

Ejemplos dplyr

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Load data

```
library(readr)

brc <- read.csv("breast-cancer.data", header=FALSE)
names <- read.csv("breast-cancer.names1.csv", header=FALSE)

names(brc) <- as.character(names$V1)
names(brc)

## [1] "Class"      "age"        "menopause"  "tumor_size" "inv_nodes"
## [6] "node_caps"  "deg_malig"  "breast"     "breast_quad" "irradiat"
```

Funciones principales de dplyr:

```
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
```

5 funciones básicas:

- filtrar filas del *dataset*: `filter`
- ordenar *dataset* seg?n ciertas variables : `arrange - top_n`
- seleccionar columnas: `select`
- crear nuevas columnas: `mutate`
- sumarizaci?n del *dataset*: `summarise`

Funcionamiento:

- 1 arg: dataset
- resto args: dice qu? hacer
- nombre columnas nunca con comillas
- resultado es un `data.frame`

Filter

```
# knitr::kable(filter(brc, deg_malig == 3), longtable = TRUE)
library(knitr)
kable(head(filter(brc, deg_malig == 3)), longtable = TRUE)
```

Class	age	menopausa	tumor_size	inv_nodes	node_caps	deg_malig	breast	breast_qual	irradiat
no-recurrence-events	30-39	premeno	30-34	0-2	no	3	left	left_low	no
no-recurrence-events	40-49	premeno	0-4	0-2	no	3	left	central	no
no-recurrence-events	50-59	ge40	25-29	0-2	no	3	left	right_up	no
no-recurrence-events	40-49	premeno	30-34	0-2	no	3	left	left_up	no
no-recurrence-events	50-59	premeno	30-34	0-2	no	3	left	left_low	no
no-recurrence-events	60-69	ge40	30-34	0-2	no	3	left	left_low	no

```
kable(head(filter(brc, deg_malig == 3, irradiat == "no")), longtable = TRUE)
```

Class	age	menopausa	tumor_size	inv_nodes	node_caps	deg_malig	breast	breast_qual	irradiat
no-recurrence-events	30-39	premeno	30-34	0-2	no	3	left	left_low	no
no-recurrence-events	40-49	premeno	0-4	0-2	no	3	left	central	no
no-recurrence-events	50-59	ge40	25-29	0-2	no	3	left	right_up	no
no-recurrence-events	40-49	premeno	30-34	0-2	no	3	left	left_up	no
no-recurrence-events	50-59	premeno	30-34	0-2	no	3	left	left_low	no
no-recurrence-events	60-69	ge40	30-34	0-2	no	3	left	left_low	no

```
kable(head(filter(brc, deg_malig == 3 & irradiat == "no" | breast == "left")), longtable = TRUE)
```

Class	age	menopaus	tumor_size	inv_nodes	node_caps	leg_malign	breast	breast_qual	irradiat
no-recurrence-events	30-39	premeno	30-34	0-2	no	3	left	left_low	no
no-recurrence-events	40-49	premeno	20-24	0-2	no	2	left	left_low	no
no-recurrence-events	60-69	ge40	15-19	0-2	no	2	left	left_low	no
no-recurrence-events	50-59	premeno	25-29	0-2	no	2	left	left_low	no
no-recurrence-events	60-69	ge40	20-24	0-2	no	1	left	left_low	no
no-recurrence-events	40-49	premeno	50-54	0-2	no	2	left	left_low	no

Ordenar *dataset*

```
brc1 <- filter(brc, deg_malign == 3 & irradiat == "no" | breast == "left")
kable(head(arrange(brc1, age, tumor_size)), longtable = TRUE)
```

Class	age	menopaus	tumor_size	inv_nodes	node_caps	leg_malign	breast	breast_qual	irradiat
no-recurrence-events	30-39	premeno	10-14	0-2	no	2	left	right_low	no
no-recurrence-events	30-39	premeno	15-19	0-2	no	1	left	left_low	no
no-recurrence-events	30-39	lt40	15-19	0-2	no	3	right	left_up	no
no-recurrence-events	30-39	premeno	15-19	0-2	no	1	left	left_low	no
recurrence-events	30-39	premeno	15-19	6-8	yes	3	left	left_low	yes
no-recurrence-events	30-39	premeno	20-24	0-2	no	2	left	right_low	no

```
kable(head(arrange(brc1, desc(age), tumor_size)), longtable = TRUE)
```

Class	age	menopaus	tumor_size	inv_nodes	node_caps	leg_malign	breast	breast_qual	irradiat
no-recurrence-events	70-79	ge40	0-4	0-2	no	1	left	right_low	no
no-recurrence-events	70-79	ge40	10-14	0-2	no	2	left	central	no
recurrence-events	70-79	ge40	15-19	9-11	?	1	left	left_low	yes
no-recurrence-events	70-79	ge40	20-24	0-2	no	3	left	left_up	no
no-recurrence-events	60-69	lt40	10-14	0-2	no	1	left	right_up	no

Class	age	menopause	tumor_size	inv_nodes	node_caps	leg_malign	breast	breast_quad	irradiat
no-recurrence-events	60-69	ge40	10-14	0-2	no	1	left	left_low	no

Ejercicio:

Extraer los 10 pacientes con menopausia con mayor tamaño del tumor y menor número de nodos invasores ordenados por edad:

```
brc2 <- filter(brc, menopause == "premeno")
# head(arrange(brc2, age, desc(tumor_size), inv_nodes), n=10)
kable((arrange(brc2, age, desc(tumor_size), inv_nodes))[1:10, ], longtable = TRUE)
```

Class	age	menopause	tumor_size	inv_nodes	node_caps	leg_malign	breast	breast_quad	irradiat
no-recurrence-events	20-29	premeno	35-39	0-2	no	2	right	right_up	no
no-recurrence-events	30-39	premeno	5-9	0-2	no	2	left	right_low	no
no-recurrence-events	30-39	premeno	40-44	0-2	no	2	right	right_up	no
no-recurrence-events	30-39	premeno	40-44	0-2	no	2	left	left_low	yes
recurrence-events	30-39	premeno	40-44	0-2	no	1	left	left_up	no
no-recurrence-events	30-39	premeno	40-44	3-5	no	3	right	right_up	yes
recurrence-events	30-39	premeno	35-39	0-2	no	3	left	left_low	no
recurrence-events	30-39	premeno	35-39	0-2	no	3	left	left_low	no
recurrence-events	30-39	premeno	35-39	9-11	yes	3	left	left_low	no
no-recurrence-events	30-39	premeno	30-34	0-2	no	3	left	left_low	no

top_n()

```
brc1 <- filter(brc, age == "40-49")
top_n(brc1, 1)
```

Selecting by irradiat

```
##
##          Class  age menopause tumor_size inv_nodes node_caps
## 1 no-recurrence-events 40-49  premeno    35-39    9-11      yes
## 2 no-recurrence-events 40-49  premeno    35-39    9-11      yes
## 3 no-recurrence-events 40-49  premeno    40-44    3-5       yes
```

```

## 4  no-recurrence-events 40-49  premeno      5-9      0-2      no
## 5  no-recurrence-events 40-49  premeno     45-49     0-2      no
## 6  no-recurrence-events 40-49  premeno     25-29     0-2      ?
## 7  no-recurrence-events 40-49  premeno     25-29     0-2      no
## 8  no-recurrence-events 40-49    ge40     40-44    15-17     yes
## 9  no-recurrence-events 40-49    ge40     30-34     0-2      no
## 10 no-recurrence-events 40-49  premeno     30-34     0-2      no
## 11 no-recurrence-events 40-49  premeno     20-24     0-2      no
## 12 no-recurrence-events 40-49  premeno     35-39     0-2     yes
## 13 no-recurrence-events 40-49  premeno     35-39     0-2     yes
## 14 no-recurrence-events 40-49  premeno     25-29     0-2      no
## 15 no-recurrence-events 40-49  premeno     20-24     6-8      no
## 16 no-recurrence-events 40-49  premeno     15-19    12-14     no
## 17 no-recurrence-events 40-49  premeno     25-29     0-2      no
## 18 no-recurrence-events 40-49  premeno     10-14     0-2      no
## 19  recurrence-events 40-49    ge40     20-24     3-5      no
## 20  recurrence-events 40-49  premeno     20-24     3-5     yes
## 21  recurrence-events 40-49  premeno     30-34    12-14     yes
## 22  recurrence-events 40-49  premeno     50-54     0-2      no
## 23  recurrence-events 40-49  premeno     30-34     0-2      no
## 24  recurrence-events 40-49  premeno     20-24     3-5     yes
## 25  recurrence-events 40-49    ge40     25-29    12-14     yes
## 26  recurrence-events 40-49  premeno     25-29     0-2      no
##    deg_malign breast breast_quad irradiat
## 1      2  right    left_up    yes
## 2      2  right    right_up   yes
## 3      3  right    left_up    yes
## 4      1  left     left_low   yes
## 5      2  left     left_low   yes
## 6      2  left     right_low  yes
## 7      3  right    left_up    yes
## 8      2  right    left_up    yes
## 9      2  left     left_up    yes
## 10     2  right    right_up   yes
## 11     3  right    left_low   yes
## 12     3  right    left_up    yes
## 13     3  right    left_low   yes
## 14     1  right    left_low   yes
## 15     2  right    left_low   yes
## 16     3  right    right_low  yes
## 17     2  left     left_up    yes
## 18     2  left     left_low   yes
## 19     3  right    left_low   yes
## 20     2  right    right_up   yes
## 21     3  left     left_up    yes
## 22     2  right    left_low   yes
## 23     1  left     left_low   yes
## 24     2  left     left_low   yes
## 25     3  left     right_low  yes
## 26     2  left     left_low   yes

```

```
head(top_n(brc1, -1))
```

```
## Selecting by irradiat
```

```
##           Class   age menopause tumor_size inv_nodes node_caps deg_malign
## 1 no-recurrence-events 40-49   premeno    20-24      0-2      no      2
## 2 no-recurrence-events 40-49   premeno    20-24      0-2      no      2
## 3 no-recurrence-events 40-49   premeno      0-4      0-2      no      2
## 4 no-recurrence-events 40-49   premeno    50-54      0-2      no      2
## 5 no-recurrence-events 40-49   premeno    20-24      0-2      no      2
## 6 no-recurrence-events 40-49   premeno      0-4      0-2      no      3
##   breast breast_quad irradiat
## 1   right   right_up      no
## 2    left   left_low      no
## 3   right   right_low     no
## 4    left   left_low      no
## 5   right   left_up      no
## 6    left   central      no
```

Seleccionar columnas

- `starts_with()`
- `ends_with()`

```
head(select(brc, age, tumor_size))
```

```
##      age tumor_size
## 1 30-39      30-34
## 2 40-49      20-24
## 3 40-49      20-24
## 4 60-69      15-19
## 5 40-49       0-4
## 6 60-69      15-19
```

```
kable(head(select(brc, breast, breast_quad, everything())))
```

breast	breast_quad	Class	age	menopause	tumor_size	inv_nodes	node_caps	deg_malign	irradiat
left	left_low	no-recurrence-events	30-39	premeno	30-34	0-2	no	3	no
right	right_up	no-recurrence-events	40-49	premeno	20-24	0-2	no	2	no
left	left_low	no-recurrence-events	40-49	premeno	20-24	0-2	no	2	no
right	left_up	no-recurrence-events	60-69	ge40	15-19	0-2	no	2	no
right	right_low	no-recurrence-events	40-49	premeno	0-4	0-2	no	2	no
left	left_low	no-recurrence-events	60-69	ge40	15-19	0-2	no	2	no

```
head(select(brc, Class:inv_nodes)) # desde:hasta
```

```
##           Class   age menopause tumor_size inv_nodes
```

```
## 1 no-recurrence-events 30-39 premeno 30-34 0-2
## 2 no-recurrence-events 40-49 premeno 20-24 0-2
## 3 no-recurrence-events 40-49 premeno 20-24 0-2
## 4 no-recurrence-events 60-69 ge40 15-19 0-2
## 5 no-recurrence-events 40-49 premeno 0-4 0-2
## 6 no-recurrence-events 60-69 ge40 15-19 0-2
```

```
head(select(brc, contains("_")))
```

```
##      tumor_size inv_nodes node_caps deg_malign breast_quad
## 1      30-34      0-2      no      3      left_low
## 2      20-24      0-2      no      2      right_up
## 3      20-24      0-2      no      2      left_low
## 4      15-19      0-2      no      2      left_up
## 5       0-4      0-2      no      2      right_low
## 6      15-19      0-2      no      2      left_low
```

```
head(select(brc, starts_with("a")))
```

```
##      age
## 1 30-39
## 2 40-49
## 3 40-49
## 4 60-69
## 5 40-49
## 6 60-69
```

```
br1 <- select(brc, breast, breast_quad, everything())
br2 <- select(br1, -c(Class, irradiat));
head(br1)
```

```
##      breast breast_quad      Class      age menopause tumor_size inv_nodes
## 1    left    left_low no-recurrence-events 30-39    premeno    30-34      0-2
## 2    right    right_up no-recurrence-events 40-49    premeno    20-24      0-2
## 3    left    left_low no-recurrence-events 40-49    premeno    20-24      0-2
## 4    right    left_up no-recurrence-events 60-69      ge40    15-19      0-2
## 5    right    right_low no-recurrence-events 40-49    premeno      0-4      0-2
## 6    left    left_low no-recurrence-events 60-69      ge40    15-19      0-2
##      node_caps deg_malign irradiat
## 1         no      3         no
## 2         no      2         no
## 3         no      2         no
## 4         no      2         no
## 5         no      2         no
## 6         no      2         no
```

```
head(br2)
```

```
##      breast breast_quad      age menopause tumor_size inv_nodes node_caps deg_malign
## 1    left    left_low 30-39    premeno    30-34      0-2      no      3
## 2    right    right_up 40-49    premeno    20-24      0-2      no      2
```

```
## 3 left left_low 40-49 premeno 20-24 0-2 no 2
## 4 right left_up 60-69 ge40 15-19 0-2 no 2
## 5 right right_low 40-49 premeno 0-4 0-2 no 2
## 6 left left_low 60-69 ge40 15-19 0-2 no 2
```

mutate():

```
br1 <- mutate(brc, distancia = 4 - deg_malig);
head(br1)
```

```
##           Class age menopause tumor_size inv_nodes node_caps deg_malig
## 1 no-recurrence-events 30-39 premeno 30-34 0-2 no 3
## 2 no-recurrence-events 40-49 premeno 20-24 0-2 no 2
## 3 no-recurrence-events 40-49 premeno 20-24 0-2 no 2
## 4 no-recurrence-events 60-69 ge40 15-19 0-2 no 2
## 5 no-recurrence-events 40-49 premeno 0-4 0-2 no 2
## 6 no-recurrence-events 60-69 ge40 15-19 0-2 no 2
## breast breast_quad irradiat distancia
## 1 left left_low no 1
## 2 right right_up no 2
## 3 left left_low no 2
## 4 right left_up no 2
## 5 right right_low no 2
## 6 left left_low no 2
```

rename():

```
br2 <- rename(br1, dist = distancia);
head(br2)
```

```
##           Class age menopause tumor_size inv_nodes node_caps deg_malig
## 1 no-recurrence-events 30-39 premeno 30-34 0-2 no 3
## 2 no-recurrence-events 40-49 premeno 20-24 0-2 no 2
## 3 no-recurrence-events 40-49 premeno 20-24 0-2 no 2
## 4 no-recurrence-events 60-69 ge40 15-19 0-2 no 2
## 5 no-recurrence-events 40-49 premeno 0-4 0-2 no 2
## 6 no-recurrence-events 60-69 ge40 15-19 0-2 no 2
## breast breast_quad irradiat dist
## 1 left left_low no 1
## 2 right right_up no 2
## 3 left left_low no 2
## 4 right left_up no 2
## 5 right right_low no 2
## 6 left left_low no 2
```

transmute():


```
head(transmute(brc2, test = deg_malig / 61.0237))
```

```
##          test
## 1 0.04916123
## 2 0.03277415
## 3 0.03277415
## 4 0.03277415
## 5 0.03277415
## 6 0.03277415
```

summarize():

Realiza un resumen de una estadística y guardarlos en un *dataframe*:

```
summarise(brc, a = mean(brc$deg_malig))
```

```
##          a
## 1 2.048951
```

```
summarise(brc, a = median(brc$deg_malig))
```

```
##          a
## 1 2
```

Agregación de datos

Agrega datos de varias columnas:

```
data("mtcars")
aggdata <- aggregate(mtcars, by=list(mtcars$cyl, mtcars$vs), FUN=mean, na.rm=TRUE)
print(aggdata)
```

```
##   Group.1 Group.2      mpg cyl  disp    hp  drat    wt    qsec vs
## 1      4      0 26.00000   4 120.30  91.0000 4.430000 2.140000 16.70000 0
## 2      6      0 20.56667   6 155.00 131.6667 3.806667 2.755000 16.32667 0
## 3      8      0 15.10000   8 353.10 209.2143 3.229286 3.999214 16.77214 0
## 4      4      1 26.73000   4 103.62  81.8000 4.035000 2.300300 19.38100 1
## 5      6      1 19.12500   6 204.55 115.2500 3.420000 3.388750 19.21500 1
##          am      gear      carb
## 1 1.0000000 5.000000 2.000000
## 2 1.0000000 4.333333 4.666667
## 3 0.1428571 3.285714 3.500000
## 4 0.7000000 4.000000 1.500000
## 5 0.0000000 3.500000 2.500000
```

pipes

`x %>% f(y) ≡ f(x, y)`

Por ejemplo, `brc %>% filter(age=="40-49") ≡ filter(brc, age=="40-49")`

```
head(filter(brc, age=="40-49"))
```

```
##           Class   age  menopause tumor_size inv_nodes node_caps deg_malign
## 1 no-recurrence-events 40-49   premeno    20-24      0-2      no      2
## 2 no-recurrence-events 40-49   premeno    20-24      0-2      no      2
## 3 no-recurrence-events 40-49   premeno      0-4      0-2      no      2
## 4 no-recurrence-events 40-49   premeno    50-54      0-2      no      2
## 5 no-recurrence-events 40-49   premeno    20-24      0-2      no      2
## 6 no-recurrence-events 40-49   premeno      0-4      0-2      no      3
##   breast breast_quad irradiat
## 1   right   right_up      no
## 2    left   left_low      no
## 3   right   right_low      no
## 4    left   left_low      no
## 5   right   left_up      no
## 6    left   central      no
```

```
brc %>%
  filter(age=="40-49") %>%
  select(tumor_size) %>%
  top_n(5)
```

Selecting by tumor_size

```
##   tumor_size
## 1      50-54
## 2      40-44
## 3      40-44
## 4      40-44
## 5         5-9
## 6      45-49
## 7      40-44
## 8      40-44
## 9      50-54
```

```
top_n(select(filter(brc, age=="40-49"), tumor_size), 5)
```

Selecting by tumor_size

```
##   tumor_size
## 1      50-54
## 2      40-44
## 3      40-44
## 4      40-44
## 5         5-9
## 6      45-49
## 7      40-44
## 8      40-44
## 9      50-54
```

`group_by()`:

```
brc %>% group_by(age)
```

```
## # A tibble: 286 x 10
## # Groups:   age [6]
##   Class age  menopause tumor_size inv_nodes node_caps deg_malig breast
##   <chr> <chr> <chr>      <chr>      <chr>      <chr>      <int> <chr>
## 1 no-r~ 30-39 premeno   30-34      0-2        no          3 left
## 2 no-r~ 40-49 premeno   20-24      0-2        no          2 right
## 3 no-r~ 40-49 premeno   20-24      0-2        no          2 left
## 4 no-r~ 60-69 ge40     15-19      0-2        no          2 right
## 5 no-r~ 40-49 premeno   0-4        0-2        no          2 right
## 6 no-r~ 60-69 ge40     15-19      0-2        no          2 left
## 7 no-r~ 50-59 premeno   25-29      0-2        no          2 left
## 8 no-r~ 60-69 ge40     20-24      0-2        no          1 left
## 9 no-r~ 40-49 premeno   50-54      0-2        no          2 left
## 10 no-r~ 40-49 premeno   20-24      0-2        no          2 right
## # ... with 276 more rows, and 2 more variables: breast_quad <chr>,
## #   irradiat <chr>
```

Podemos usar `group_by()` junto con `summarise()`:

```
brc %>%
  group_by(age) %>%
  summarise(mean_deg_malig=mean(deg_malig))
```

```
## 'summarise()' ungrouping output (override with '.groups' argument)
```

```
## # A tibble: 6 x 2
##   age  mean_deg_malig
##   <chr>          <dbl>
## 1 20-29            2
## 2 30-39          2.14
## 3 40-49          2.07
## 4 50-59          2.07
## 5 60-69          1.98
## 6 70-79          1.5
```

```
brc %>%
  group_by(age,tumor_size) %>%
  summarise(mean_deg_malig=mean(deg_malig),
            sd_deg_malig=sd(deg_malig))
```

```
## 'summarise()' regrouping output by 'age' (override with '.groups' argument)
```

```
## # A tibble: 46 x 4
## # Groups:   age [6]
##   age  tumor_size mean_deg_malig sd_deg_malig
##   <chr> <chr>          <dbl>          <dbl>
```

##	1	20-29	35-39	2	NA
##	2	30-39	0-4	2	0
##	3	30-39	10-14	1.5	0.707
##	4	30-39	15-19	1.8	1.10
##	5	30-39	20-24	2.33	0.516
##	6	30-39	25-29	2.17	0.753
##	7	30-39	30-34	2.14	0.690
##	8	30-39	35-39	3	0
##	9	30-39	40-44	2	0.816
##	10	30-39	5-9	2	NA
##	#	... with 36 more rows			