

dataframe basic manipulation

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dataframe is a collection of different class of data

Create a dataframe

```
quarters<-rep(paste('q',seq=(1:4),sep=''),3)
year<-rep(seq(2000,2002),each=4)
mydataframe<-cbind.data.frame(year,quarters)
mydataframe
```

```
##   year quarters
## 1  2000      q1
## 2  2000      q2
## 3  2000      q3
## 4  2000      q4
## 5  2001      q1
## 6  2001      q2
## 7  2001      q3
## 8  2001      q4
## 9  2002      q1
## 10 2002      q2
## 11 2002      q3
## 12 2002      q4
```

access each column by calling the column name

```
mydataframe$year
```

```
## [1] 2000 2000 2000 2000 2001 2001 2001 2001 2002 2002 2002 2002
```

```
mydataframe$quarters
```

```
## [1] q1 q2 q3 q4 q1 q2 q3 q4 q1 q2 q3 q4
## Levels: q1 q2 q3 q4
```

access each column by indexing location

```
mydataframe[,1]
```

```
## [1] 2000 2000 2000 2000 2001 2001 2001 2001 2002 2002 2002 2002
```

```
mydataframe[,2]
```

```
## [1] q1 q2 q3 q4 q1 q2 q3 q4 q1 q2 q3 q4
## Levels: q1 q2 q3 q4
```

access rows by indexing location

```
mydataframe[1:5,] # first 5 rows
```

```
##   year quarters
## 1 2000       q1
## 2 2000       q2
## 3 2000       q3
## 4 2000       q4
## 5 2001       q1
```

select rows based on column values, using slicing, or subset() function

```
mydataframe[quarters=='q1',]
```

```
##   year quarters
## 1 2000       q1
## 5 2001       q1
## 9 2002       q1
```

```
subset(mydataframe,quarters=='q1')
```

```
##   year quarters
## 1 2000       q1
## 5 2001       q1
## 9 2002       q1
```

using logic operator |(or) &(and)

```
mydataframe[quarters=='q1'|quarters=='q2',]
```

```
##   year quarters
## 1 2000       q1
## 2 2000       q2
## 5 2001       q1
## 6 2001       q2
## 9 2002       q1
## 10 2002      q2
```

```
mydataframe[quarters=='q1'&year==2000,]
```

```
##   year quarters
## 1 2000       q1
```

how to change row / column names:

```
row.names(mydataframe)
```

```
## [1] "1" "2" "3" "4" "5" "6" "7" "8" "9" "10" "11" "12"
```

```
row.names(mydataframe)<-letters[1:12]
row.names(mydataframe)
```

```
## [1] "a" "b" "c" "d" "e" "f" "g" "h" "i" "j" "k" "l"
```

```
colnames(mydataframe)
```

```
## [1] "year"      "quarters"
```

```
colnames(mydataframe)[1]<- 'YEAR' #change the first column's name
mydataframe
```

```
##   YEAR quarters
## a 2000      q1
## b 2000      q2
## c 2000      q3
## d 2000      q4
## e 2001      q1
## f 2001      q2
## g 2001      q3
## h 2001      q4
## i 2002      q1
## j 2002      q2
## k 2002      q3
## l 2002      q4
```

create a new column in a dataframe:

```
set.seed(1)
mydataframe$newcolumn1<-rnorm(12,mean = 0,sd = 1)
mydataframe$newcolumn2<-runif(12,min = 0,max = 1)
mydataframe
```

```
##   YEAR quarters newcolumn1 newcolumn2
## a 2000      q1 -0.6264538 0.26722067
## b 2000      q2  0.1836433 0.38611409
## c 2000      q3 -0.8356286 0.01339033
## d 2000      q4  1.5952808 0.38238796
## e 2001      q1  0.3295078 0.86969085
## f 2001      q2 -0.8204684 0.34034900
## g 2001      q3  0.4874291 0.48208012
## h 2001      q4  0.7383247 0.59956583
## i 2002      q1  0.5757814 0.49354131
## j 2002      q2 -0.3053884 0.18621760
## k 2002      q3  1.5117812 0.82737332
## l 2002      q4  0.3898432 0.66846674
```

if our newcolumn3 is newcolumn1+newcolumn2

```
mydataframe$newcolumn3<-mydataframe$newcolumn1+mydataframe$newcolumn2
mydataframe
```

```
##   YEAR quarters newcolumn1 newcolumn2 newcolumn3
## a 2000      q1 -0.6264538 0.26722067 -0.3592331
## b 2000      q2  0.1836433 0.38611409  0.5697574
## c 2000      q3 -0.8356286 0.01339033 -0.8222383
## d 2000      q4  1.5952808 0.38238796  1.9776688
## e 2001      q1  0.3295078 0.86969085  1.1991986
## f 2001      q2 -0.8204684 0.34034900 -0.4801194
## g 2001      q3  0.4874291 0.48208012  0.9695092
## h 2001      q4  0.7383247 0.59956583  1.3378905
## i 2002      q1  0.5757814 0.49354131  1.0693227
## j 2002      q2 -0.3053884 0.18621760 -0.1191708
## k 2002      q3  1.5117812 0.82737332  2.3391545
## l 2002      q4  0.3898432 0.66846674  1.0583100
```

if our newcolumn4 is an index column : when newcolumn1>0, it is 1, otherwise 0

```
mydataframe$newcolumn5<- ifelse(mydataframe$newcolumn1>0,1,0)
mydataframe
```

```
##   YEAR quarters newcolumn1 newcolumn2 newcolumn3 newcolumn5
## a 2000      q1 -0.6264538 0.26722067 -0.3592331          0
## b 2000      q2  0.1836433 0.38611409  0.5697574          1
## c 2000      q3 -0.8356286 0.01339033 -0.8222383          0
## d 2000      q4  1.5952808 0.38238796  1.9776688          1
## e 2001      q1  0.3295078 0.86969085  1.1991986          1
## f 2001      q2 -0.8204684 0.34034900 -0.4801194          0
## g 2001      q3  0.4874291 0.48208012  0.9695092          1
## h 2001      q4  0.7383247 0.59956583  1.3378905          1
## i 2002      q1  0.5757814 0.49354131  1.0693227          1
## j 2002      q2 -0.3053884 0.18621760 -0.1191708          0
## k 2002      q3  1.5117812 0.82737332  2.3391545          1
## l 2002      q4  0.3898432 0.66846674  1.0583100          1
```

Here we introduce how to two data manipulation packages in R

dplyr *tidyr* ##### install packages and library the packages

```
install.packages('dplyr')
install.packages('tidyr')
library.packages('dplyr')
library.packages('tidyr')
```

what if we have a list of packages needs to install and library? Someone posted in Github <https://gist.github.com/stevenworthington/3178163>

```
ipak <- function(pkg){
  new.pkg <- pkg[!(pkg %in% installed.packages()[, "Package"])]
  if (length(new.pkg))
```

```

      install.packages(new.pkg, dependencies = TRUE)
    sapply(pkg, require, character.only = TRUE)
  }

```

usage

```

packages_toinstall <- c("ggplot2", "plyr", "reshape2", "RColorBrewer", "scales", "grid", 'dplyr', 'tidyr')
ipak(packages_toinstall)

```

library multiple packages

```

packages_toload<-c("ggplot2", "plyr", "reshape2", "RColorBrewer", "scales", "grid")
lapply(packages_toload, require, character.only = TRUE)

```

```

packages_toload<-c('dplyr','tidyr')
lapply(packages_toload,require,character.only=TRUE)

```

```

## [[1]]
## [1] TRUE
##
## [[2]]
## [1] TRUE

```

piping

using the output of the previous step as the input of the current step

```
input %>% function1 () %>% function2 ()
```

```
#equal to : function2(function1(input))
```

```
iris
```

```

##      Sepal.Length Sepal.Width Petal.Length Petal.Width   Species
## 1         5.1         3.5         1.4         0.2     setosa
## 2         4.9         3.0         1.4         0.2     setosa
## 3         4.7         3.2         1.3         0.2     setosa
## 4         4.6         3.1         1.5         0.2     setosa
## 5         5.0         3.6         1.4         0.2     setosa
## 6         5.4         3.9         1.7         0.4     setosa
## 7         4.6         3.4         1.4         0.3     setosa
## 8         5.0         3.4         1.5         0.2     setosa
## 9         4.4         2.9         1.4         0.2     setosa
## 10        4.9         3.1         1.5         0.1     setosa
## 11        5.4         3.7         1.5         0.2     setosa
## 12        4.8         3.4         1.6         0.2     setosa
## 13        4.8         3.0         1.4         0.1     setosa
## 14        4.3         3.0         1.1         0.1     setosa
## 15        5.8         4.0         1.2         0.2     setosa
## 16        5.7         4.4         1.5         0.4     setosa
## 17        5.4         3.9         1.3         0.4     setosa
## 18        5.1         3.5         1.4         0.3     setosa

```

## 19	5.7	3.8	1.7	0.3	setosa
## 20	5.1	3.8	1.5	0.3	setosa
## 21	5.4	3.4	1.7	0.2	setosa
## 22	5.1	3.7	1.5	0.4	setosa
## 23	4.6	3.6	1.0	0.2	setosa
## 24	5.1	3.3	1.7	0.5	setosa
## 25	4.8	3.4	1.9	0.2	setosa
## 26	5.0	3.0	1.6	0.2	setosa
## 27	5.0	3.4	1.6	0.4	setosa
## 28	5.2	3.5	1.5	0.2	setosa
## 29	5.2	3.4	1.4	0.2	setosa
## 30	4.7	3.2	1.6	0.2	setosa
## 31	4.8	3.1	1.6	0.2	setosa
## 32	5.4	3.4	1.5	0.4	setosa
## 33	5.2	4.1	1.5	0.1	setosa
## 34	5.5	4.2	1.4	0.2	setosa
## 35	4.9	3.1	1.5	0.2	setosa
## 36	5.0	3.2	1.2	0.2	setosa
## 37	5.5	3.5	1.3	0.2	setosa
## 38	4.9	3.6	1.4	0.1	setosa
## 39	4.4	3.0	1.3	0.2	setosa
## 40	5.1	3.4	1.5	0.2	setosa
## 41	5.0	3.5	1.3	0.3	setosa
## 42	4.5	2.3	1.3	0.3	setosa
## 43	4.4	3.2	1.3	0.2	setosa
## 44	5.0	3.5	1.6	0.6	setosa
## 45	5.1	3.8	1.9	0.4	setosa
## 46	4.8	3.0	1.4	0.3	setosa
## 47	5.1	3.8	1.6	0.2	setosa
## 48	4.6	3.2	1.4	0.2	setosa
## 49	5.3	3.7	1.5	0.2	setosa
## 50	5.0	3.3	1.4	0.2	setosa
## 51	7.0	3.2	4.7	1.4	versicolor
## 52	6.4	3.2	4.5	1.5	versicolor
## 53	6.9	3.1	4.9	1.5	versicolor
## 54	5.5	2.3	4.0	1.3	versicolor
## 55	6.5	2.8	4.6	1.5	versicolor
## 56	5.7	2.8	4.5	1.3	versicolor
## 57	6.3	3.3	4.7	1.6	versicolor
## 58	4.9	2.4	3.3	1.0	versicolor
## 59	6.6	2.9	4.6	1.3	versicolor
## 60	5.2	2.7	3.9	1.4	versicolor
## 61	5.0	2.0	3.5	1.0	versicolor
## 62	5.9	3.0	4.2	1.5	versicolor
## 63	6.0	2.2	4.0	1.0	versicolor
## 64	6.1	2.9	4.7	1.4	versicolor
## 65	5.6	2.9	3.6	1.3	versicolor
## 66	6.7	3.1	4.4	1.4	versicolor
## 67	5.6	3.0	4.5	1.5	versicolor
## 68	5.8	2.7	4.1	1.0	versicolor
## 69	6.2	2.2	4.5	1.5	versicolor
## 70	5.6	2.5	3.9	1.1	versicolor
## 71	5.9	3.2	4.8	1.8	versicolor
## 72	6.1	2.8	4.0	1.3	versicolor

## 73	6.3	2.5	4.9	1.5 versicolor
## 74	6.1	2.8	4.7	1.2 versicolor
## 75	6.4	2.9	4.3	1.3 versicolor
## 76	6.6	3.0	4.4	1.4 versicolor
## 77	6.8	2.8	4.8	1.4 versicolor
## 78	6.7	3.0	5.0	1.7 versicolor
## 79	6.0	2.9	4.5	1.5 versicolor
## 80	5.7	2.6	3.5	1.0 versicolor
## 81	5.5	2.4	3.8	1.1 versicolor
## 82	5.5	2.4	3.7	1.0 versicolor
## 83	5.8	2.7	3.9	1.2 versicolor
## 84	6.0	2.7	5.1	1.6 versicolor
## 85	5.4	3.0	4.5	1.5 versicolor
## 86	6.0	3.4	4.5	1.6 versicolor
## 87	6.7	3.1	4.7	1.5 versicolor
## 88	6.3	2.3	4.4	1.3 versicolor
## 89	5.6	3.0	4.1	1.3 versicolor
## 90	5.5	2.5	4.0	1.3 versicolor
## 91	5.5	2.6	4.4	1.2 versicolor
## 92	6.1	3.0	4.6	1.4 versicolor
## 93	5.8	2.6	4.0	1.2 versicolor
## 94	5.0	2.3	3.3	1.0 versicolor
## 95	5.6	2.7	4.2	1.3 versicolor
## 96	5.7	3.0	4.2	1.2 versicolor
## 97	5.7	2.9	4.2	1.3 versicolor
## 98	6.2	2.9	4.3	1.3 versicolor
## 99	5.1	2.5	3.0	1.1 versicolor
## 100	5.7	2.8	4.1	1.3 versicolor
## 101	6.3	3.3	6.0	2.5 virginica
## 102	5.8	2.7	5.1	1.9 virginica
## 103	7.1	3.0	5.9	2.1 virginica
## 104	6.3	2.9	5.6	1.8 virginica
## 105	6.5	3.0	5.8	2.2 virginica
## 106	7.6	3.0	6.6	2.1 virginica
## 107	4.9	2.5	4.5	1.7 virginica
## 108	7.3	2.9	6.3	1.8 virginica
## 109	6.7	2.5	5.8	1.8 virginica
## 110	7.2	3.6	6.1	2.5 virginica
## 111	6.5	3.2	5.1	2.0 virginica
## 112	6.4	2.7	5.3	1.9 virginica
## 113	6.8	3.0	5.5	2.1 virginica
## 114	5.7	2.5	5.0	2.0 virginica
## 115	5.8	2.8	5.1	2.4 virginica
## 116	6.4	3.2	5.3	2.3 virginica
## 117	6.5	3.0	5.5	1.8 virginica
## 118	7.7	3.8	6.7	2.2 virginica
## 119	7.7	2.6	6.9	2.3 virginica
## 120	6.0	2.2	5.0	1.5 virginica
## 121	6.9	3.2	5.7	2.3 virginica
## 122	5.6	2.8	4.9	2.0 virginica
## 123	7.7	2.8	6.7	2.0 virginica
## 124	6.3	2.7	4.9	1.8 virginica
## 125	6.7	3.3	5.7	2.1 virginica
## 126	7.2	3.2	6.0	1.8 virginica

```
## 127      6.2      2.8      4.8      1.8 virginica
## 128      6.1      3.0      4.9      1.8 virginica
## 129      6.4      2.8      5.6      2.1 virginica
## 130      7.2      3.0      5.8      1.6 virginica
## 131      7.4      2.8      6.1      1.9 virginica
## 132      7.9      3.8      6.4      2.0 virginica
## 133      6.4      2.8      5.6      2.2 virginica
## 134      6.3      2.8      5.1      1.5 virginica
## 135      6.1      2.6      5.6      1.4 virginica
## 136      7.7      3.0      6.1      2.3 virginica
## 137      6.3      3.4      5.6      2.4 virginica
## 138      6.4      3.1      5.5      1.8 virginica
## 139      6.0      3.0      4.8      1.8 virginica
## 140      6.9      3.1      5.4      2.1 virginica
## 141      6.7      3.1      5.6      2.4 virginica
## 142      6.9      3.1      5.1      2.3 virginica
## 143      5.8      2.7      5.1      1.9 virginica
## 144      6.8      3.2      5.9      2.3 virginica
## 145      6.7      3.3      5.7      2.5 virginica
## 146      6.7      3.0      5.2      2.3 virginica
## 147      6.3      2.5      5.0      1.9 virginica
## 148      6.5      3.0      5.2      2.0 virginica
## 149      6.2      3.4      5.4      2.3 virginica
## 150      5.9      3.0      5.1      1.8 virginica
```

```
iris %>% group_by(Species)
```

```
## # A tibble: 150 x 5
## # Groups:   Species [3]
##   Sepal.Length Sepal.Width Petal.Length Petal.Width Species
##         <dbl>         <dbl>         <dbl>         <dbl> <fctr>
## 1         5.1         3.5         1.4         0.2 setosa
## 2         4.9         3.0         1.4         0.2 setosa
## 3         4.7         3.2         1.3         0.2 setosa
## 4         4.6         3.1         1.5         0.2 setosa
## 5         5.0         3.6         1.4         0.2 setosa
## 6         5.4         3.9         1.7         0.4 setosa
## 7         4.6         3.4         1.4         0.3 setosa
## 8         5.0         3.4         1.5         0.2 setosa
## 9         4.4         2.9         1.4         0.2 setosa
## 10        4.9         3.1         1.5         0.1 setosa
## # ... with 140 more rows
```

using slice

```
iris%>%slice(1:5)
```

```
## # A tibble: 5 x 5
##   Sepal.Length Sepal.Width Petal.Length Petal.Width Species
##         <dbl>         <dbl>         <dbl>         <dbl> <fctr>
## 1         5.1         3.5         1.4         0.2 setosa
## 2         4.9         3.0         1.4         0.2 setosa
## 3         4.7         3.2         1.3         0.2 setosa
```



```
## 4          4.6          3.1          1.5          0.2 setosa
## 5          5.0          3.6          1.4          0.2 setosa
```

using summarize

```
iris%>%group_by(Species)%>%summarize(n_obs=n(),min_length=min(Sepal.Length),max_length=max(Sepal.Length))
```

```
## # A tibble: 3 x 6
##   Species n_obs min_length max_length min_petal_width max_petal_width
##   <fctr> <int>      <dbl>      <dbl>          <dbl>          <dbl>
## 1 setosa   50       4.3       5.8           0.1           0.6
## 2 versicolor 50       4.9       7.0           1.0           1.8
## 3 virginica 50       4.9       7.9           1.4           2.5
```

using select

```
iris%>%group_by(Species)%>%summarize(n_obs=n(),min_length=min(Sepal.Length),max_length=max(Sepal.Length))
```

```
## # A tibble: 3 x 2
##   Species n_obs
##   <fctr> <int>
## 1 setosa   50
## 2 versicolor 50
## 3 virginica 50
```

There are many other useful methods you can call from the packages for example:

```
rename,
mutate
filter,
left_join,
right_join,
inner_join,
full_join .....
```