25.28	237.46	157.59	66.69	
##	Carnitine	Citrate	Creatine	Creatinine	Dimethylamine	Ethanolamine	Formate	
## 1	265.07	3714.50	196.37	16481.60	632.70	645.48	441.42	
## 2	120.30	2617.57	212.72	15835.35	607.89	487.85	252.14	
## 3	25.03	862.64	221.41	24587.66	735.10	407.48	249.64	
## 4	200.34	13629.61	85.63	20952.22	1064.22	820.57	468.72	
## 5	84.77	854.06	105.64	6768.26	242.26	365.04	114.43	
## 6	40.04	1958.63	200.34	15677.78	614.00	459.44	314.19	
##	Fucose	Fumarate	Glucose	Glutamine	Glycine	Glycolate	Guanidoacetate	Hippurate
## 1	336.97	7.69	395.44	871.31	2038.56	685.40	154.47	4582.50
## 2	198.34	18.92	8690.62	601.85	1107.65	651.97	109.95	1737.15
## 3	186.79	7.10	1352.89	301.87	620.17	141.17	183.09	4315.64
## 4	407.48	96.54	862.64	1685.81	5064.45	70.81	102.51	757.48
## 5	26.05	19.69	6836.29	432.68	395.44	26.58	52.98	1152.86
## 6	123.97	5.05	512.86	298.87	482.99	428.38	57.97	3568.85
##	Histidine	Hypoxanthine	Isoleucine	Lactate	Leucine	Lysine	Methylamine	
## 1	925.19		97.51	5.58	106.70	42.10	146.94	52.46
## 2	845.56		82.27	8.17	368.71	77.48	284.29	23.57
## 3	284.29		114.43	9.30	749.95	31.50	97.51	18.73
## 4	1043.15		223.63	37.71	368.71	103.54	290.03	48.91
## 5	327.01		66.69	40.04	3640.95	101.49	122.73	27.94
## 6	459.44		62.80	8.17	113.30	28.79	120.30	36.97
##	Methylguanidine	N.N.Dimethylglycine	O.Acetylcarnitine	Pantothenate				
## 1	9.97		23.34		52.98		25.79	
## 2	7.69		87.36		50.40		186.79	
## 3	4.66		24.53		5.58		145.47	
## 4	141.17		40.04		254.68		42.52	
## 5	5.31		46.06		45.60		74.44	
## 6	43.38		24.29		13.46		35.52	
##	Pyroglutamate	Pyruvate	Quinolinolate	Serine	Succinate	Sucrose	Tartrate	Taurine
## 1	437.03	21.12	165.67	284.29	154.47	45.15	97.51	1919.85
## 2	437.03	36.97	72.97	391.51	244.69	459.44	32.79	1261.43
## 3	713.37	29.37	192.48	295.89	142.59	160.77	16.28	4272.69
## 4	566.80	64.07	86.49	1248.88	144.03	111.05	837.15	1525.38
## 5	184.93	12.30	38.09	206.44	68.72	75.19	4.53	468.72
## 6	432.68	32.79	112.17	387.61	33.45	336.97	24.05	2059.05
##	Threonine	Trigonelline	Trimethylamine.N.oxide	Tryptophan	Tyrosine	Uracil		
## 1	184.93	943.88		2121.76	259.82	290.03	111.05	
## 2	198.34	208.51		639.06	83.10	167.34	46.99	
## 3	109.95	192.48		1152.86	82.27	60.34	31.50	
## 4	376.15	992.27		1450.99	235.10	323.76	30.57	
## 5	64.07	86.49		172.43	103.54	142.59	44.26	

```
## 6      105.64      862.64      880.07      239.85      127.74      29.67
## Valine Xylose cis.Aconitate myo.Inositol trans.Aconitate pi.Methylhistidine
## 1 86.49 72.24 237.46 135.64 51.94 157.59
## 2 109.95 192.48 333.62 376.15 217.02 307.97
## 3 59.15 2164.62 330.30 86.49 58.56 145.47
## 4 102.51 125.21 1863.11 247.15 75.94 249.64
## 5 160.77 186.79 101.49 749.95 98.49 84.77
## 6 36.97 89.12 287.15 129.02 121.51 399.41
## tau.Methylhistidine
## 1 160.77
## 2 130.32
## 3 83.93
## 4 254.68
## 5 79.84
## 6 68.72
```

```
# Obtener un resumen estadístico de las columnas
summary(cachexia)
```

```
## Patient.ID      Muscle.loss      X1.6.Anhydro.beta.D.glucose
## Length:77      Length:77      Min. : 4.71
## Class :character Class :character 1st Qu.: 28.79
## Mode :character Mode :character Median : 45.60
## Mean :105.63
## 3rd Qu.:141.17
## Max. :685.40
## X1.Methylnicotinamide X2.Aminobutyrate X2.Hydroxyisobutyrate X2.Oxoglutarate
## Min. : 6.42 Min. : 1.28 Min. : 4.85 Min. : 5.53
## 1st Qu.: 15.80 1st Qu.: 5.26 1st Qu.:15.80 1st Qu.: 22.42
## Median : 36.60 Median : 10.49 Median :32.46 Median : 55.15
## Mean : 71.57 Mean : 18.16 Mean :37.25 Mean : 145.09
## 3rd Qu.: 73.70 3rd Qu.: 19.49 3rd Qu.:54.60 3rd Qu.: 92.76
## Max. :1032.77 Max. :172.43 Max. :93.69 Max. :2465.13
## X3.Aminoisobutyrate X3.Hydroxybutyrate X3.Hydroxyisovalerate X3.Indoxylsulfate
## Min. : 2.61 Min. : 1.70 Min. : 0.92 Min. : 27.66
## 1st Qu.: 11.70 1st Qu.: 5.99 1st Qu.: 5.26 1st Qu.: 82.27
## Median : 22.65 Median : 11.70 Median : 12.55 Median : 144.03
## Mean : 76.76 Mean : 21.72 Mean : 21.65 Mean : 218.88
## 3rd Qu.: 56.26 3rd Qu.: 29.96 3rd Qu.: 30.27 3rd Qu.: 333.62
## Max. :1480.30 Max. :175.91 Max. :164.02 Max. :1043.15
## X4.Hydroxyphenylacetate Acetate Acetone Adipate
## Min. : 15.49 Min. : 3.49 Min. : 2.29 Min. : 1.55
## 1st Qu.: 41.68 1st Qu.: 16.28 1st Qu.: 4.95 1st Qu.: 6.11
## Median : 70.11 Median : 39.65 Median : 7.10 Median : 10.18
## Mean :112.02 Mean : 66.14 Mean : 11.43 Mean : 24.76
## 3rd Qu.:145.47 3rd Qu.: 86.49 3rd Qu.: 10.49 3rd Qu.: 19.11
## Max. :796.32 Max. :411.58 Max. :206.44 Max. :327.01
## Alanine Asparagine Betaine Carnitine
## Min. : 16.78 Min. : 6.69 Min. : 2.29 Min. : 2.18
## 1st Qu.: 78.26 1st Qu.: 20.49 1st Qu.: 28.79 1st Qu.: 14.44
## Median : 194.42 Median : 42.10 Median : 64.72 Median : 23.81
## Mean : 273.56 Mean : 62.28 Mean : 90.32 Mean : 52.09
## 3rd Qu.: 399.41 3rd Qu.: 89.12 3rd Qu.:127.74 3rd Qu.: 60.95
## Max. :1312.91 Max. :273.14 Max. :391.51 Max. :487.85
## Citrate Creatine Creatinine Dimethylamine
```


##	Min.	: 59.74	Min.	: 2.75	Min.	: 1002	Min.	: 41.26
##	1st Qu.:	788.40	1st Qu.:	17.64	1st Qu.:	3498	1st Qu.:	142.59
##	Median :	1790.05	Median :	44.26	Median :	7631	Median :	304.90
##	Mean :	2235.35	Mean :	126.83	Mean :	8734	Mean :	358.17
##	3rd Qu.:	3071.74	3rd Qu.:	117.92	3rd Qu.:	12333	3rd Qu.:	454.86
##	Max.	:13629.61	Max.	:1863.11	Max.	:33860	Max.	:1556.20
##	Ethanolamine		Formate		Fucose		Fumarate	
##	Min.	: 16.12	Min.	: 6.42	Min.	: 5.70	Min.	: 0.79
##	1st Qu.:	86.49	1st Qu.:	53.52	1st Qu.:	29.37	1st Qu.:	2.23
##	Median :	204.38	Median :	95.58	Median :	61.56	Median :	4.10
##	Mean :	276.26	Mean :	147.40	Mean :	88.67	Mean :	8.44
##	3rd Qu.:	407.48	3rd Qu.:	167.34	3rd Qu.:	123.97	3rd Qu.:	7.85
##	Max.	:1436.55	Max.	:1480.30	Max.	:407.48	Max.	:96.54
##	Glucose		Glutamine		Glycine		Glycolate	
##	Min.	: 26.84	Min.	: 23.34	Min.	: 38.09	Min.	: 5.42
##	1st Qu.:	80.64	1st Qu.:	113.30	1st Qu.:	262.43	1st Qu.:	50.91
##	Median :	210.61	Median :	225.88	Median :	528.48	Median :	130.32
##	Mean :	559.85	Mean :	306.87	Mean :	880.72	Mean :	187.99
##	3rd Qu.:	407.48	3rd Qu.:	445.86	3rd Qu.:	1096.63	3rd Qu.:	267.74
##	Max.	:8690.62	Max.	:1685.81	Max.	:5064.45	Max.	:720.54
##	Guanidoacetate		Hippurate		Histidine		Hypoxanthine	
##	Min.	: 7.03	Min.	: 92.76	Min.	: 14.15	Min.	: 3.78
##	1st Qu.:	33.78	1st Qu.:	492.75	1st Qu.:	66.69	1st Qu.:	20.70
##	Median :	64.72	Median :	1224.15	Median :	174.16	Median :	40.04
##	Mean :	86.37	Mean :	2286.84	Mean :	292.64	Mean :	61.10
##	3rd Qu.:	108.85	3rd Qu.:	2921.93	3rd Qu.:	419.89	3rd Qu.:	83.93
##	Max.	:561.16	Max.	:19341.34	Max.	:1863.11	Max.	:265.07
##	Isoleucine		Lactate		Leucine		Lysine	
##	Min.	: 1.790	Min.	: 7.32	Min.	: 2.51	Min.	: 10.49
##	1st Qu.:	3.900	1st Qu.:	35.52	1st Qu.:	9.12	1st Qu.:	30.27
##	Median :	7.170	Median :	81.45	Median :	19.11	Median :	69.41
##	Mean :	8.709	Mean :	158.46	Mean :	24.36	Mean :	108.79
##	3rd Qu.:	11.250	3rd Qu.:	139.77	3rd Qu.:	31.19	3rd Qu.:	121.51
##	Max.	:40.040	Max.	:3640.95	Max.	:103.54	Max.	:788.40
##	Methylamine		Methylguanidine		N.N.Dimethylglycine		O.Acetylcarnitine	
##	Min.	: 1.51	Min.	: 1.70	Min.	: 0.79	Min.	: 1.23
##	1st Qu.:	5.26	1st Qu.:	4.26	1st Qu.:	7.03	1st Qu.:	3.94
##	Median :	14.73	Median :	7.85	Median :	21.98	Median :	11.47
##	Mean :	17.38	Mean :	15.32	Mean :	26.35	Mean :	19.73
##	3rd Qu.:	24.05	3rd Qu.:	19.30	3rd Qu.:	40.04	3rd Qu.:	20.91
##	Max.	:52.46	Max.	:141.17	Max.	:120.30	Max.	:254.68
##	Pantothenate		Pyroglutamate		Pyruvate		Quinolinate	
##	Min.	: 2.59	Min.	: 21.33	Min.	: 0.90	Min.	: 5.21
##	1st Qu.:	11.13	1st Qu.:	68.72	1st Qu.:	4.85	1st Qu.:	26.58
##	Median :	22.65	Median :	157.59	Median :	13.46	Median :	51.42
##	Mean :	44.88	Mean :	211.45	Mean :	21.29	Mean :	66.44
##	3rd Qu.:	41.26	3rd Qu.:	301.87	3rd Qu.:	29.08	3rd Qu.:	87.36
##	Max.	:692.29	Max.	:1064.22	Max.	:184.93	Max.	:259.82
##	Serine		Succinate		Sucrose		Tartrate	
##	Min.	: 16.12	Min.	: 1.72	Min.	: 6.49	Min.	: 2.20
##	1st Qu.:	83.10	1st Qu.:	8.58	1st Qu.:	19.30	1st Qu.:	6.89
##	Median :	142.59	Median :	30.88	Median :	40.85	Median :	12.94
##	Mean :	197.69	Mean :	60.23	Mean :	113.23	Mean :	40.00
##	3rd Qu.:	270.43	3rd Qu.:	74.44	3rd Qu.:	94.63	3rd Qu.:	25.79

```
## Max. :1248.88 Max. :589.93 Max. :2079.74 Max. :837.15
## Taurine Threonine Trigonelline Trimethylamine.N.oxide
## Min. : 17.81 Min. : 8.25 Min. : 10.07 Min. : 55.7
## 1st Qu.: 99.48 1st Qu.: 31.82 1st Qu.: 53.52 1st Qu.: 175.9
## Median : 249.64 Median : 64.07 Median : 114.43 Median : 383.8
## Mean : 525.12 Mean : 95.36 Mean : 270.44 Mean : 652.2
## 3rd Qu.: 665.14 3rd Qu.:137.00 3rd Qu.: 340.36 3rd Qu.: 735.1
## Max. :4272.69 Max. :450.34 Max. :2252.96 Max. :5486.2
## Tryptophan Tyrosine Uracil Valine
## Min. : 8.67 Min. : 4.22 Min. : 3.10 Min. : 4.10
## 1st Qu.: 21.33 1st Qu.: 23.57 1st Qu.: 11.94 1st Qu.: 12.18
## Median : 46.99 Median : 60.34 Median : 27.39 Median : 33.12
## Mean : 66.24 Mean : 81.76 Mean : 35.56 Mean : 35.67
## 3rd Qu.: 96.54 3rd Qu.:113.30 3rd Qu.: 44.26 3rd Qu.: 50.40
## Max. :259.82 Max. :539.15 Max. :179.47 Max. :160.77
## Xylose cis.Aconitate myo.Inositol trans.Aconitate
## Min. : 10.07 Min. : 12.94 Min. : 11.59 Min. : 4.90
## 1st Qu.: 29.96 1st Qu.: 36.23 1st Qu.: 30.27 1st Qu.: 12.43
## Median : 50.40 Median : 129.02 Median : 78.26 Median : 26.84
## Mean : 100.93 Mean : 204.22 Mean :135.40 Mean : 40.63
## 3rd Qu.: 89.12 3rd Qu.: 254.68 3rd Qu.:167.34 3rd Qu.: 57.40
## Max. :2164.62 Max. :1863.11 Max. :854.06 Max. :217.02
## pi.Methylhistidine tau.Methylhistidine
## Min. : 11.36 Min. : 8.00
## 1st Qu.: 67.36 1st Qu.: 27.39
## Median : 162.39 Median : 68.72
## Mean : 370.29 Mean : 89.69
## 3rd Qu.: 387.61 3rd Qu.:130.32
## Max. :2697.28 Max. :317.35
```

Comprobaremos si hay algún valor faltante.

```
# Visualizar si hay filas o columnas con NA
anyNA(cachexia)
```

```
## [1] FALSE
```

Observamos que en nuestro caso no hay ninguno.

Análisis utilizando el objeto SummarizedExperiment

La información que ya hemos comprobado como las dimensiones o estructura del dataset también puede comprobarse a través del contenedor que hemos creado.

```
# Mostrar la estructura del contenedor
contenedor_cachexia
```

```
## class: SummarizedExperiment
## dim: 63 77
## metadata(0):
## assays(1): counts
## rownames(63): X1.6.Anhydro.beta.D.glucose X1.Methylnicotinamide ...
## pi.Methylhistidine tau.Methylhistidine
## rowData names(1): Metabolitos
## colnames: NULL
## colData names(2): Patient.ID Muscle.loss
```

```
# Visualizar los nombres de los metabolitos
```

```
rowData(contenedor_cachexia)
```

```
## DataFrame with 63 rows and 1 column
##                               Metabolitos
##                               <character>
## X1.6.Anhydro.beta.D.glucose X1.6.Anhydro.beta.D...
## X1.Methylnicotinamide       X1.Methylnicotinamide
## X2.Aminobutyrate            X2.Aminobutyrate
## X2.Hydroxyisobutyrate       X2.Hydroxyisobutyrate
## X2.Oxoglutarate             X2.Oxoglutarate
## ...                         ...
## cis.Aconitate               cis.Aconitate
## myo.Inositol                myo.Inositol
## trans.Aconitate             trans.Aconitate
## pi.Methylhistidine          pi.Methylhistidine
## tau.Methylhistidine         tau.Methylhistidine
```

```
# Visualizar los primeros datos de expresión
```

```
assay(contenedor_cachexia)[1:10, 1:10]
```

```
##           [,1] [,2] [,3] [,4] [,5] [,6] [,7]
## X1.6.Anhydro.beta.D.glucose 40.85 62.18 270.43 154.47 22.20 212.72 151.41
## X1.Methylnicotinamide       65.37 340.36 64.72  52.98 73.70  31.82  36.60
## X2.Aminobutyrate            18.73  24.29  12.18 172.43  15.64  18.36   8.67
## X2.Hydroxyisobutyrate       26.05  41.68  65.37  74.44  83.93  80.64  42.52
## X2.Oxoglutarate             71.52  67.36  23.81 1199.91  33.12  47.94 223.63
## X3.Aminoisobutyrate        1480.30 116.75  14.30  555.57  29.67  17.46  56.26
## X3.Hydroxybutyrate          56.83  43.82   5.64  175.91  76.71  31.82  11.59
## X3.Hydroxyisovalerate       10.07  79.84  23.34  25.03  69.41  35.16  25.79
## X3.Indoxylsulfate           566.80 368.71 665.14 411.58 165.67 183.09 223.63
## X4.Hydroxyphenylacetate     120.30 432.68 292.95 214.86  97.51 132.95  59.15
##           [,8] [,9] [,10]
## X1.6.Anhydro.beta.D.glucose 31.50 51.42 117.92
## X1.Methylnicotinamide       6.82 30.27  52.46
## X2.Aminobutyrate            4.18  7.54  19.49
## X2.Hydroxyisobutyrate       12.94 34.81  72.24
## X2.Oxoglutarate             25.03 80.64  73.70
## X3.Aminoisobutyrate         8.67 17.99  57.97
## X3.Hydroxybutyrate          1.73  9.03  26.84
## X3.Hydroxyisovalerate       8.76  3.25  28.50
## X3.Indoxylsulfate           111.05 391.51 116.75
## X4.Hydroxyphenylacetate     33.78 145.47  50.40
```

```
# Mostrar los metadatos
```

```
colData(contenedor_cachexia)
```

```
## DataFrame with 77 rows and 2 columns
##      Patient.ID Muscle.loss
##      <character> <character>
## 1      PIF_178    cachexic
## 2      PIF_087    cachexic
## 3      PIF_090    cachexic
## 4    NETL_005_V1    cachexic
## 5      PIF_115    cachexic
```

```
## ...
## 73 NETCR_019_V2 control
## 74 NETL_012_V1 control
## 75 NETL_012_V2 control
## 76 NETL_003_V1 control
## 77 NETL_003_V2 control
```

```
# Obtener el resumen estadístico de los datos de expresión
summary(datos_expresion)
```

```
##          V1          V2          V3          V4
## Min.   : 5.58   Min.   : 7.69   Min.   : 4.44   Min.   : 25.03
## 1st Qu.: 52.72   1st Qu.: 78.66   1st Qu.: 31.50   1st Qu.: 102.51
## Median : 154.47   Median : 208.51   Median : 141.17   Median : 247.15
## Mean   : 699.86   Mean   : 708.30   Mean   : 771.79   Mean   : 1021.28
## 3rd Qu.: 416.24   3rd Qu.: 412.10   3rd Qu.: 308.03   3rd Qu.: 673.71
## Max.   :16481.60   Max.   :15835.35   Max.   :24587.66   Max.   :20952.22
##          V5          V6          V7          V8
## Min.   : 4.53   Min.   : 5.05   Min.   : 2.10   Min.   : 1.73
## 1st Qu.: 44.26   1st Qu.: 35.34   1st Qu.: 26.73   1st Qu.: 7.14
## Median : 84.77   Median : 113.30   Median : 91.84   Median : 18.17
## Mean   : 441.22   Mean   : 537.48   Mean   : 400.85   Mean   : 82.77
## 3rd Qu.: 196.62   3rd Qu.: 325.58   3rd Qu.: 223.63   3rd Qu.: 52.52
## Max.   :6836.29   Max.   :15677.78   Max.   :8022.46   Max.   :2208.35
##          V9          V10         V11         V12
## Min.   : 2.41   Min.   : 9.12   Min.   : 4.26   Min.   : 7.17
## 1st Qu.: 14.63   1st Qu.: 43.82   1st Qu.: 31.98   1st Qu.: 38.77
## Median : 39.65   Median : 117.92   Median : 83.93   Median : 127.74
## Mean   : 207.80   Mean   : 478.07   Mean   : 367.52   Mean   : 650.75
## 3rd Qu.: 102.00   3rd Qu.: 405.50   3rd Qu.: 182.20   3rd Qu.: 283.05
## Max.   :6634.24   Max.   :8690.62   Max.   :8433.78   Max.   :19341.34
##          V13         V14         V15         V16
## Min.   : 6.05   Min.   : 3.49   Min.   : 1.48   Min.   : 2.230
## 1st Qu.: 36.30   1st Qu.: 28.09   1st Qu.: 5.17   1st Qu.: 5.965
## Median : 83.93   Median : 71.52   Median : 17.46   Median : 18.360
## Mean   : 484.70   Mean   : 355.17   Mean   : 53.48   Mean   : 56.669
## 3rd Qu.: 218.11   3rd Qu.: 152.94   3rd Qu.: 40.65   3rd Qu.: 37.155
## Max.   :15677.78   Max.   :12209.87   Max.   :1480.30   Max.   :1635.980
##          V17         V18         V19         V20
## Min.   : 3.29   Min.   : 3.39   Min.   : 2.92   Min.   : 3.10
## 1st Qu.: 17.91   1st Qu.: 26.06   1st Qu.: 18.82   1st Qu.: 39.26
## Median : 64.07   Median : 78.26   Median : 74.44   Median : 82.27
## Mean   : 318.71   Mean   : 424.13   Mean   : 356.55   Mean   : 461.16
## 3rd Qu.: 164.90   3rd Qu.: 193.47   3rd Qu.: 176.13   3rd Qu.: 296.43
## Max.   :9701.15   Max.   :10198.54   Max.   :6974.39   Max.   :11158.98
##          V21         V22         V23         V24
## Min.   : 4.85   Min.   : 5.26   Min.   : 4.35   Min.   : 1.55
## 1st Qu.: 28.80   1st Qu.: 44.97   1st Qu.: 39.41   1st Qu.: 8.85
## Median : 64.72   Median : 98.49   Median : 75.94   Median : 17.81
## Mean   : 460.75   Mean   : 645.12   Mean   : 546.23   Mean   : 153.92
## 3rd Qu.: 210.62   3rd Qu.: 397.55   3rd Qu.: 267.15   3rd Qu.: 53.80
## Max.   :9798.65   Max.   :14328.42   Max.   :13359.73   Max.   :5943.18
##          V25         V26         V27         V28
## Min.   : 4.71   Min.   : 4.57   Min.   : 6.42   Min.   : 2.41
## 1st Qu.: 16.20   1st Qu.: 25.03   1st Qu.: 69.42   1st Qu.: 31.00
```

##	Median :	31.19	Median :	72.97	Median :	196.37	Median :	97.51
##	Mean :	183.79	Mean :	350.55	Mean :	1237.54	Mean :	516.61
##	3rd Qu.:	108.31	3rd Qu.:	186.53	3rd Qu.:	641.11	3rd Qu.:	330.37
##	Max. :	4865.87	Max. :	8349.86	Max. :	33860.35	Max. :	11271.13
##	V29		V30		V31		V32	
##	Min. :	0.790	Min. :	10.07	Min. :	1.82	Min. :	2.69
##	1st Qu.:	6.425	1st Qu.:	46.06	1st Qu.:	13.33	1st Qu.:	31.82
##	Median :	15.180	Median :	115.58	Median :	45.15	Median :	70.81
##	Mean :	62.813	Mean :	738.89	Mean :	199.61	Mean :	376.69
##	3rd Qu.:	29.370	3rd Qu.:	336.99	3rd Qu.:	119.20	3rd Qu.:	267.74
##	Max. :	1737.150	Max. :	21590.31	Max. :	4188.09	Max. :	11731.12
##	V33		V34		V35		V36	
##	Min. :	2.32	Min. :	3.19	Min. :	2.08	Min. :	2.01
##	1st Qu.:	14.82	1st Qu.:	28.64	1st Qu.:	20.19	1st Qu.:	12.94
##	Median :	37.34	Median :	61.56	Median :	45.60	Median :	24.05
##	Mean :	227.97	Mean :	327.88	Mean :	191.82	Mean :	148.51
##	3rd Qu.:	104.17	3rd Qu.:	153.72	3rd Qu.:	110.89	3rd Qu.:	64.39
##	Max. :	5431.66	Max. :	8349.86	Max. :	5014.05	Max. :	4315.64
##	V37		V38		V39		V40	
##	Min. :	5.53	Min. :	4.01	Min. :	3.67	Min. :	2.18
##	1st Qu.:	42.32	1st Qu.:	49.70	1st Qu.:	21.02	1st Qu.:	14.88
##	Median :	101.49	Median :	116.75	Median :	62.80	Median :	50.91
##	Mean :	496.29	Mean :	581.79	Mean :	270.28	Mean :	198.65
##	3rd Qu.:	290.56	3rd Qu.:	330.69	3rd Qu.:	177.69	3rd Qu.:	121.56
##	Max. :	13359.73	Max. :	16481.60	Max. :	7631.20	Max. :	3533.34
##	V41		V42		V43		V44	
##	Min. :	5.47	Min. :	7.32	Min. :	1.95	Min. :	4.01
##	1st Qu.:	32.62	1st Qu.:	50.91	1st Qu.:	21.66	1st Qu.:	36.88
##	Median :	98.49	Median :	119.10	Median :	48.42	Median :	94.63
##	Mean :	502.98	Mean :	697.47	Mean :	279.24	Mean :	579.72
##	3rd Qu.:	234.00	3rd Qu.:	404.56	3rd Qu.:	144.90	3rd Qu.:	242.27
##	Max. :	12332.58	Max. :	19930.37	Max. :	7115.28	Max. :	14764.78
##	V45		V46		V47		V48	
##	Min. :	2.53	Min. :	6.62	Min. :	1.120	Min. :	0.90
##	1st Qu.:	61.26	1st Qu.:	45.40	1st Qu.:	7.885	1st Qu.:	9.68
##	Median :	120.30	Median :	127.74	Median :	27.390	Median :	21.98
##	Mean :	745.91	Mean :	525.02	Mean :	143.280	Mean :	72.36
##	3rd Qu.:	337.24	3rd Qu.:	497.73	3rd Qu.:	68.400	3rd Qu.:	44.70
##	Max. :	22247.84	Max. :	14328.42	Max. :	2864.070	Max. :	1702.75
##	V49		V50		V51		V52	
##	Min. :	6.89	Min. :	1.21	Min. :	1.28	Min. :	1.51
##	1st Qu.:	47.94	1st Qu.:	7.15	1st Qu.:	7.03	1st Qu.:	7.30
##	Median :	121.51	Median :	15.18	Median :	18.92	Median :	18.54
##	Mean :	639.13	Mean :	76.81	Mean :	71.90	Mean :	170.47
##	3rd Qu.:	306.53	3rd Qu.:	42.85	3rd Qu.:	44.26	3rd Qu.:	62.80
##	Max. :	15063.05	Max. :	2392.27	Max. :	2489.91	Max. :	4817.45
##	V53		V54		V55		V56	
##	Min. :	6.17	Min. :	7.10	Min. :	1.36	Min. :	0.79
##	1st Qu.:	27.12	1st Qu.:	35.34	1st Qu.:	7.58	1st Qu.:	5.56
##	Median :	106.70	Median :	101.49	Median :	14.30	Median :	11.25
##	Mean :	396.24	Mean :	343.37	Mean :	64.50	Mean :	54.02
##	3rd Qu.:	250.44	3rd Qu.:	231.62	3rd Qu.:	35.17	3rd Qu.:	24.31
##	Max. :	9996.60	Max. :	7480.09	Max. :	1480.30	Max. :	1064.22
##	V57		V58		V59		V60	

```
## Min. : 1.97 Min. : 4.39 Min. : 4.10 Min. : 1.77
## 1st Qu.: 20.19 1st Qu.: 29.52 1st Qu.: 26.57 1st Qu.: 15.34
## Median : 54.05 Median : 87.36 Median : 49.40 Median : 35.87
## Mean : 289.17 Mean : 347.33 Mean : 361.04 Mean : 137.42
## 3rd Qu.: 115.00 3rd Qu.: 234.28 3rd Qu.: 202.39 3rd Qu.: 79.56
## Max. : 6974.39 Max. : 8266.78 Max. : 11849.01 Max. : 3827.63
## V61 V62 V63 V64
## Min. : 4.31 Min. : 1.23 Min. : 1.14 Min. : 2.05
## 1st Qu.: 21.45 1st Qu.: 4.00 1st Qu.: 15.93 1st Qu.: 10.48
## Median : 62.18 Median : 13.46 Median : 46.06 Median : 23.57
## Mean : 357.12 Mean : 42.80 Mean : 316.91 Mean : 159.57
## 3rd Qu.: 177.72 3rd Qu.: 28.08 3rd Qu.: 107.86 3rd Qu.: 56.26
## Max. : 10614.75 Max. : 1339.43 Max. : 7785.36 Max. : 5115.34
## V65 V66 V67 V68
## Min. : 1.55 Min. : 3.29 Min. : 6.23 Min. : 3.03
## 1st Qu.: 5.78 1st Qu.: 22.43 1st Qu.: 50.41 1st Qu.: 9.30
## Median : 15.96 Median : 49.90 Median : 100.48 Median : 24.05
## Mean : 63.52 Mean : 240.74 Mean : 467.35 Mean : 97.24
## 3rd Qu.: 34.65 3rd Qu.: 125.86 3rd Qu.: 284.31 3rd Qu.: 63.47
## Max. : 1571.84 Max. : 6768.26 Max. : 13359.73 Max. : 2121.76
## V69 V70 V71 V72
## Min. : 3.10 Min. : 0.920 Min. : 1.21 Min. : 1.230
## 1st Qu.: 45.45 1st Qu.: 6.795 1st Qu.: 10.54 1st Qu.: 6.145
## Median : 152.93 Median : 17.990 Median : 26.05 Median : 17.460
## Mean : 511.55 Mean : 110.017 Mean : 118.81 Mean : 56.986
## 3rd Qu.: 323.05 3rd Qu.: 60.700 3rd Qu.: 60.37 3rd Qu.: 30.725
## Max. : 13493.99 Max. : 2298.470 Max. : 3165.29 Max. : 1002.250
## V73 V74 V75 V76
## Min. : 3.67 Min. : 1.84 Min. : 2.69 Min. : 2.51
## 1st Qu.: 22.12 1st Qu.: 10.70 1st Qu.: 9.30 1st Qu.: 14.88
## Median : 58.56 Median : 21.33 Median : 24.05 Median : 34.12
## Mean : 342.26 Mean : 142.84 Mean : 147.55 Mean : 159.46
## 3rd Qu.: 180.37 3rd Qu.: 63.44 3rd Qu.: 59.20 3rd Qu.: 90.47
## Max. : 10097.06 Max. : 3789.54 Max. : 3498.19 Max. : 3498.19
## V77
## Min. : 1.62
## 1st Qu.: 12.55
## Median : 24.29
## Mean : 121.70
## 3rd Qu.: 80.70
## Max. : 2864.07
```

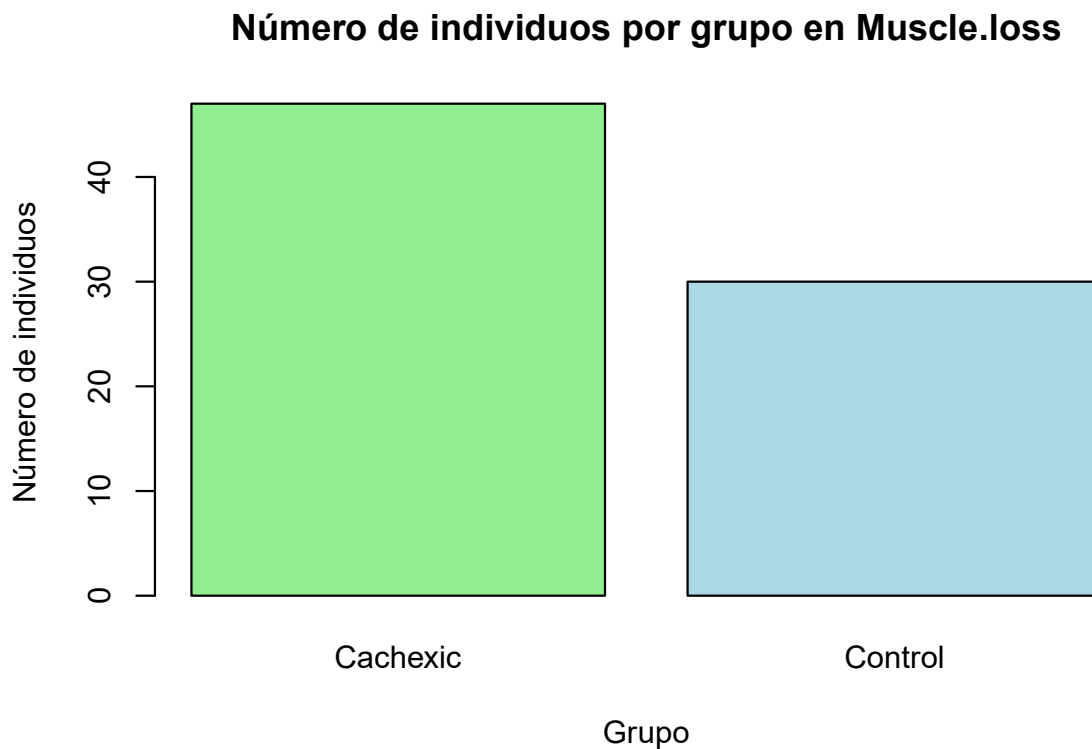
Análisis de las variables

Según la variable 'Muscle.loss' los individuos se clasifican en dos grupos (cachexic y control), por lo que resultaría interesante saber cuantos individuos hay para cada grupo.

```
# Calcular el número de individuos pertenecientes a cada grupo (cachexic y control)
muscle_loss <- colData(contenedor_cachexia)$Muscle.loss
conteo_grupos <- table(muscle_loss)
print(conteo_grupos)

## muscle_loss
## cachexic control
## 47 30
```

```
# Representar en un gráfico de barras el número de individuos de cada grupo
barplot(conteo_grupos,
      main="Número de individuos por grupo en Muscle.loss",
      xlab="Grupo",
      ylab="Número de individuos",
      col=c("lightgreen", "lightblue"),
      names.arg=c("Cachexic", "Control"))
```



Hemos obtenido que hay 47 individuos con cachexia y 30 individuos pertenecientes al grupo de control.

Una vez sabemos esto, podemos analizar qué variables presentan diferencias significativas entre los grupos, para lo que realizaremos una prueba de Wilcoxon.

```
# Crear una lista para almacenar los resultados
resultados <- list()

# Iterar sobre cada metabolito
for (i in 1:nrow(datos_expresion)) {
  metabolito <- datos_expresion[i, ]

  # Prueba de Wilcoxon
  test <- wilcox.test(metabolito ~ muscle_loss)

  # Almacenar los p-values y convertir los resultados a un data frame
  resultados[[rownames(datos_expresion)[i]]] <- test$p.value
}
```

```
## Warning in wilcox.test.default(x = DATA[[1L]], y = DATA[[2L]], ...): cannot
```

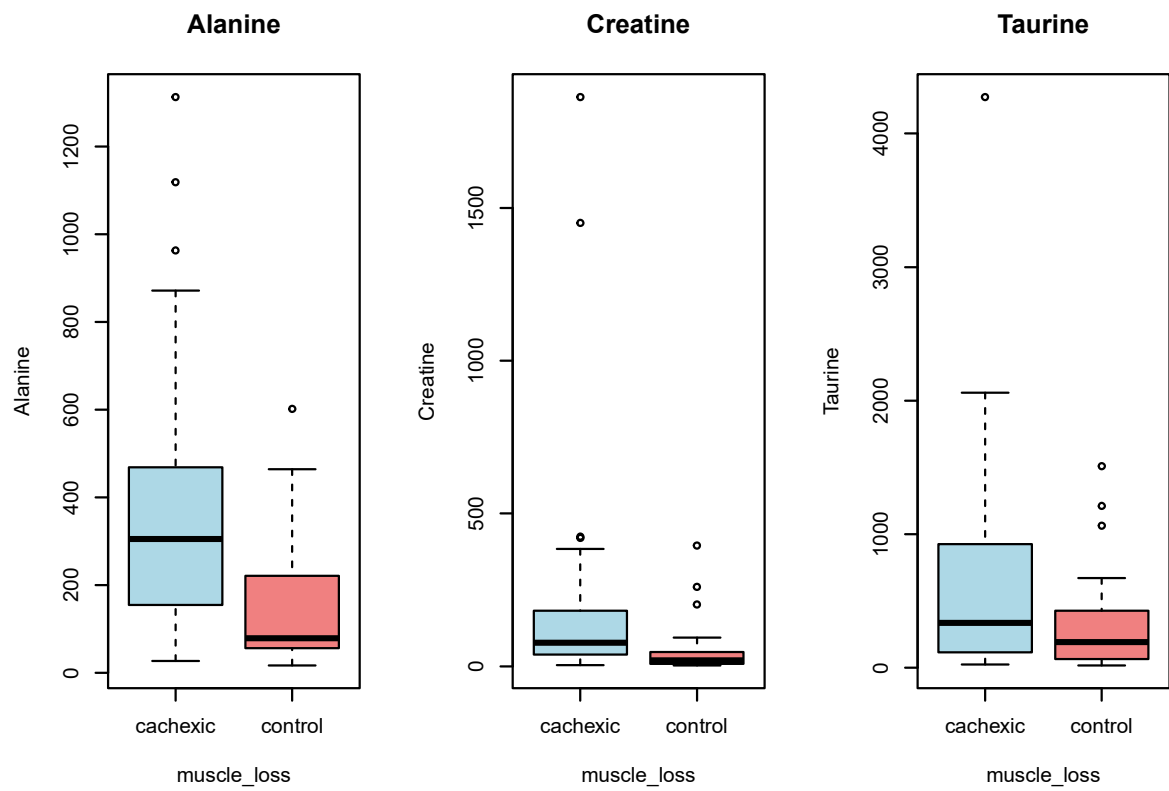

## Guanidoacetate	Guanidoacetate	2.011183e-02
## Hippurate	Hippurate	4.420227e-03
## Histidine	Histidine	2.326531e-03
## Lactate	Lactate	1.852242e-03
## Leucine	Leucine	5.772980e-05
## Lysine	Lysine	7.129366e-04
## Methylamine	Methylamine	1.694445e-04
## N.N.Dimethylglycine	N.N.Dimethylglycine	3.770829e-05
## O.Acetylcarnitine	O.Acetylcarnitine	1.875475e-02
## Pyroglutamate	Pyroglutamate	2.507912e-04
## Pyruvate	Pyruvate	2.960075e-03
## Quinolate	Quinolate	1.526936e-05
## Serine	Serine	9.276199e-04
## Succinate	Succinate	1.289705e-04
## Sucrose	Sucrose	2.613634e-04
## Taurine	Taurine	3.395917e-02
## Threonine	Threonine	8.944166e-04
## Trigonelline	Trigonelline	7.727431e-03
## Trimethylamine.N.oxide	Trimethylamine.N.oxide	8.742793e-03
## Tryptophan	Tryptophan	4.940642e-04
## Tyrosine	Tyrosine	1.442573e-03
## Valine	Valine	4.414600e-05
## Xylose	Xylose	4.308938e-04
## cis.Aconitate	cis.Aconitate	2.085041e-04
## myo.Inositol	myo.Inositol	1.087008e-04
## trans.Aconitate	trans.Aconitate	1.442260e-03
## pi.Methylhistidine	pi.Methylhistidine	1.875553e-02
## tau.Methylhistidine	tau.Methylhistidine	8.220042e-03

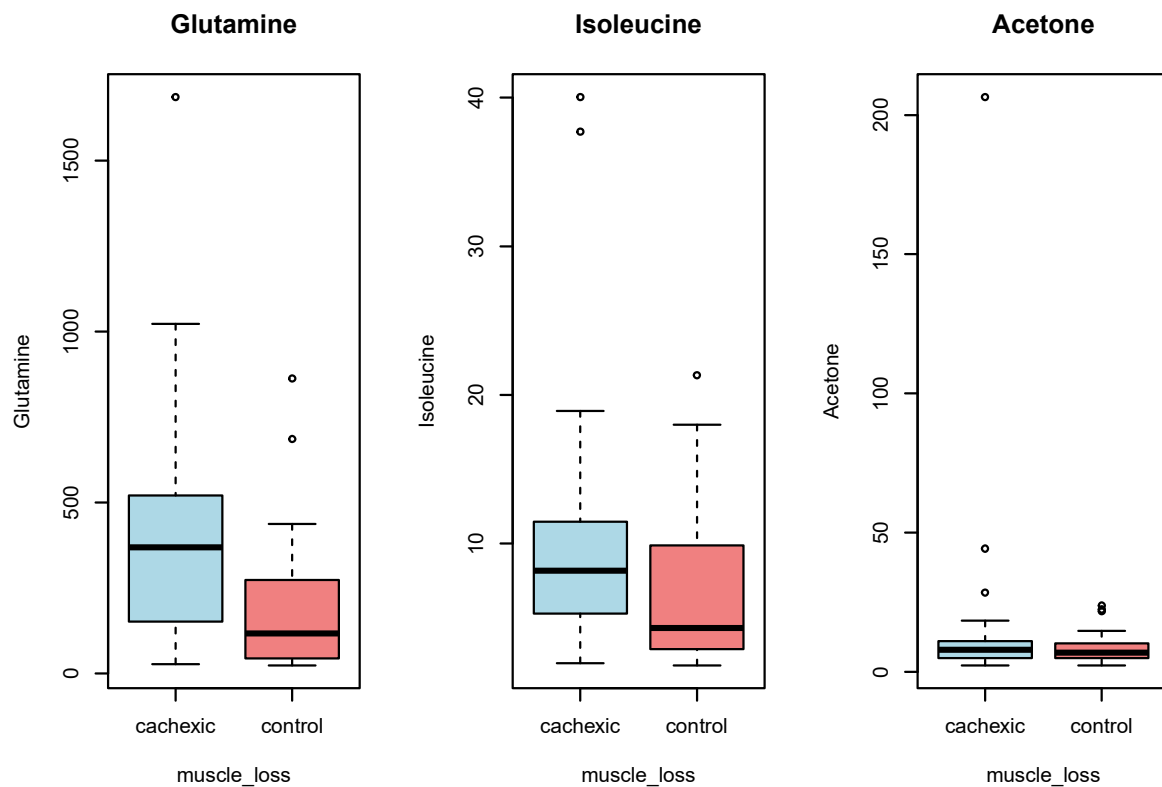
En este rápido análisis hemos obtenido que 55 de las 63 variables presentan diferencias significativas entre los grupos con un nivel de significancia de $p < 0.05$. Sería necesario un análisis en mayor profundidad para comprobar esto con certeza, pero parece que hay diferencias entre los dos grupos de individuos.

Podemos representar gráficamente algunas de las variables. Por ejemplo, representaremos 4 que han resultado significativas en el anterior análisis (Alanine, Creatine, Taurine y Glutamine) y 2 que no (Isoleucine y Acetone).

```
# Seleccionar los metabolitos que vamos a representar
metabolitos_seleccionados <- c("Alanine", "Creatine", "Taurine", "Glutamine", "Isoleucine", "Acetone")

# Crear los boxplot
par(mfrow=c(1, 3))
for (metabolito in metabolitos_seleccionados[1:6]) {
  boxplot(datos_expresion[metabolito, ] ~ muscle_loss,
    main=paste(metabolito),
    ylab=metabolito,
    col=c("lightblue", "lightcoral"))
}
```



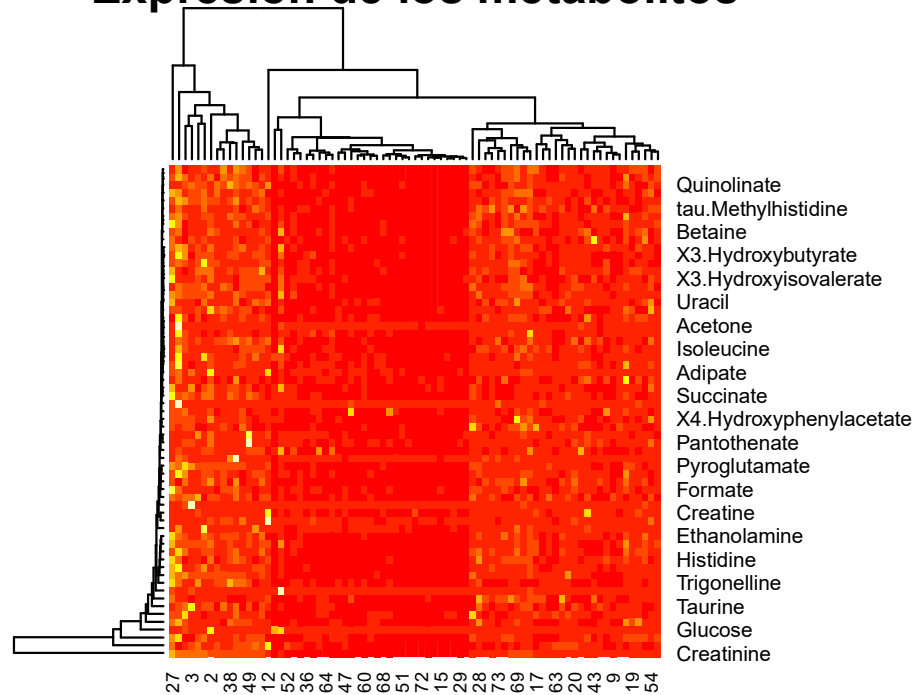


```
par(mfrow=c(1, 2))
```

También se podrían llevar a cabo análisis que estudien las variables en su conjunto, por ejemplo, a través de un heatmap. Realizaremos un heatmap con los datos de expresión de los metabolitos y una matriz de correlación.

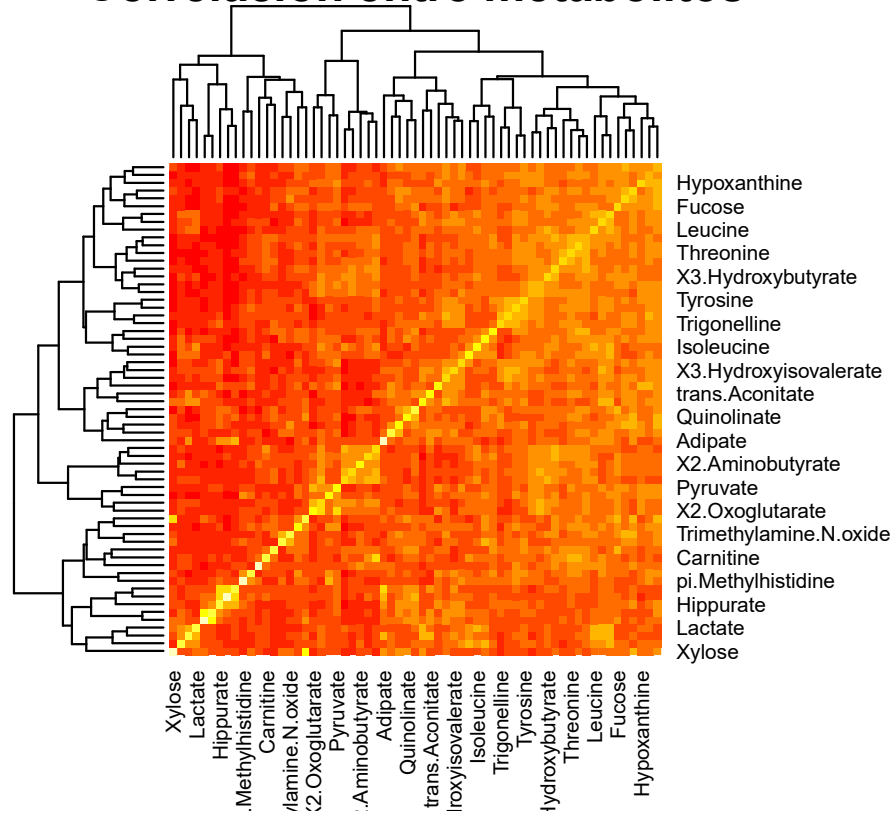
```
# Representar el heatmap
heatmap(datos_expresion, main="Expresión de los metabolitos", col=heat.colors(10), scale="row")
```

Expresión de los metabolitos



```
# Representar la matriz de correlación
cor_metabolitos <- cor(t(datos_expresion), use="pairwise.complete.obs")
heatmap(cor_metabolitos, main="Correlación entre metabolitos", col=heat.colors(10))
```

Correlación entre metabolitos



4. Reposición de los datos en github

Por último, creamos un repositorio de github donde se adjuntaran todos los entregables de esta PEC. En mi caso, corresponde al siguiente enlace: <https://github.com/DanielAcostaBarrios/Acosta-Barrios-Daniel-PEC1>