

# SANS\_FONS\_MGLORIA\_PAC1\_do

2025-04-02

## R Markdown

Anàlisis de les metadades

L'objecte que proporciona la crida està formada alhora per dos objectes “summarizedexperiment”:m1,m2. S'han obtingut per selecció diferencial en l'assay cromatogràfic. Les anàlisis proporcionades són de la selecció amb ió positiu dels metabolits.

L'objecte m1 té 46 mostres que corresponen als 46 ratolins tractats o no amb el tractament immunològic i en diferents dietes que contenen fibra. Es corresponen amb “samples” en l'objecte summarized experiment m1. Té 327 files que corresponen a “features” del objecte summarizedexperiment m1. Mijçant la funció “dimnames” obtenim la numeració de les mostres (samples) i la numeració dels metabolits(features). Amb la funció colData obtenim els tractaments i les condicions del assaig.

```
se=do_query(
  context='study',
  input_item="study_id",
  input_value="ST003789",
  output_item="SummarizedExperiment"
)
assay(se)
```

```
## $AN006228
## class: SummarizedExperiment
## dim: 327 46
## metadata(8): data_source study_id ... description subject_type
## assays(1): ''
## rownames(327): ME1000003 ME1000004 ... ME1000326 ME1000327
## rowData names(3): metabolite_name metabolite_id refmet_name
## colnames(46): 12 14 ... 9 90
## colData names(8): local_sample_id study_id ... Diet Treatment
##
## $AN006229
## class: SummarizedExperiment
## dim: 266 46
## metadata(8): data_source study_id ... description subject_type
## assays(1): ''
## rownames(266): ME1000329 ME1000330 ... ME1000592 ME1000593
## rowData names(3): metabolite_name metabolite_id refmet_name
## colnames(46): 12 14 ... 9 90
## colData names(8): local_sample_id study_id ... Diet Treatment
```

```
m1<-se$AN006228
m1
```

```
## class: SummarizedExperiment
## dim: 327 46
## metadata(8): data_source study_id ... description subject_type
## assays(1): ''
## rownames(327): ME1000003 ME1000004 ... ME1000326 ME1000327
## rowData names(3): metabolite_name metabolite_id refmet_name
## colnames(46): 12 14 ... 9 90
## colData names(8): local_sample_id study_id ... Diet Treatment
```

```
m2<-se$AN006229
m2
```

```
## class: SummarizedExperiment
## dim: 266 46
## metadata(8): data_source study_id ... description subject_type
## assays(1): ''
## rownames(266): ME1000329 ME1000330 ... ME1000592 ME1000593
## rowData names(3): metabolite_name metabolite_id refmet_name
## colnames(46): 12 14 ... 9 90
## colData names(8): local_sample_id study_id ... Diet Treatment
```

```
dim(m1)
```

```
## [1] 327 46
```

```
dimnames(m1)
```

```
## [[1]]
## [1] "ME1000003" "ME1000004" "ME1000005" "ME1000001" "ME1000002" "ME1000006"
## [7] "ME1000007" "ME1000008" "ME1000009" "ME1000010" "ME1000011" "ME1000012"
## [13] "ME1000013" "ME1000014" "ME1000015" "ME1000016" "ME1000017" "ME1000018"
## [19] "ME1000019" "ME1000020" "ME1000021" "ME1000023" "ME1000022" "ME1000024"
## [25] "ME1000025" "ME1000026" "ME1000027" "ME1000028" "ME1000029" "ME1000031"
## [31] "ME1000030" "ME1000032" "ME1000033" "ME1000034" "ME1000035" "ME1000036"
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## [43] "ME1000043" "ME1000044" "ME1000045" "ME1000046" "ME1000047" "ME1000048"
## [49] "ME1000049" "ME1000050" "ME1000051" "ME1000052" "ME1000053" "ME1000054"
## [55] "ME1000055" "ME1000056" "ME1000057" "ME1000058" "ME1000059" "ME1000060"
## [61] "ME1000061" "ME1000062" "ME1000063" "ME1000064" "ME1000065" "ME1000066"
## [67] "ME1000067" "ME1000068" "ME1000069" "ME1000070" "ME1000071" "ME1000072"
## [73] "ME1000073" "ME1000074" "ME1000075" "ME1000076" "ME1000077" "ME1000078"
## [79] "ME1000079" "ME1000080" "ME1000081" "ME1000082" "ME1000083" "ME1000084"
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## [91] "ME1000091" "ME1000092" "ME1000093" "ME1000094" "ME1000095" "ME1000096"
## [97] "ME1000097" "ME1000098" "ME1000099" "ME1000100" "ME1000101" "ME1000102"
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## [115] "ME1000115" "ME1000116" "ME1000117" "ME1000118" "ME1000119" "ME1000120"
## [121] "ME1000121" "ME1000122" "ME1000123" "ME1000124" "ME1000125" "ME1000126"
## [127] "ME1000127" "ME1000128" "ME1000129" "ME1000130" "ME1000131" "ME1000132"
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## [283] "ME1000282" "ME1000283" "ME1000284" "ME1000285" "ME1000286" "ME1000287"
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## [295] "ME1000294" "ME1000295" "ME1000296" "ME1000297" "ME1000298" "ME1000299"
## [301] "ME1000300" "ME1000301" "ME1000302" "ME1000303" "ME1000304" "ME1000305"
## [307] "ME1000306" "ME1000307" "ME1000308" "ME1000309" "ME1000310" "ME1000311"
## [313] "ME1000312" "ME1000313" "ME1000314" "ME1000315" "ME1000316" "ME1000317"
## [319] "ME1000318" "ME1000319" "ME1000320" "ME1000321" "ME1000322" "ME1000323"
## [325] "ME1000324" "ME1000326" "ME1000327"
##
## [[2]]
## [1] "12" "14" "2" "22" "23" "24" "26" "27" "28" "29" "30" "32" "33" "37" "4"
## [16] "40" "41" "43" "44" "45" "46" "49" "50" "52" "55" "56" "57" "6" "60" "61"
## [31] "63" "65" "7" "71" "72" "73" "75" "78" "79" "80" "83" "86" "88" "89" "9"
## [46] "90"
```

```
colData(m1)
```

```
## DataFrame with 46 rows and 8 columns
## local_sample_id study_id sample_source mb_sample_id raw_data
## <character> <character> <character> <character> <character>
## 12 12 ST003789 Serum SA412477 12.MZXML
## 14 14 ST003789 Serum SA412473 14.MZXML
## 2 2 ST003789 Serum SA412472 2.MZXML
## 22 22 ST003789 Serum SA412468 22.MZXML
## 23 23 ST003789 Serum SA412469 23.MZXML
## ... ... ... ...
## 86 86 ST003789 Serum SA412497 86.MZXML
## 88 88 ST003789 Serum SA412498 88.MZXML
## 89 89 ST003789 Serum SA412499 89.MZXML
## 9 9 ST003789 Serum SA412475 9.MZXML
## 90 90 ST003789 Serum SA412500 90.MZXML
## Sample_source Diet Treatment
## <factor> <factor> <factor>
## 12 Serum Chow isotype control
## 14 Serum Chow isotype control
## 2 Serum Chow isotype control
## 22 Serum Chow anti-PD-1
## 23 Serum Chow anti-PD-1
## ... ... ...
## 86 Serum HC_In8 anti-PD-1
## 88 Serum HC_In8 anti-PD-1
## 89 Serum HC_In8 anti-PD-1
## 9 Serum Chow isotype control
## 90 Serum HC_In8 anti-PD-1
```