

R For Data Science Cheat Sheet

Tidyverse for Beginners

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Tidyverse

The **tidyverse** is a powerful collection of R packages that are actually data tools for transforming and visualizing data. All packages of the tidyverse share an underlying philosophy and common APIs.

The core packages are:



- **ggplot2**, which implements the grammar of graphics. You can use it to visualize your data.



- **dplyr** is a grammar of data manipulation. You can use it to solve the most common data manipulation challenges.



- **tidyr** helps you to create tidy data or data where each variable is in a column, each observation is a row and each value is a cell.



- **readr** is a fast and friendly way to read rectangular data.



- **purrr** enhances R's functional programming (FP) toolkit by providing a complete and consistent set of tools for working with functions and vectors.



- **tibble** is a modern re-imaging of the data frame.



- **stringr** provides a cohesive set of functions designed to make working with strings as easy as possible



- **forcats** provide a suite of useful tools that solve common problems with factors.

You can install the complete tidyverse with:

```
> install.packages("tidyverse")
```

Then, load the core tidyverse and make it available in your current R session by running:

```
> library(tidyverse)
```

Note: there are many other tidyverse packages with more specialised usage. They are not loaded automatically with `library(tidyverse)`, so you'll need to load each one with its own call to `library()`.

Useful Functions

```
> tidyverse_conflicts()
```

Conflicts between tidyverse and other packages

```
> tidyverse_deps()
```

List all tidyverse dependencies

```
> tidyverse_logo()
```

Get tidyverse logo, using ASCII or unicode characters

```
> tidyverse_packages()
```

List all tidyverse packages

```
> tidyverse_update()
```

Update tidyverse packages

Loading in the data

```
> library(datasets)
```

Load the datasets package

```
> library(gapminder)
```

Load the gapminder package

```
> attach(iris)
```

Attach iris data to the R search path

dplyr

Filter

`filter()` allows you to select a subset of rows in a data frame.

```
> iris %>%  
  filter(Species=="virginica")  
> iris %>%  
  filter(Species=="virginica",  
         Sepal.Length > 6)
```

Select iris data of species "virginica"
Select iris data of species "virginica" and sepal length greater than 6.

Arrange

`arrange()` sorts the observations in a dataset in ascending or descending order based on one of its variables.

```
> iris %>%  
  arrange(Sepal.Length)  
> iris %>%  
  arrange(desc(Sepal.Length))
```

Sort in ascending order of sepal length
Sort in descending order of sepal length

Combine multiple dplyr verbs in a row with the pipe operator `%>%`:

```
> iris %>%  
  filter(Species=="virginica") %>%  
  arrange(desc(Sepal.Length))
```

Filter for species "virginica" then arrange in descending order of sepal length

Mutate

`mutate()` allows you to update or create new columns of a data frame.

```
> iris %>%  
  mutate(Sepal.Length=Sepal.Length*10)  
> iris %>%  
  mutate(SLMm=Sepal.Length*10)
```

Change Sepal.Length to be in millimeters
Create a new column called SLMm

Combine the verbs `filter()`, `arrange()`, and `mutate()`:

```
> iris %>%  
  filter(Species=="Virginica") %>%  
  mutate(SLMm=Sepal.Length*10) %>%  
  arrange(desc(SLMm))
```

Summarize

`summarize()` allows you to turn many observations into a single data point.

```
> iris %>%  
  summarize(medianSL=median(Sepal.Length))  
> iris %>%  
  filter(Species=="virginica") %>%  
  summarize(medianSL=median(Sepal.Length))
```

Summarize to find the median sepal length
Filter for virginica then summarize the median sepal length

You can also summarize multiple variables at once:

```
> iris %>%  
  filter(Species=="virginica") %>%  
  summarize(medianSL=median(Sepal.Length),  
            maxSL=max(Sepal.Length))
```

`group_by()` allows you to summarize within groups instead of summarizing the entire dataset:

```
> iris %>%  
  group_by(Species) %>%  
  summarize(medianSL=median(Sepal.Length),  
            maxSL=max(Sepal.Length))  
> iris %>%  
  filter(Sepal.Length>6) %>%  
  group_by(Species) %>%  
  summarize(medianPL=median(Petal.Length),  
            maxPL=max(Petal.Length))
```

Find median and max sepal length of each species
Find median and max petal length of each species with sepal length > 6

ggplot2

Scatter plot

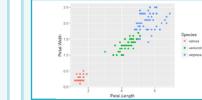
Scatter plots allow you to compare two variables within your data. To do this with ggplot2, you use `geom_point()`

```
> iris_small <- iris %>%  
  filter(Sepal.Length > 5)  
> ggplot(iris_small, aes(x=Petal.Length,  
                         y=Petal.Width)) +  
  geom_point()
```

Compare petal width and length

Additional Aesthetics

• Color



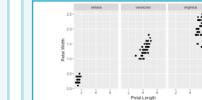
```
> ggplot(iris_small, aes(x=Petal.Length,  
                         y=Petal.Width,  
                         color=Species)) +  
  geom_point()
```

• Size



```
> ggplot(iris_small, aes(x=Petal.Length,  
                         y=Petal.Width,  
                         color=Species,  
                         size=Sepal.Length)) +  
  geom_point()
```

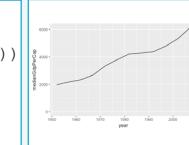
Faceting



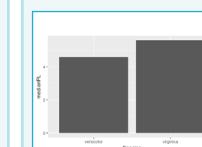
```
> ggplot(iris_small, aes(x=Petal.Length,  
                         y=Petal.Width)) +  
  geom_point() +  
  facet_wrap(~Species)
```

Line Plots

```
> by_year <- gapminder %>%  
  group_by(year) %>%  
  summarize(medianGdpPerCap=median(gdpPerCap))  
> ggplot(by_year, aes(x=year,  
                         y=medianGdpPerCap)) +  
  geom_line() +  
  expand_limits(y=0)
```



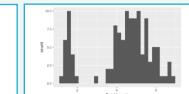
Bar Plots



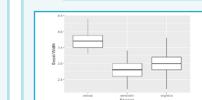
```
> by_species <- iris %>%  
  filter(Sepal.Length>6) %>%  
  group_by(Species) %>%  
  summarize(medianPL=median(Petal.Length))  
> ggplot(by_species, aes(x=Species,  
                         y=medianPL)) +  
  geom_col()
```

Histograms

```
> ggplot(iris_small, aes(x=Petal.Length)) +  
  geom_histogram()
```



Box Plots



```
> ggplot(iris_small, aes(x=Species,  
                         y=Sepal.Width)) +  
  geom_boxplot()
```

