

Read file -> Tidy up the file -> Transform -> visualize

Installing

#install the tidyverse package
install.packages("tidyverse")

Do this once

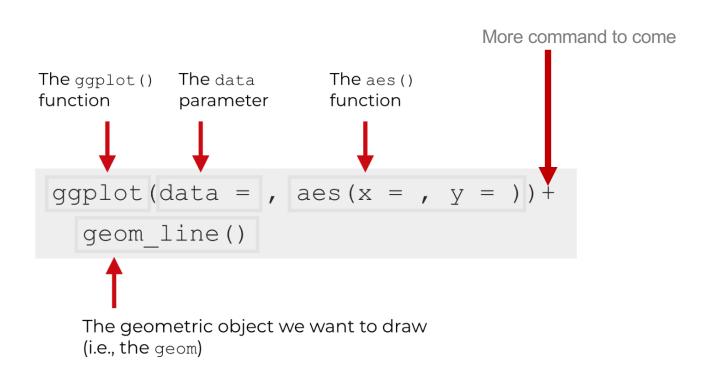
Loading

load the tidyverse package
library(tidyverse)

Do at the top of every* script

```
surveys <- read_csv("data/portal_data_joined.csv")</pre>
```

Notice the underscore!





- Select and filter
- Mutate
- Summarise
- Group by

Subset Observations (Rows)



dplyr::filter(iris, Sepal.Length > 7)

Extract rows that meet logical criteria.

dplyr::distinct(iris)

Remove duplicate rows.

dplyr::sample_frac(iris, 0.5, replace = TRUE)

Randomly select fraction of rows.

dplyr::sample_n(iris, 10, replace = TRUE)

Randomly select n rows.

dplyr::slice(iris, 10:15)

Select rows by position.

dplyr::top_n(storms, 2, date)

Select and order top n entries (by group if grouped data).

	Logic in R - ?(Comparison, ?base	::Logic
<	Less than	!=	Not equal to
>	Greater than	%in%	Group membership
==	Equal to	is.na	Is NA
<=	Less than or equal to	!is.na	Is not NA
>=	Greater than or equal to	&, ,!,xor,any,all	Boolean operators

Subset Variables (Columns)



dplyr::select(iris, Sepal.Width, Petal.Length, Species)

Select columns by name or helper function.



Base R - dataFrame[ROW , COLUMN]



Use pipe for passing data to a new command

Often confused with the **ggplot +** which is used to add more lines to a plot

- Installing tidyverse once load every time
- Importing tables with dplyr
- Basic ggplot syntax
- select and filter
- Pipes. %>%

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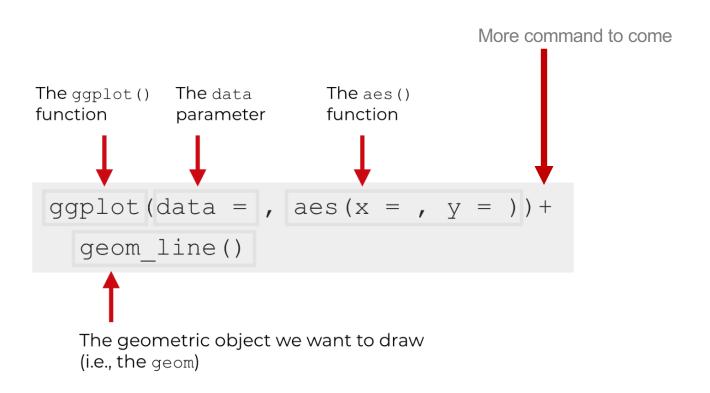
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Base R - dataFrame[ROW , COLUMN]



Use **pipe** for passing **data** to a new command

Often confused with the **ggplot +** which is used to add more lines to a plot

Transforming tables with dplyr

- creating new columns
- •
- sorting data tables
- summarizing data tables
- Frequencies
- Grouping data

More ggplot

- Plotting data (lines plots)
- Faceting
- Customising plots
- Exporting tables to file
- Boxplots and histograms



dplyr::mutate(iris, sepal = Sepal.Length + Sepal. Width)

Compute and append one or more new columns.

dplyr::mutate_each(iris, funs(min_rank))

Apply window function to each column.

dplyr::transmute(iris, sepal = Sepal.Length + Sepal. Width)

Compute one or more new columns. Drop original columns.



Mutate uses **window functions**, functions that take a vector of values and return another vector of values, such as:

Possible functions include:

dplyr::lead

Copy with values shifted by 1.

dplyr::lag

Copy with values lagged by 1.

dplyr::dense_rank

Ranks with no gaps.

dplyr::min_rank

Ranks. Ties get min rank.

dplyr::percent_rank

Ranks rescaled to [0, 1].

dplyr::row_number

Ranks. Ties got to first value.

dplyr::ntile

Bin vector into n buckets.

dplyr::between

Are values between a and b?

dplyr::cume_dist

Cumulative distribution.

dplyr::arrange(mtcars, desc(mpg))

Order rows by values of a column (high to low).

Summarise Data



dplyr::summarise(iris, avg = mean(Sepal.Length))

Summarise data into single row of values.

dplyr::summarise_each(iris, funs(mean))

Apply summary function to each column.

dplyr::count(iris, Species, wt = Sepal.Length)

Count number of rows with each unique value of variable (with or without weights).



Summarise uses **summary functions**, functions that take a vector of values and return a single value, such as:

Possible functions include:

dplyr::first

First value of a vector.

dplyr::last

Last value of a vector.

dplyr::nth

Nth value of a vector.

dplyr::n

of values in a vector.

dplyr::n_distinct

of distinct values in a vector.

IQR

IQR of a vector.

min

Minimum value in a vector.

max

Maximum value in a vector.

mean

Mean value of a vector.

median

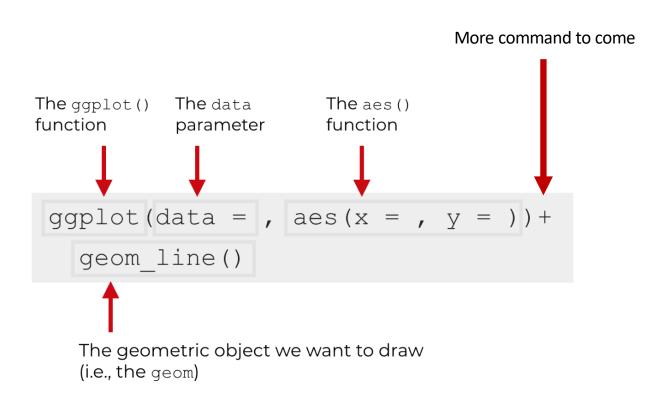
Median value of a vector.

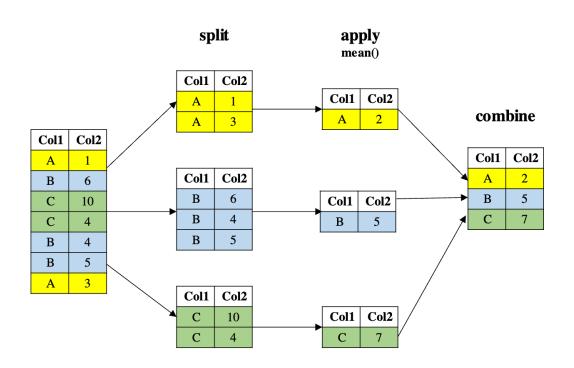
var

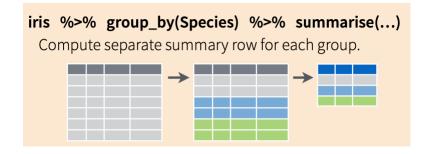
Variance of a vector.

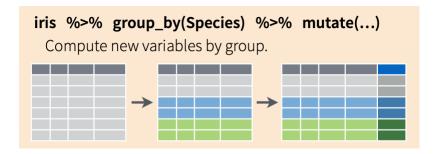
sd

Standard deviation of a vector.



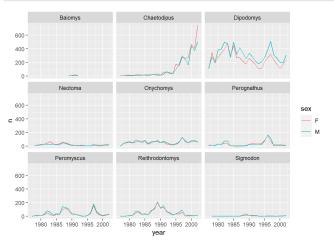






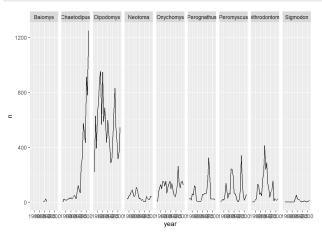
facet_wrap





facet_grid

```
ggplot(data = yearly_counts, mapping = aes(x = year, y = n)) +
   geom_line() +
   #display the genera as columns
   facet_grid(cols = vars(genus))
```



Major labels

```
labs(title = "Observed genera through time",
    x = "Year of observation",
    y = "Number of animals")
```

Colour and legend

Everything else

```
theme(axis.text.x = element_text(size=7, angle=90, vjust=0.5),
    axis.text.y = element_text(size=7),
    strip.text=element_text(size=7, angle=45)) +
```