

Summary

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Resource: cheat Sheets

This page features a large number of cheat sheets on different topics: Cheat Sheets

R Basics

The **if** statement: example

```
my_number <-12

if (my_number < 20){
  x <- sprintf('%i is less than 20', my_number)
  print(x)
}
```

```
## [1] "12 is less than 20"
```

The for loop: example

```
my_vector <- runif(5)
for (x in my_vector) {
  y <- x * 3
  print(y)
}
```

```
## [1] 0.623995
## [1] 2.54174
## [1] 1.017559
## [1] 0.6667724
## [1] 0.5250116
```

One very common use of the `for` loop is to iterate a bit of code exactly `n` times.

```
number_of_time_i_want_to_repeat_this <- 10
for (x in 1:10) {
  print('This is being repeated!')
}
```

```
## [1] "This is being repeated!"
## [1] "This is being repeated!"
## [1] "This is being repeated!"
## [1] "This is being repeated!"
## [1] "This is being repeated!"
## [1] "This is being repeated!"
## [1] "This is being repeated!"
## [1] "This is being repeated!"
## [1] "This is being repeated!"
## [1] "This is being repeated!"
```

The while loop: example

```
i <- 1
while (i < 6) {
  print(i)
  i <- i + 1
}
```

```
## [1] 1
## [1] 2
## [1] 3
## [1] 4
## [1] 5
```

Reading data

Reading data from excel using `readxl`

```
library(tidyverse)
library(readxl)
# Reading the first sheet
data <- read_excel("data/transit-data.xlsx")
# Reading a specific range from a specific sheet
data <- read_excel("data/transit-data.xlsx", sheet = 'info', range = 'B1:C7')
```

Reading data from text files

Reading comma separated files

```
data <- read_csv('data/pakistan_intellectual_capital.csv')

## New names:
## Rows: 1142 Columns: 13
## -- Column specification
## ----- Delimiter: "," chr
## (10): Teacher Name, University Currently Teaching, Department, Province ... dbl
## (3): ...1, S#, Year
## i Use `spec()` to retrieve the full column specification for this data. i
## Specify the column types or set `show_col_types = FALSE` to quiet this message.
## * `` -> `...1`
```

Reading tab separated files

```
data <- read_tsv('data/films.dat')

## Rows: 100 Columns: 6
## -- Column specification -----
## Delimiter: "\t"
## chr (1): Title
## dbl (5): Year, Length, Cast, Rating, Description
##
## i Use `spec()` to retrieve the full column specification for this data.
## i Specify the column types or set `show_col_types = FALSE` to quiet this message.
```

Reading files separated by specific character

```
# The columns of this file are separated by a space.
data <- read_delim('data/wages1833bis.csv', delim = ' ')

## New names:
## Rows: 51 Columns: 6
## -- Column specification
## ----- Delimiter: " " dbl
## (6): ...1, age, mnum, mwage, fnum, fwage
## i Use `spec()` to retrieve the full column specification for this data. i
## Specify the column types or set `show_col_types = FALSE` to quiet this message.
## * `` -> `...1`
```

Some interesting options when using read_csv(), read_tsv(), or read_delim()

No column names?

If data has no column names, use col_names = FALSE

```
read_csv("1,2,3\n4,5,6", col_names = FALSE)
```

```
## Rows: 2 Columns: 3
## -- Column specification -----
## Delimiter: ","
## dbl (3): X1, X2, X3
##
```

```
## i Use `spec()` to retrieve the full column specification for this data.
## i Specify the column types or set `show_col_types = FALSE` to quiet this message.

## # A tibble: 2 x 3
##       X1     X2     X3
##   <dbl> <dbl> <dbl>
## 1     1     2     3
## 2     4     5     6
```

You can also directly set the column names in this case.

```
read_csv("1,2,3\n4,5,6", col_names = c("x", "y", "z"))
```

```
## Rows: 2 Columns: 3
## -- Column specification -----
## Delimiter: ","
## dbl (3): x, y, z
##
## i Use `spec()` to retrieve the full column specification for this data.
## i Specify the column types or set `show_col_types = FALSE` to quiet this message.

## # A tibble: 2 x 3
##       x     y     z
##   <dbl> <dbl> <dbl>
## 1     1     2     3
## 2     4     5     6
```

Specifying missing data

Data cleaning and data operations

Another option that commonly needs tweaking is `na`: this specifies the value (or values) that are used to represent missing values in your file:

```
read_csv("a,b,c\n1,2,.", na = ".")
```

```
## Rows: 1 Columns: 3
## -- Column specification -----
## Delimiter: ","
## dbl (2): a, b
## lgl (1): c
##
## i Use `spec()` to retrieve the full column specification for this data.
## i Specify the column types or set `show_col_types = FALSE` to quiet this message.

## # A tibble: 1 x 3
##       a     b c
##   <dbl> <dbl> <lgl>
## 1     1     2 NA
```

Renaming variables to R acceptable format

Human readable names are very handy for coding. **Indeed, in general, only dots and underscores are allowed in variable names.** There is a quick way to solve this.

```
data <- read_excel("data/transit-data.xlsx", sheet='transport data', skip = 1)
colnames(data)
```

```
## [1] "sender location" "sender latitude" "sender longitude"
```

```
## [4] "receiver location" "receiver latitude" "receiver longitude"
## [7] "date" "number of items"
```

```
colnames(data) <- make.names(colnames(data))
colnames(data)
```

```
## [1] "sender.location" "sender.latitude" "sender.longitude"
## [4] "receiver.location" "receiver.latitude" "receiver.longitude"
## [7] "date" "number.of.items"
```

Creating new variables using mutate()

```
data <- read_csv('data/pizzasize.csv')
```

```
## Rows: 250 Columns: 5
## -- Column specification -----
## Delimiter: ","
## chr (3): Store, CrustDescription, Topping
## dbl (2): ID, Diameter
##
## i Use `spec()` to retrieve the full column specification for this data.
## i Specify the column types or set `show_col_types = FALSE` to quiet this message.
```

```
data <- mutate(data, surface = 3.14 * (Diameter/2)^2)
```

Selecting columns using select()

```
data <- read_excel("data/transit-data.xlsx", sheet = 'transport data', skip=1)
colnames(data) <- make.names(colnames(data))
subset <- select(data, date, sender.latitude)
head(subset)
```

```
## # A tibble: 6 x 2
##   date sender.latitude
##   <chr>          <dbl>
## 1 5729           51.0
## 2 5741           51.0
## 3 5743           51.0
## 4 5752           51.0
## 5 5757           51.0
## 6 5765           51.0
```

Filtering using filter()

```
subset <- filter(data, sender.latitude < 50)
subset <- filter(subset, sender.latitude > 32)
subset <- filter(subset, sender.longitude > 0)
subset <- select(subset, sender.latitude, sender.longitude)
summary(subset)
```

```
## sender.latitude sender.longitude
## Min. :47.54 Min. : 2.130
## 1st Qu.:47.54 1st Qu.: 9.212
## Median :48.12 Median : 9.300
## Mean :48.24 Mean : 9.020
```

```
## 3rd Qu.:48.80    3rd Qu.:10.739
## Max.      :49.57    Max.      :16.320
```

Splitting and uniting variable values using `separate()` and `unite()`

```
patient_data <- read_tsv('data/inpatient.tsv')

## Rows: 163065 Columns: 12
## -- Column specification -----
## Delimiter: "\t"
## chr (10): DRG Definition, Provider Name, Provider Street Address, Provider C...
## dbl (2): Provider Id, Total Discharges
##
## i Use `spec()` to retrieve the full column specification for this data.
## i Specify the column types or set `show_col_types = FALSE` to quiet this message.

colnames(patient_data) <- make.names(colnames(patient_data))
# Split a column into two columns, based on a character
test1 <- separate(patient_data, Hospital.Referral.Region.Description, into=c('state', 'city'), sep='-')

## Warning: Expected 2 pieces. Additional pieces discarded in 791 rows [710, 711,
## 718, 830, 853, 1918, 1921, 1927, 1930, 1940, 2055, 2086, 3398, 3401, 3411, 3415,
## 3594, 3630, 5424, 5425, ...].

# Join two columns
test2 <- unite(patient_data, 'combined', c(Provider.Id, Provider.Name), sep='_+', remove = FALSE)
```

String manipulation using the `stringr` library

It does happen that you need to clean textual data. The `stringr` package has a bunch of functions to make your life easier (but not easy). I will run through some examples but do have a look at the cheatsheet as well.

Grouping and summarizing data using `group_by()` and `summarize()`

The `group_by()` function takes a tibble and returns the same tibble, but with some extra information so that any subsequent function can act on each unique combination defined in the `group_by()`.

```
car_data <- read_delim('data/cars.txt', delim = ' ')

## Rows: 93 Columns: 26
## -- Column specification -----
## Delimiter: " "
## chr (6): make, model, type, cylinders, rearseat, luggage
## dbl (20): min_price, mid_price, max_price, mpg_city, mpg_hgw, airbag, drive,...
##
## i Use `spec()` to retrieve the full column specification for this data.
## i Specify the column types or set `show_col_types = FALSE` to quiet this message.

grouped <- group_by(car_data, type, make)
summaries <- summarise(grouped, mean.length = mean(length))

## `summarise()` has grouped output by 'type'. You can override using the
## `.groups` argument.

summaries

## # A tibble: 81 x 3
## # Groups:   type [6]
```

```
##   type    make      mean.length
##   <chr>   <chr>      <dbl>
## 1 Compact Audi      180
## 2 Compact Chevrolet 183
## 3 Compact Chrysler  183
## 4 Compact Dodge     181
## 5 Compact Ford      177
## 6 Compact Honda     185
## 7 Compact Mazda     184
## 8 Compact Mercedes-Benz 175
## 9 Compact Nissan    181
## 10 Compact Oldsmobile 188
## # ... with 71 more rows
## # i Use `print(n = ...)` to see more rows
```

You can ask for more than one summary statistic.

```
summaries <- summarise(grouped, mean.length = mean(length), max.length = max(length), std_rpm = sd(rpm))
```

```
## `summarise()` has grouped output by 'type'. You can override using the
## `.groups` argument.
```

```
summaries
```

```
## # A tibble: 81 x 5
## # Groups:   type [6]
##   type    make      mean.length max.length std_rpm
##   <chr>   <chr>      <dbl>      <dbl>   <dbl>
## 1 Compact Audi      180        180     NA
## 2 Compact Chevrolet 183        184     0
## 3 Compact Chrysler  183        183     NA
## 4 Compact Dodge     181        181     NA
## 5 Compact Ford      177        177     NA
## 6 Compact Honda     185        185     NA
## 7 Compact Mazda     184        184     NA
## 8 Compact Mercedes-Benz 175        175     NA
## 9 Compact Nissan    181        181     NA
## 10 Compact Oldsmobile 188        188     NA
## # ... with 71 more rows
## # i Use `print(n = ...)` to see more rows
```

Converting to wide format using `pivot_wider()`

The result can be reshaped into a wide format. While this format is often not suited for plotting or analysis, it might make it easier to look at the data. Here is a quick visual:

country	year	cases
Angola	1999	800
Angola	2000	750
Angola	2001	925
Angola	2002	1020
India	1999	20100
India	2000	25650
India	2001	26800
India	2002	27255
Mongolia	1999	450
Mongolia	2000	512
Mongolia	2001	510
Mongolia	2002	586

country	1999	2000	2001	2002
Angola	800	750	925	1020
India	20100	25650	26800	27255
Mongolia	450	512	510	586

Pivot data wider

```
data %>%
  pivot_wider(
    names_from = "year",
    values_from = "cases"
  )
```

```
car_data <- read_delim('data/cars.txt', delim = ' ')
```

```
## Rows: 93 Columns: 26
## -- Column specification -----
## Delimiter: " "
## chr (6): make, model, type, cylinders, rearseat, luggage
## dbl (20): min_price, mid_price, max_price, mpg_city, mpg_hgw, airbag, drive,...
##
## i Use `spec()` to retrieve the full column specification for this data.
## i Specify the column types or set `show_col_types = FALSE` to quiet this message.
```

```
grouped <- group_by(car_data, type, make)
summaries <- summarise(grouped, mean.length = mean(length))
```

```
## `summarise()` has grouped output by 'type'. You can override using the
## `.groups` argument.
```

```
wide <- pivot_wider(summaries, id_cols = make, names_from = type, values_from = mean.length)
head(wide)
```

```
## # A tibble: 6 x 7
##   make      Compact Large Midsize Small Sporty  Van
##   <chr>      <dbl> <dbl>   <dbl> <dbl>   <dbl> <dbl>
## 1 Audi         180    NA     193    NA     NA     NA
## 2 Chevrolet    183   214     198    NA    186    186
## 3 Chrysler     183   203      NA    NA     NA     NA
## 4 Dodge        181    NA     192   173    180    175
## 5 Ford         177   212     192   156   180.    176
## 6 Honda        185    NA      NA    173   175     NA
```

Making data longer (melting data) using `pivot_longer()`

Here is a quick graphic:

country	1999	2000	2001	2002
Angola	800	750	925	1020
India	20100	25650	26800	27255
Mongolia	450	512	510	586

country	year	cases
Angola	1999	800
Angola	2000	750
Angola	2001	925
Angola	2002	1020
India	1999	20100
India	2000	25650
India	2001	26800
India	2002	27255
Mongolia	1999	450
Mongolia	2000	512
Mongolia	2001	510
Mongolia	2002	586

Pivot data longer

```
data %>%
  pivot_longer(
    cols = 1999:2002,
    names_to = "year",
    values_to = "cases"
  )
```

Let's look at some data:

```
head(relig_income, 5)
```

```
## # A tibble: 5 x 11
##   religion      `<$10k` $10-2~1 $20-3~2 $30-4~3 $40-5~4 $50-7~5 $75-1~6 $100--7
##   <chr>         <dbl>   <dbl>   <dbl>   <dbl>   <dbl>   <dbl>   <dbl>   <dbl>
## 1 Agnostic         27     34     60     81     76    137    122    109
## 2 Atheist          12     27     37     52     35     70     73     59
## 3 Buddhist         27     21     30     34     33     58     62     39
## 4 Catholic        418    617    732    670    638   1116    949    792
## 5 Don't know/re~    15     14     15     11     10     35     21     17
## # ... with 2 more variables: `>150k` <dbl>, `Don't know/refused` <dbl>, and
## #   abbreviated variable names 1: ` $10-20k`, 2: ` $20-30k`, 3: ` $30-40k`,
## #   4: ` $40-50k`, 5: ` $50-75k`, 6: ` $75-100k`, 7: ` $100-150k`
## # i Use `colnames()` to see all variable names
```

This data is in a wider format. But we can easily melt it to a long format.

```
new <- pivot_longer(relig_income, cols = !religion)
head(new, 5)
```

```
## # A tibble: 5 x 3
##   religion name      value
##   <chr>   <chr>   <dbl>
## 1 Agnostic <$10k      27
## 2 Agnostic $10-20k    34
## 3 Agnostic $20-30k    60
## 4 Agnostic $30-40k    81
## 5 Agnostic $40-50k    76
```

You can specify names for the new columns while melting.

```
new <- pivot_longer(relig_income, !religion, names_to = "income", values_to = "count")
head(new, 5)
```

```
## # A tibble: 5 x 3
##   religion income  count
##   <chr>   <chr>   <dbl>
```

```
## 1 Agnostic <$10k      27
## 2 Agnostic $10-20k    34
## 3 Agnostic $20-30k    60
## 4 Agnostic $30-40k    81
## 5 Agnostic $40-50k    76
```

Merging data using the `_join()` functions

The different merge operations are illustrated in the image below. The various operations differ in the way they handle rows missing in the left or right tibble. In the image below, the merge is done by the variable ID.

ID X1			ID X2		
1	a1		2	b1	
2	a2		3	b2	

inner_join			left_join			right_join			full_join			semi_join		anti_join	
ID	X1	X2	ID	X1	X2	ID	X1	X2	ID	X1	X2	ID	X1	ID	X1
2	a2	b1	1	a1	NA	2	a2	b1	1	a1	NA	2	a2	1	a1
			2	a2	b1	3	NA	b2	2	a2	b1				
									3	NA	b2				