

Contents

1	Introduction 7					
	1.1	References & Resources	7			
2	Import or Create Data					
	2.1	Create Data	9			
	2.2	Local Drive	0			
	2.3	Internet	1			
	2.4	Database	3			
	2.5	Notes & References	5			
3	Tidy		7			
	3.1	Explore Raw Data	7			
	3.2	Tidy Data	8			
	3.3	Prepare Data for Analysis	0			
4	Tra	nsform 2	5			
	4.1	Arithmetic & Summary Statistics	5			
	4.2	Create New Variables or Modify Existing Ones	8			
	4.3	Dates and Datetimes	8			
	4.4	Factors	1			
	4.5	Merge or Append Data	1			
	4.6	Narrow in on Observations of Interest	2			
	47	Test 3	3			

4	CONTENTS	

5	Visualize				
	5.1	Interfaces	35		
	5.2	Graphs, Charts, and Plots	35		
	5.3	Graphs, Charts, and Plots - Statistics	40		
6	Mod	lel	41		
	6.1	Cluster Analysis	41		
	6.2	General	41		
	6.3	Regression	42		
	6.4	Simulation & Prediction	43		
	6.5	Strings	43		
7	Con	nmunicate	45		
	7.1	CSS	45		
	7.2	Export	45		
	7.3	Format Output	46		
	7.4	Graphics	47		
	7.5	Plots	47		
	7.6	RMarkdown	47		
8	Program 49				
	8.1	Characters	49		
	8.2	Conditionals & Control Flows	49		
	8.3	Environment and Workspace	50		
	8.4	Evaluation (Standard and Non-standard) $\ \ \ldots \ \ \ldots \ \ \ldots$	53		
	8.5	Functionals	54		
	8.6	Functions	55		
	8.7	Learn About an Object	56		
	8.8	Loops	57		
	8.9	Optimization	57		
	8.10	Pipes	57		
	8.11	Popups	57		

CONTENTS	5
8.12 Selecting & Subsetting	58
8.13 Style Guide	59
8.14 System Commands	59
8.15 Version Control	59

6 CONTENTS

Introduction

In *R for Data Science*, Garrett Grolemund and Hadley Wickham outline the tools needed to tackle about 80% of the tasks required in a typical data science project ("Introduction"). Those tools, illustrated in the diagram below and the organizing template for this reference manual, are:

- Import
- Tidy
- Transform
- Visualize
- Model
- Communicate
- \bullet Program

PhantomJS not found. You can install it with webshot::install_phantomjs(). If it is installed,

1.1 References & Resources

- For an introduction to the R programming language, see the R Project for Statistical Computing's "What is R?" and Wikipedia's "R (programming language)."
- To download R, go to r-project.org and choose the cloud CRAN Mirror option.
- To program in the R language on a user-friendly platform, download the RStudio IDE.
- The R Project for Statistical Computing

- Library of R Packages
- Getting Help with R
- The R Manuals
- Frequently Asked Questions
- Books Related to R
- Documentation

• RStudio

- RStudio Cheat Sheets
- Webinars and Videos On Demand
- Online learning
- RStudio Blog

• Online Manuals

- R for Data Science
- Advanced R by Hadley Wickham + Hadley's second edition draft is available here.
- R Packages by Hadley Wickham
- The R Inferno by Patrick Burns
- The tidyverse style guide
- Efficient R Programming
- Free R Reading Material
- $-\ What\ They\ Forgot\ to\ Teach\ You\ About\ R$ by Jenny Bryan and Jim Hester

• Other Online Resources

- DataCamp
- RDocumentation
- R Bloggers
 - \ast "Tutorials for learning R"
- Regular Expressions 101

Import or Create Data

2.1 Create Data

```
{base}
  • array()
  • c()
       - See base::vector().
  • data.frame()
  • dir.create()
  • factor()
  • list()
  • matrix()
  • seq()
  • vector()
       - Preferable to base::c() when creating an empty vector ("Five Tips
         to Improve Your R Code").
{stats}
  • rnorm()
{tibble}
  • add_row()
  • tibble()
  • tribble()
```

2.2 Local Drive

```
{base}
  • attach()
       - Allows objects in the database to be accessed by giving their names
         (e.g., height rather than women$height).
  • file.choose()
  • file.size()
   • load()
       - Reload datasets saved with base::save().
  • readRDS()
       - Restore an R object written with base::saveRDS().
{data.table}
  • fread()
       - Similar to utils::read.table(), but faster and more convenient for
         large data sets.
{foreign}
  • read.spss()
{haven}
  • read_sas()
{readr}
  • read_csv()
  • read_csv2()
  • read_delim()
  • rread_tsv()
{readxl}
  • excel_sheets()
  • read_excel()
   • read_xls()
```

2.3. INTERNET 11

• read_xlsx()

{utils}

- data()
 - Load specified data sets, or list the available data sets.
 - Use this function to load the data sets that accompany R packages, such as openintro::hsb2, openintro::email50,and gapminder::gapminder.
- read.csv()
- read.csv2()
- read.delim()
- read.delim2()
- read.table()

{XLConnect}

• readWorksheetFromFile()

2.3 Internet

{httr}

- GET()
 - Get a URL.

{jsonlite}

• read_json()

{readr}

- read_csv()
- read_csv2()
- read_delim()
- read_tsv()

{rjson}

• fromJSON()

destfile = "chis_2009.Rdata",

```
- Convert JSON to R.
{utils}
   • download.file()
       - See example below.
   • unzip()
       - See example below.
2.3.1 Examples
utils::download.file():
download.file(
  "https://assets.datacamp.com/production/repositories/5028/datasets/a55843f83746968c7
  destfile = paste0(getwd(), "/Snake River Visits.rds")
)
# Option 1:
snake_river_visits <- readRDS(file.choose())</pre>
# Option 2:
                   <- pasteO(getwd(), "/Snake River Visits.rds")</pre>
snake_river_path
snake_river_visits <- readRDS(snake_river_path)</pre>
utils::download.file() for .Rdata files:
# Example 1:
download.file(
  "https://assets.datacamp.com/production/repositories/236/datasets/7f714f993f1ad4c3d2
  destfile = "datacamp_iris_dataset.Rdata"
load("datacamp_iris_dataset.Rdata")
# Example 2:
download.file(
  "https://assets.datacamp.com/production/repositories/235/datasets/3b6fc2923b59905858
```

The documentation for `download.file` indicates that the function will
automatically include `mode = "wb"` for .Rdata files. That may have happened
in Example 1, but didn't happen in Example 2, which is why I've included it.

2.4. DATABASE 13

- dbClearResult()
- dbConnect()
- dbCreateTable()
- dbDataType()
- dbDisconnect()
- dbExecute()
- dbFetch()
- dbGetQuery()
- dbListTables()
- dbReadTable()
- dbRemoveTable()
- dbSendQuery()
- dbSendStatement()
- Id()
 - See example below.
- SQL()
 - See example below.

{dbplyr}

```
• in_schema()
```

• tbl()

{odbc}

- dbConnect()
 - See example below for specifying a database.
 - See example below for troubleshooting hang-ups.

2.4.1 Examples

```
DBI::Id() vs DBI::SQL():
```

The following are comparable methods of accessing <database>.<schema_name>.<table_name>:

```
dbCreateTable(
   conn,
   name =
     Id(catalog = "database_name", schema = "schema_name", table = "table_name"),
   fields = sample_data_frame
)

dbCreateTable(
   conn,
   name = SQL("database_name.schema_name.table_name"),
   fields = sample_data_frame
)
```

odbc::dbConnect():

```
# Specify database:
conn_1 <- dbConnect(odbc::odbc(), dsn = "sql") # Default database: Actuary

conn_2 <- dbConnect(odbc::odbc(), dsn = "sql", Database = "Staging")

# Troubleshoot hang-ups
# It appears that RStudio's attempt to load information into the Connection Pane,
# via `odbc::dbConnect`, can sometimes cause the call to hang, indefinitely.
# Use the following code to access the database in such a situation:

options(connectionObserver = FALSE)</pre>
```

2.4.2 References

• "Databases using R"

2.4.3 SQL and R

2.4.3.1 Tips and Troubleshooting Hints

- Temp Tables
 - Use SET NOCOUNT ON in the SQL query to prevent R confusing the SQL rowcount return value with the temporary table.
- Try case-sensitive syntax (e.g., when querying a database other than the default).
- \bullet Try selecting specific columns, rather than SELECT *, as SQL BLOB columns cause trouble.

2.4.3.2 SQL References

- "10 Easy Steps to a Complete Understanding of SQL"
- "Query Planning"

2.5 Notes & References

- "Data structures" (Wickham, Advanced R)
- {openintro} contains data sets useful for practicing and teaching.
- "Vectors" (Grolemund & Wickham, R for Data Science)

Tidy

"Tidying your data means storing it in a consistent form that matches the semantics of the dataset with the way it is stored. In brief, when your data is tidy, each column is a variable and each row is an observation. Tidying data is important because the consistent structure lets you focus your struggle on questions about the data, not fighting to get the data into the right form for different functions."

- Garrett Grolemund & Hadley Wickham, R for Data Science

3.1 Explore Raw Data

{base}

- attr()
 - attr(x, "names") is the same as base::names(x).
 - Use attr(x, "names") <- value to set attribute values.
- attribute()s
- class()
- colnames()
- dim()
- dimnames()
- format()
- length()
- levels()
 - Levels default to alphabetical order, so be careful when renaming them (i.e., don't accidentally set the "F" level equal to "Male" rather than "Female").

```
• mode()
   • names()
   • nchar()
   • order()
   • rownames()
   • summary()
   • typeof()
{graphics}
   • hist()
   • plot()
{skimr}
   • skim()
       - Similar to tibble::glimpse() and utils::str().
{tibble}
  • glimpse()
       - Similar to skimr::skim() and utils::str().
{utils}
   • head()
   • str()
       - Similar to skimr::skim() and tibble::glimpse().
   • tail()
       Tidy Data
3.2
{base}
  • duplicated()
       - See also data.table::duplicated()
   • unique()
       - See also data.table::unique()
```

3.2. TIDY DATA 19

```
{data.table}
   • anyDuplicated()
   • duplicated()
       - See also base::duplicated()
   • unique()
       - See also base::unique()
{janitor}
   • clean_names()
   • get_dupes()
       - Prefer over base::duplicated() and data.table::duplicated().
   • remove_empty()
{purrr}
   • transpose()
       - Turn a list-of-lists inside-out.
{splitstackshape}
   • cSplit()
       - See also tidyr::separate()
{tibble}
   • rowid_to_column()

    Add a column of sequential row IDs.

       -\, Useful when a separate ID is required to manipulate rows or columns,
         such as when using tidyr::pivot_longe()r
   • rownames()
{tidyr}
   • nest()
       - Nest repeated values in a list-variable.
```

- Helpful when separating a data frame in preparation to model the data for each grouping.
- replace_na()
- separate()
- pivot_longer()
- pivot_wider()
- unite()
- unnest()

3.3 Prepare Data for Analysis

3.3.1 Coerce

```
{base}
```

- as.array()
- as.data.frame()
 - See also tibble::as_tibble().
- as.Date()
- as.factor()
- as.list()
- as.matrix()
- as.POSIX*()
- factor()
 - Rearrange the order of factors by using the levels argument. For example, rearrange the order of "Bad,"Good," and "Neutral" using 'levels = c("Bad", "Neutral", "Good").
- is.na()
 - Use is.na <- to set elements to NA.
- unclass()

{methods}

• as()

{tibble}

- as_tibble()
 - Preferable to base::as.data.frame().
- enframe()
 - Preferable to using tibble::as_tibble() to coerce a vector to a data frame.

3.3.2 Dates and Datetimes

{anytime}

• anytime(): Parse POSIXct or Date objects from input data.

{base}

- as.Date()
- OlsonNames()
 - Displays available time zones.
- as.POSIXct.*()
- as.POSIXlt.*()
- strptime()
 - Date-time conversion to and from character.
- Sys.timezone()
 - See also base::OlsonNames().

{fasttime}

- fastPOSIXct()
 - Convert strings into POSIXct object (string must be in year, month, day, hour, minute, second format.)

{hms}

- as.hms()
- hms()
 - Store time-of-day values as hms class.
- is.hms()

{lubridate}

- as_date()
- fast_strptime()
 - Fast C parser of numeric formats only that accepts explicit format arguments (just as base::strptime()).
 - Note that the format argument must match the input exactly, including any non-white space characters (such as "T" and "Z").

- make_date()
 - Create dates from numeric representations.
- make_datetime()
 - Create date-times from numeric representations.
- parse_date_time()
 - This function can be slow, because it is designed to be forgiving and flexible. If the dates you are working with are in a consistent format (ideally ISO 8601), use one of the following: fasttime::fastPOSIXct() or lubridate::parse_date_time2().
- parse_date_time2()
 - Fast C parser of numeric orders.
- ymd(): Parse dates with year, month, and day components.
 - Related formats: ydm(), mdy(), myd(), dmy(), dym(), yq().
- ymd_hms(): Parse date-times with year, month, day, hour, minute, and second components.
 - Related formats: ymd_hm(), ymd_h(), dmy_hms(), dmy_hm(),
 dmy_h(), mdy_hms(), mdy_hm(), mdy_h(), ydm_hms(), ydm_hm(),
 ydm_h().

3.3.3 Filter

{dplyr}

• filter()

{purrr}

• keep()

{stats}

• na.omit()

3.3.4 Strings

```
{base}
```

- cat()
 - Concatenate and print.
- chartr()
 - Change certain characters.
- gregexpr()
- grep()
- grepl()
- gsub()
- regexec()
- regexpr()
- sub()
- tolower()
 - See also stringr::str_to_lower().
- toupper(): Convert to uppercase.
 - See also stringr::str_to_upper().

{qdap}

• check_spelling()

{qdapDictionaries}

- DICTIONARY()
 - Nettalk Corpus syllable data set.
- GradyAugmented()
 - Augmented list of Grady Ward's English Words and Mark Kantrowitz's Names List.
 - Mark Kantrowitz's *Names* list is available in full here.

{stringr}

- str_detect
 - Control the pattern argument options with regex() (e.g., str_detect(x, regex(pattern, ignore_case = TRUE)).
- str_remove()
- str_to_lower()
- str_to_title()
- str_trim()
- str_to_upper()

3.3.5 Test

```
{base}
```

- all()
- any()
 - Use any(is.na(data.frame)) to determine if there are any NA values in a data frame.
- anyNA()
 - Possibly faster implementation of base::any(is.na(x)).
- exists()
- is.array()
- is.data.frame()
- is.matrix()
- is.vector()
- setequal()
 - Check two vectors for equality.
- sum()
 - Test whether all elements of a vector do or do not meet a certain condition, use as follows: sum(email\$num_char < 0).

{purrr}

• every()

{stats}

- complete.cases()
 - $-\,$ Find complete cases (i.e., rows without NA values).

{tibble}

• is_tibble()

Transform

"Transformation includes narrowing in on observations of interest (like all people in one city, or all data from the last year), creating new variables that are functions of existing variables (like computing velocity from speed and time), and calculating a set of summary statistics (like counts or means)."

- Garrett Grolemund & Hadley Wickham, R for Data Science

4.1 Arithmetic & Summary Statistics

{assertive}

• is_divisible_by()

{base}

- abs()
- colMeans()
- colSums()
- diff()
- IQR()
- max()
- mean()
 - Note that base::mean() can be used to calculate a percentage when used in dplyr::summarize(), as follows:

```
by_country <-
  votes %>%
  group_by(country) %>%
  summarize(total = n(),
             percent_yes = mean(vote == 1))
  • median()
   • min()
   • Mode: Use table to view the mode of a data set.
   • Operators:
       - Arithmetic Operators: +, -, *, /, %%
       - Comparison Operators: \langle , \rangle, \langle =, \rangle = = , !=.
            * Use identical() and (all.equal)) rather than == and != in
              tests where a single TRUE or FALSE is required (such as if ex-
              pressions).
  • range()
       - Use diff(range()) to get the range as a measure of variability.
   • round()
   • rowMeans()
   • rowSums()
  • scale()
   • sd()
  • signif()
       - Round values to a specified number of significant digits.
   • sqrt()
  • sum()
   • summary()
   • var()
{dplyr}
  • count()
  • group_by()
   • n()
       - Must be used within summarize(), mutate(), or filter().
  • n_distinct()
   • near()
       - Compare two numeric vectors (safer than using ==).
   • summarize()
```

```
- Use mean(variable == value) to get a percentage (see the above
         example for mean().)
       - See dplyr::n_distinct().
   • tally()
   • top_n()
{dummies}
   • dummy.data.frame()
       - Dummify a data frame (useful when needing to calculate the Jaccard
         index for categorical data)
{magrittr}
   • extract()
       - x \%\% extract(y) is equivalent to x[y].
   • extract2()
   • multiply_by()
       - x \% multiply_by(y) is equivalent to x * y.
   • raise_to_power()
       - x \% raise_to_power(y) is equivalent to x^y.
{stats}
   • aggregate()
       - Compute summary statistics of data subsets.
   • cor()
   • cov()
   • cutree()
       - Pair with stats::hclust().
   • dist()
   hclust()
   • lag()
   • rnorm()
   • var()
```

4.2 Create New Variables or Modify Existing Ones

countrycode

• countrycode(): Convert country codes into country names.

dplyr

- mutate(): Add new variables.
 - mutate() can also be used to modify existing variables. To change the case of a character variable, for example, do something like:
 - Child function: transmute() (drops existing variables).
- recode(): Recode values (the numeric alternative to using if_else or case_when()).

Example: dplyr::mutate

```
df <-
    df %>%
    mutate(var_name = str_to_lower(var_name))

# Add a T/F vector, which can then be referenced to highlight certain
# information in a plot:

df <-
    df %>%
    mutate(hilight = attributed_provider_id = 95597)
```

4.3 Dates and Datetimes

- base::date
 - Get the current system date and time.
- base::difftime
 - Time intervals and differences.
 - base::difftime is the function behind the operator when used with dates and datetimes (e.g., time_1 - time_2 is equivalent to difftime(time_1, time_2)). The advantage of using difftime over -, however, is the units argument because it allows you to specify the unit of time in which the difference is calculated.

- base::months
 - Extract the month names.
- base::quarters
 - Extract the calendar quarters.
- base::seq.Date
- base::Sys.Date
 - Get the current date in the current time zone.
- base::Sys.time
 - Get the absolute date-time value (which can be converted to various time zones and may return different days).
- base::weekdays
 - Extract weekday names.
- lubridate::ceiling_date
- lubridate::date
 - Get or set the date component of a date-time.
- lubridate::day
 - Get or set the day component of a datetime.
- lubridate::floor date
- lubridate::month
 - Get or set the month component of a datetime.
- lubridate::now
 - The current time (as a POSIXct object).
- lubridate::quarter
 - Get or set the fiscal quarter or semester component of a datetime.
- lubridate::round_date
- Time spans: Duration
 - Use when you are interested in seconds elapsed.
 - lubridate::ddays
 - lubridate::dhours
 - lubridate::dminutes
 - lubridate::dseconds
 - lubridate::dweeks
 - lubridate::dyears
- Time spans: Interval
 - Use when you have a start and end.

```
- %--%
```

- int_aligns
- int_diff
- int_end
- int_flip
- int_length
- int_overlaps
- int_shift
- int_standardize
- int_start
- interval
- is.interval
- Time spans: Period
 - Use when you are interested in human units.
 - lubridate::day
 - lubridate::hour
 - lubridate::minute
 - lubridate::month
 - lubridate::second
 - lubridate::week
 - iubiiuate..weer
 - lubridate::year
- Time zones:
 - lubridate::force_tz
 - * Change the time zone without changing the clock time.
 - lubridate::tz
 - * Extract the time zone from a datetime.
 - lubridate::with_tz
 - * View the same instant in a different time zone.
- lubridate::today
- lubridate::%m+% & %m-%
 - Add and subtract months to a date without exceeding the last day of the new month.
- lubridate::%within%
 - Test whether a date or interval falls within an interval.
- lubridate::year
 - Get or set the year component of a datetime.

4.4. FACTORS 31

4.4 Factors

forcats

- fct_drop(): Drop levels.
- fct_reorder(): Reorder levels, based on the value of another variable.
- fct rev(): Reverse levels.

stats

- reorder(): Reorder levels of a factor.
 - Useful within the aes() argument in a ggplot() call.

4.5 Merge or Append Data

base

- append(): Add elements to a vector.
- cbind(): Combine objects by column.
- intersect(): Combine data shared in common between two datasets.
 - Similar to dplyr::semi_join().
- merge(): Merge two data frames.
 - dplyr::join functions are an alternative to merge().
- rbind(): Combine objects by row.
- setdiff(): Find the difference between two vectors.
 - Similar to dplyr::anti_join().
- union(): Combine two datasets without duplicating values.

dplyr

- bind(): Bind multiple data frames by row and column.
 - Child functions: bind_rows(), bind_cols(), combine().
- Join Functions: Join two tables.
 - Filtering Joins:
 - * anti_join(): Return all rows from x where there are not matching values in y, keeping just columns from x.

* semi_join(): Return all rows from x where there are matching values in y, keeping just columns from x. A semi join differs from an inner join because an inner join will return one row of x for each matching row of y, where a semi join will never duplicate rows of x.

- Mutating Joins:

- * full_join(): Return all rows and all columns from both x and y. Where there are not matching values, returns NA for the one missing.
- * inner_join(): Return all rows from x where there are matching values in y, and all columns from x and y. If there are multiple matches between x and y, all combination of the matches are returned.
- * left_join(): Return all rows from x, and all columns from x and y. Rows in x with no match in y will have NA values in the new columns. If there are multiple matches between x and y, all combinations of the matches are returned.
- * right_join(): Return all rows from y, and all columns from x and y. Rows in x with no match in y will have NA values in the new columns. If there are multiple matches between y and x, all combinations of the matches are returned.

tibble

- add_column(): Add columns to a data frame.
- add_row(): Add rows to a data frame.

4.6 Narrow in on Observations of Interest

{base}

- droplevels: Drop unused levels from factors.
 - This function will keep levels that have even 1 or 2 counts. If you want to remove levels with low counts from a data set in order to simplify your analysis, first filter out those rows and then use droplevels.
- order: Arrange in ascending or descending order.
 - Alternative: dplyr::arrange.
- prop.table: Express table entries as proportions of the marginal table.
 - The input is a table produced by table.
 - As these are proportions of the whole, sum(prop.table(table_name))
 = 1.

4.7. TEST 33

- Specify conditional proportions on rows or columns by using the margin argument.
- table: Build a table of the counts at each combination of factor levels.
 - Use prop.table to see the table entries expressed as proportions.
- dplr::all equal(): Compare data frames.
- dplyr::arrange(): Arrange rows by variable, in ascending order.
 - Related functions: arrange_all(), arrange_at(), arrange_if().
- dplyr::distinct(): Select distinct rows.
- dplyr::filter(): Return rows with matching conditions.
 - Use %in% when using multiple | conditions. The following two commands, for example, are equivalent:

```
ilo_data %>%
  filter(country %in% c("Sweden", "Switzerland"))

ilo_data %>%
  filter(country == "Sweden" | country == "Switzerland")
```

- dplyr::rename(): Rename variables by name (a modification of select).
- dplyr::sample_n(): Sample n rows from a table.
- dplyr::select(): Select/rename variables.
 - Helper functions include: starts_wit, ends_with, contains, matches, num_range, and one_of. See ?dplyr::select_helpers.
 - Helper functions take strings (e.g., contains("work") rather than contains(work)).
- dplyr::slice(): Choose rows by position.
- dplyr::transmute(): Create or transform variables.
 - Like a combination of dplyr::select and dplyr::mutate.

4.7 Test

- base::identical(): Test objects for exact equality.
- base::match(): Value matching.
 - %in% is the more intuitive binary operator.
- base::setequal(): Check two vectors for equality.
- base::which(): Determine which indices are TRUE.
 - This function is often unnecessary, according to the DataCamp article "Five Tips to Improve Your R Code".

- base::which.min()/base::which.max(): Where is the min()/max() or first TRUE/FALSE?
- %in%: See base::match().

Visualize

"Visualisation is a fundamentally human activity. A good visualisation will show you things that you did not expect, or raise new questions about the data. A good visualisation might also hint that you're asking the wrong question, or you need to collect different data. Visualisations can surprise you, but don't scale particularly well because they require a human to interpret them."

- Garrett Grolemund & Hadley Wickham, R for Data Science ***

5.1 Interfaces

• shiny

5.2 Graphs, Charts, and Plots

• base::abline()

Add straight lines to a plot.

base::plot()base::points()

- Add points to a plot.

- diagram
- DiagrammeR
- GGally::ggally_box()
- GGally::ggpairs()
 - Scatter plot matrix (SPLOM).
- GGally::ggparcoord()
 - Parallel coordinate plots.
- ggbeeswarm::geom beeswarm()
- ggplot2::aes()
 - Arguments include color, fill, size, labels, alpha, shape (1-20 accept color attributes and 21-25 accept color and fill attributes), linewidth, linetype, and group.
 - Use ?pch to see options for shape.
 - Note that aes can be called within geom_* rather than prior to.
 - The default shape for points does not have a fill attribute, which
 means that mapping a categorical variable onto fill won't result in
 multiple colors.
 - Helper functions to include in the call when needing to modify the data include: stats::reorder().
- ggplot2::coord_cartesian()
 - Zoom a plot in or out without changing the underlying data.
- ggplot2::coord_flip()
 - Flip the x and y axes.
- ggplot2::coord_polar()
 - Use to convert a stacked bar chart to a pie chart.
- ggplot2::element_blank()
- ggplot2::element_line()
- ggplot2::element rect()
- ggplot2::element_text()
- ggplot2::facet_grid()
 - Lay out panels in a grid.
 - ROWS ~ COLS: When using a tilde function, the variable on the left specifies the rows and the variable on the right specifies the columns.
 When faceting in only one direction (e.g., only on rows), use . to specify nothing for the unused direction.
- ggplot2::facet_wrap()
 - Wrap a 1D ribbon of panels into 2D (observe a variable, conditional on another variable).

```
• ggplot2::geom_abline()
```

Add reference lines to a plot.

```
• ggplot2::geom_bar()
```

- Create a bar chart, where the height of the bar is proportional to the number of cases in each group.
- ggplot2::geom_boxplot()
- ggplot2::geom_col()
 - Create a bar chart, where the height of the bar represents values in the data.
- ggplot2::geom_density()
 - Create a kernal density estimate (a smoothed version of a histogram).
 - Consider using ggplot2::geom_rug() with ggplot2::geom_density() in order to be transparent about smoothed data.
- ggplot2::geom_dotplot()
 - Create a histogram out of dots.
- ggplot2::geom_errorbar()
 - Shortcut for geom_bar(stat = "identity").
- ggplot2::geom_histogram()
 - Use y = stat(density) to rescale the y-axis from counts to a probability estimate.
- ggplot2::geom_hline() *ggplot2::geom_jitter()
 - Useful with ggplot2::geom_boxplot().
- ggplot2::geom_path()
 - Connect observations in the order in which they appear.
- ggplot2::geom_point()
- ggplot2::geom_rug()
 - Useful with ggplot2::geom_density().
- ggplot2::geom_smooth()
 - Smoothed conditional means; aids the eye in seeing patterns in the presence of overplotting.
- ggplot2::geom_text()
- ggplot2::geom_tile()
 - Heat maps.
- ggplot2::geom_violin()

```
• ggplot2::geom_vline()
• ggplot2::ggplot()
• ggplot2::ggtitle()
• ggplot2::group()
    - Used within aes().
    - Note that group is usually unnecessary when specifying color,
       shape, fill, or linetype within aes, or when using facets. See
       "Aesthetics: grouping" for more information.
• ggplot2::labs()
    - Modify axis, legend, and plot labels.
• ggplot2::position_identity()
    - Don't adjust position.
• ggplot2::position_dodge()

    Dodge overlapping objects side-to-side.

• ggplot2::position_nudge()
    - Nudge points a fixed distance.
• ggplot2::position_stack()
    - Stack overlapping objects on top of each other, as counts.
• ggplot2::position_fill()
    - Stack overlapping objects on top of each other, as densities.
• ggplot2::position_jitter()
• ggplot2::position_jitterdodge()
• ggplot2::scale_x_continuous()
• ggplot2::scale_y_*()
• ggplot2::scale_color_manual()
• ggplot2::scale_fill_*()
• ggplot2::scale_shape_*()
• ggplot2::scale_size()
• ggplot2::scale_linetype_*()
• ggplot2::stat_bin()
• ggplot2::stat_bin2d()
• ggplot2::stat_bindot()
• ggplot2::stat_binhex()
• ggplot2::stat_boxplot()
• ggplot2::stat_contour()
```

ggplot2::stat_quantile()ggplot2::stat_smooth()ggplot2::stat_sum()ggplot2::theme()

```
- See example for grid::unit().
• ggplot2::theme_get()
• ggplot2::theme_set()
• ggplot2::theme_update()
    - "When you call 'theme_update' and assign it to an object (e.g., called
       old), that object stores the current default theme, and the argu-
       ments update the default theme. If you want to restore the previous
       default theme, you can get it back by using theme update again."
       -DataCamp
• ggplot2::xlab()
• ggplot2::xlim()
• ggplot2::ylab()
• ggridges::geom_density_ridges()
    - Create a ridgeline plot.
• graphics::boxplot()
    - Create a box-and-whisker plot.
• graphics::hist()
• graphics::par()
• graphics::stripchart()

    One dimensional scatter plots.

    - Prefer over ggplot when creating one-dimensional plots.
• grDevices::colorRamp()

    Color interpolation.

• grDevices::colorRampPalette()

    Color interpolation.

• grid::unit()
    - Create a unit object, to specify locations and dimensions in a coor-
       dinate system.

    See example below.

• plotly::plot_ly()
```

- Make the ColorBrewer color palettes available as R palattes.

5.2.0.1 Examples

• RColorBrewer::brewer.pal()

• grid::unit

```
# Increase spacing between facets in a `ggplot` object:

plot +
   theme(
    panel.spacing.x = unit(2, "cm"),
    panel.margin = unit(c(1, 2, 1, 1), "cm"))
```

5.3 Graphs, Charts, and Plots - Statistics

```
ggplot2::mean_cl_boot
ggplot2::mean_cl_normal
ggplot2::sdl
ggplot2::median_hilow
ggplot2::stat_function

Compute y values from a function of x values.

ggplot2::qq

Perfrm calculations for a quantile-quantile plot.

ggplot2::qq_line
ggplot2::summary
Summarize y values at distinct x values.
```

Chapter 6

Model

"Models are complementary tools to visualisation. Once you have made your questions sufficiently precise, you can use a model to answer them. Models are a fundamentally mathematical or computational tool, so they generally scale well. ... But every model makes assumptions, and by its very nature a model cannot question its own assumptions. That means a model cannot fundamentally surprise you. - Garrett Grolemund & Hadley Wickham, R for Data Science" ***

6.1 Cluster Analysis

{stats}

- cutree()
 - Pair with stats::hclust().
- dist()
- hclust()
- kmeans()

6.2 General

- base::sampledplyr::sample_nstats::coef
 - Extract model coefficients.

stats::corstats::covstats::cov2cor

- Scale a covariance matrix into a correlation matrix.

• stats::var

6.3 Regression

- broom::augment
 - Augment data with information from an object.
- broom::glance
 - Construct a single row summary of a model, fit, or other object.
- broom::tidy
 - Turn an object into a tidy tibble.
- {mgcv::gam
 - $-\,$ Generalized additive models (GAMs) with integrated smoothness estimation.
- mixtools::
- stats::df.residual
 - Get the residual degrees of freedom.
- stats::lm
 - Fit linear models.
- stats::p.adjust
 - Adjust p-values for multiple comparisons.
- stats::predict
 - Model predictions.
- stats::residuals
 - Extract model residuals.
- tidyr::nest
 - Nest repeated values in a list-variable.
 - Helpful when separating a data frame in preparation to model the data for each grouping.

6.4 Simulation & Prediction

- base::set.seedstats::predict
 - Model prediction.
 - Use with lm class objects and new data to predict new values (e.g., predict(model, newdata)).
- stats::rnorm

6.5 Strings

- base::agrep
- fuzzyjoin::stringdist_join
- fuzzyjoin::stringdist_anti_join
- fuzzyjoin::stringdist_full_join
- fuzzyjoin::stringdist_inner_join
- fuzzyjoin::stringdist_left_join
- fuzzyjoin::stringdist_right_join
- fuzzyjoin::stringdist_semi_join
- fuzzywuzzyR::FuzzMatcher

Chapter 7

Communicate

"The last step of data science is communication, an absolutely critical part of any data analysis project. It doesn't matter how well your models and visualisation have led you to understand the data unless you can also communicate your results to others."

- Garrett Grolemund & Hadley Wickham, R for Data Science ***

7.1 CSS

Center Title

7.2 Export

• base::file.path

• base::print

Use the include.rownames = FALSE argument to remove row numbers (or names) from the output.

base::savebase::saveRDS

 See "A better way of saving and loading objects in R" to understand the differences between save and saveRDS.

- readr::write_csvreadr::write_delim
 - About twice as fast as utils::write.csv, and never writes row names.

• readr::write_excel_csv

readr::write_tsvutils::write.csvutils::write.csv2utils::write.table

- Prefer readr::write_delim to utils::write.table.

• XLConnect::loadWorkbook

7.3 Format Output

base::formatDT::datatable

- Use options = list(dom = 't') to remove the search bar.
- Use options = list(dom = "t", lengthchange = FALSE) to remove the search bar and box that allows the user to choose how many rows to see.
- Use options = list(columnDefs = list(list(className =
 "db-center", targets = (column_position)))) to center
 column values (where 0 indicates the first column).
- DT::formatCurrency
- gt::gt
 - Alternative to knitr::kable and DT::datatable, styled after ggplot2.

kableExtra::kable_stylingkableExtra::add_footnote

knitr::kablelubridate::stamp

- Format dates and times based on human-friendly templates.
- scales::dollar

7.4. GRAPHICS 47

7.4 Graphics

**

- knitr::include_graphics
 - Embed external images in 'knitr' documents.
 - Preferable to the ![alt text or image title](path/to/image)
 Markdown syntax for embedding an external image, as include_graphics offers more control over the attributes of the image.

7.5 Plots

• ggplot2::element_*: Specify the display of how non-data components of a plot are drawn.

• ggplot2::labs: Modify axis, legend, and plot labels.

- Child functions: xlab, ylab, ggtitle

• ggplot2::theme: Customize the non-data components of a plot.

See example below.

- See ?theme_classic for a list of predefined themes.

Example: ggplot2::theme

```
ggplot(plot_data_2006) +
  geom_histogram(aes(x = working_hours)) +
  labs(
    x = "Working hours per week",
    y = "Number of countries") +
  theme(
    text = element_text(family = "Bookman", color = "gray25"))
```

7.6 RMarkdown

• See RMarkdown: The Definitive Guide

[•] See Mozzila Developer Network for CSS help.

Chapter 8

Program

"Surrounding [the tools for importing, tidying, transforming, visualising, modeling, and communicating data] is programming. Programming is a cross-cutting tool that you use in every part of a project. You don't need to be an expert programmer to be a data scientist, but learning more about programming pays off because becoming a better programmer allows you to automate common tasks, and solve new problems with greater ease."

- Garrett Grolemund & Hadley Wickham, R for Data Science

8.1 Characters

References: * ?Quotes

8.2 Conditionals & Control Flows

base

• Control (access documentation using "?Control")

- if (cond) expr

* The key difference between if (cond) expr and ifelse is that if (cond) expr will evaluate only the first element of an object with length > 1. See the documentation for each function and

the ateucher's GitHub example titled "R: if vs ifelse" to learn more.

- for (var in seq) expr
- while (cond) expr
- repeat expr
- break
- next
- identical(): Test objects for exact equality.
 - Use identical() rather than == and != in if and while statements to test for equality.
- 'ifelse(): Conditional element selection. +dplyr::if_else()' is more strict by checking the object type.
- stop(): Stop execution of the expression and execute an error action.
 - Useful in combination with an if statement when you want to generate helpful error messages.
- stopifnot(): Ensure the truth of an R expression.
 - Prefer base::stop() to base::stopifnot().

dplyr

- case_when(): A general vectorized if.
- if_else(): Vectorized if.

8.3 Environment and Workspace

- base::.libPaths()
- base::baseenve()
 - The environment of the base package; it's enclosing environment ("parent environment") is the empty environment.
- base::dir()
 - List the files in a directory.
- base::dir.create()
 - Create a file path in the computer's file system.
- base::emptyenv()
 - The empty environment, which is the ancestor of all environments and the only environment without an enclosing environment.
- base::environment()

- The current environment.
- base::environmentName()
- base::exists()
 - Remember that R will look for an object in parent environments until it reaches the empty environment, so use inherits = FALSE to limit the search to only the current environment.
- base::getOption()
 - Set and examine global options.
- base::getwd()
 - Get the working directory.
- base::globalenv()
 - The environment in which you normally work, it's enclosing environment is the last package attached with library() or require().
- base::history()
 - Display the previous 25 commands.
- base::install.packages()
 - Install packages from repositories or local files.
- base::library()
 - Load and attach packages, returning an error if the packages does not exist.
- base::list.files()
 - List the files in a directory/folder.
- base::loadedNamespaces()
 - Return the loaded name spaces.
- base::loadhistory()
 - Recall command history.
- base::ls()
 - List objects in the specified environment.
- base::list2env()
 - From a list, build or add to an environment.
- base::new.env()
 - Create a new environment.
- base::options()

- Set and examine global options.
- base::parent.env()
 - Return the enclosing environment of the environment listed as an argument.
 - base::parent.env() returns information that can be unhelpful, so use with base::environmentName(), as follows: parent.env(environment_name) %>% environmentName().
- base::q()
 - Terminate an R session.
- base::R.version()
 - Version information.
- base::R.version.string()
 - Version information.
 - Same call as R.version\$version.string().
- base::require()
 - Load and attach packages, returning FALSE if the package does not exist.
- base::rm()
 - Remove objects from a specified environment.
- base::savehistory()
 - Save command history (default value is ".Rhistory").
- base::save.image()
 - Save the current workspace.
- base::search()
 - Return a list of attached packages and R objects.
- base::searchpaths()
 - Return the path to attached packages.
- base::setwd()
 - Set the working directory file path.
 - When using Windows, use "/" instead of "".
- base::Sys.getenv()
 - See also Sys.setenv().
- base::Sys.info()

- Extract system and user information.
- Example: Sys.info()[c("sysname", "release")].
- base::Sys.setenv()
 - See also Sys.setenv().
- gdata::object.size()
 - Report the space allocated for an object.
 - See also utils::object.size().
- here::here()
- installr::updateR()
 - Check for the latest R version; downloads and installs new R versions.
- pryr::where(): Find where a name is defined.
- utils::ls.str(): List objects and their structure.
- utils::object.size(): Report the space allocated for an object.
 - See also gdata::object.size().
- utils::sessionInfo(): Collect information about the current R session.

References:

• "Environments" (Hadley Wickham, Advanced R)

8.4 Evaluation (Standard and Non-standard)

base

- cat(): Concatenate and print.
- print(): Print the argument to the Console. + A shortcut toprint()' is to place the code you want printed inside parentheses.
- quote(): Return the argument, unevaluated.
- writeLines(): Display quotes and backslashes as they would be read, rather than as R stores them (i.e., see the raw contents of the string, as the print() representation is not the same as the string itself).

rlang

- Quosures
 - enquo(), new_quosure(), quo().

References:

- "Non-standard evaluation" (Hadley Wickham, Advanced R)
- "Non-standard evaluation" (Hadley Wickham, lazyeval package vignette)
- "Programming with dplyr" (dplyr.tidyverse.org)

8.5 Functionals

{base}

- Apply Functions
 - apply: Apply functions over array margins.
 - lapply: Apply a function over a list or vector.
 - sapply: Apply a function over a list or vector and return a vector or matrix.
 - vapply: A safer version of sapply, as it requires the output type to be predetermined.
 - mapply: Apply a function to multiple list or vector arguments.
 - rapply: Recursively apply a function to a list.
 - tapply: Apply a function over a ragged array.

{purrr}

- map: Apply a function to each element of a vector.
 - tidyr::unnest is useful in changing the list-column output of map into rows.
- map2: Map over multiple inputs simultaneously.
- map_if: Apply a function to elements of that match a condition.
- possibly: Usese a default value whenever an error occurs.
- quietly: Capture side effects in a list with components result, output, messages, and warnings.
- safely: Capture side effects in a list with components result and error.
- transpose: Transpose a list (turn a list-of-lists inside-out).

8.6. FUNCTIONS 55

8.6 Functions

• assertive::assert_*(): Check whether the input is * (e..g, assert_is_numeric()) and throw and error if the input does not meet the condition.

- assertive::coerce_to(): Coerce the input to a different class, with a warning.
- assertive::is_*(): Checks whether the input matches the condition specified by * (e.g., assertive::is_non_positive()).
- assertive::use_first(): Use only the first element of a vector.
- base::do.call(): Execute a function call from a name or a function and a list of arguments to be passed to the function.
- base::invisible(): Return a (temporarily) invisible copy of an object.
- base::match.arg(): Argument verification.
 - Useful when matching a character argument specified in the function signature. For example,

```
## function (x, n, p = NULL, alternative = c("two.sided", "less",
## "greater"), conf.level = 0.95, correct = TRUE)
## NULL

# The body of `prop.test` contains the following line of code:
# `alternative <- match.arg(alternative), which reassigns it to the selected</pre>
```

• base::message()

character vector.

- Generate a diagnostic message.
- Preferable to generating a message using cat.
- base::return()
 - Return a value from a function.
 - Useful in if statements where one condition is simple and the other is complex (see section 19.6.1 "Explicit return statements" in Hadley Wickham's R for Data Science.
- base::setNames()
 - Set the names in an object.
 - Useful in function writing; see documentation.
- base::stop()
 - Stop execution of the expression and execute an error action.

- Useful in combination with an if statement when you want to generate helpful error messages.
- base::stopifnot()
 - Ensure the truth of an R expression.
 - See section 19.5.2 "Checking values" in Hadley Wickham's *R for Data Science* for a discussion of stop() versus stopifnot().
 - Consider functions from {assertive} as an alternative to stopifnot and stop.
- base::unlist()
 - Flatten lists.
 - Useful when using purrr's map functions, which return objects as type list.
- zeallot::%<-%: Multiple assignment operator (see example below).

8.6.0.1 Examples

• zeallot::%<-%:

```
session <- function() {
  list(
    r_version = R.version.string,
    operating_system = Sys.info()[c("sysname", "release")],
    loaded_pkgs = loadedNamespaces()
  )
}
c(vrsn, os, pkgs) %<-% session()</pre>
```

8.7 Learn About an Object

- ?object_name
- ??object_name
- base::args(): Display the argument names and default values of a function.
- base::attributes(): View or assign an objects attributes (e.g., class(), dim(), dimnames(), names(), row.names()).
- base::body(): Get or set the body of a function.
- base::colnames(): Retrieve or set column names.
- base::dim(): Retrieve or set the dimnames of an object.
- base::dimnames(): Retrieve or set the dimension names of an object.

8.8. LOOPS 57

- base::formals(): Get or set the formal arguments of a function.
- base::help(): Get the topic documentation.
- base::help.search(): Search the help system for documentation matching a given character string.
- base::vignette(): View a specified package vignette.
- base::rownames(): Retrieve or set row names.

8.8 Loops

- base::seq(): Sequence generation (this functions makes length() unnecessary).
- base::seq_along(): In for loops, safer than using base::ncol() or base::nrow().

8.9 Optimization

microbenchmark::microbenchmark(): Sub-millisecond accurate timing of expression evaluations. + A more accurate replacement of system.time(replicate(1000, expr)).

8.10 Pipes

- magrittr::%<>%: Compound assignment-pipe operator.
- magrittr::%>%: Forward-pipe operator.
- magrittr:: %\$%: Expositions-pipe operator.
- magrittr::add(): +, for pipes.
- magrittr::and(): &, for pipes.
- magrittr::extract():[, for pipes (see also purrr::pluck()).
- magrittr::extract2(): [[, for pipes.
- magrittr::freduce(): Apply a list of functions sequentially.
- magrittr::is_in(): %in%, for pipes.
- magrittr::multiply_by(): *, for pipes.
- magrittr::or(): |, for pipes.
- magrittr::raise_to_power(): ^, for pipes.
- magrittr::subtract(): -, for pipes.

8.11 Popups

• svDialogs::dlg_message(): Display a modal message box (works in Windows, MacOS, and Linux).

• tcltk::tk_messageBox(): Display a generic message box using Tk (Windows-specific).

8.12 Selecting & Subsetting

```
.$variable_name: See example below.
.[["variable_name"]]: See example below.
base::subset()
dplyr::first()
dplyr::las()t
dplyr::rename()
dplyr::rename()
dplyr::select(): Helper functions include contains(), ends_with(), matches(), num_range(), one_of(), starts_with().
ggplot2::cut_number
magrittr::extract()
magrittr::extract2()
purrr:pluck(): See also magrittr::extract2().
```

8.12.0.1 Examples

• .\$variable_name:

```
ui_summary_table <-
aws_vendors %>%
filter(str_detect(vendor_name, "UTAH INTERACTIVE")) %>%
.$vendor_id %>%
map(query_summary_table) %>%
bind_rows()
```

• .[["variable_name"]]:

8.12.0.2 References

• "Indexing lists in #rstats. Inspired by Residence Inn" (Hadley Wickham, Twitter, 14 September 2015)

8.13 Style Guide

- "The Tidyverse Style Guide" by Hadley Wickham.
- General Layout and Ordering (taken from the outdated "Google's R Style Guide")
 - Title
 - Author
 - File description (e.g., purpose of program, inputs, outputs)
 - source() and library() statements
 - Function definitions
 - Executed statements.

8.14 System Commands

```
base::shell()base::system()base::system2()sys::exec()sys::exec_r()
```

8.15 Version Control

- base::update.packages()
 - 00LOCK error when updating a package: Use update.packages(ask
 = FALSE, checkBuilt = TRUE, INSTALL_opts = "--no-lock")
- installr::updateR()
 - Remove old versions from Windows:
 - * Settings > Apps and Features > uninstall old version
 - * Control Panel > All Control Panel Items > Programs and Features > uninstall old versions
- packrat::snapshot(): Capture and store the packages and versions in use.
- packrat::restore(): Load the most recent snapshot to the project's private library.

• Packrat still seems to be under development. The idea sounds good, but in practice the packrat package has caused lots of problems.

8.15.0.1 Git

- A successful Git branching model
- git-scm.com
- Git and GitHub by Hadley Wickham
- Pro Git by Scott Chacon and Ben Straub
- Understanding the GitHub flow
- git branch: List, create, or delete branches.
 - git branch -d <branch_name>: Delete a local branch.
 - * See "Delete branch in RStudio pop-up" for help removing branches in RStudio after removing them from Git.