Notes for the BAN400 Exam

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1 Functions

1.1 Basic functions

Function	Package	Description
mean()	base	Calculates the mean of a vector of numbers
median()	base	Calculates the median of a vector of numbers
sd()	base	Calculates the standard deviation of a vector
var() sum()	base base	of numbers Calculates the variance of a vector of numbers Calculates the sum of a vector of numbers
c()	base	Creates vector
length()	base	The number of elements in a vector or list
ncol()	base	Number of columns of data frame or matrix
nrow()	base	Number of rows of data frame or matrix
min()	base	The smallest value in a set
max()	base	The largest value in a set

1.2 Math

Function	Package	Description
sqrt()	base	Calculates square root of number or vector of
abs()	base	numbers Calculates absolute value of number or vector
sign()	base	of numbers Returns the sign of x
round()	base	Rounds x to n decimal places
ceiling()	base	Rounds up to the nearest integer
floor()	base	Rounds down to the nearest integer
cumsum()	base	Cumulative sum
cor()	base	Correlation

1.3 Reading data

Function	Package	Description
read_csv()	readr	Read csv-file (comma separated values)
read_csv2()	readr	Uses semicolons as separators and keeps commas
read_delim()	readr	Read file with columns separated by any
read_excel() read_fwf()	readxl readr	delimiter Read data from excel files Reads fixed width data
read_tsv() readLines	readr base	Reads tab separated values Read some or all text lines from a connection

1.4 Data wrangling

Function	Package	Description
head()	base	Returns the first few rows of a data frame or
tail()	base	vector Returns the first few rows of a data frame or
filter() select()	dplyr dplyr	vector Returns elements that satisfy conditions Choose specific columns from a data frame
arrange()	dplyr	Sorts rows of a data frame by specified columns
sort() mutate() transmutate()	base dplyr dplyr	Sorts a vector in ascending or descending order Adds or modifies columns in a data frame Creates a new data frame containing only the
summarise()	dplyr	specified computations Summary statistics for columns in a data frame (typically used with grouped data)
group_by()	dplyr	Group data by one or more variables
ungroup()	dplyr	Ungroup data such that subsequent operations to
<pre>left_join()</pre>	dplyr	apply to the entire dataset Returns all values from the first data frame with all columns and values from the second
inner_join()	dplyr	data frame where there is a match Joins two data frames by keeping only records
right_join()	dplyr	that match in both data sets Returns all values from the second data frame with matching columns and values from the first
full_join()	dplyr	data frame where there is a match Returns all values and columns from both data frames and filling in NA where there is no match
semi_join()	dplyr	Filters the first data frame keeping only rows
anti_join()	dplyr	with matching keys in the second data frame Filters the first data frame to keep only rows with no match in the second data frame

1.5 Machine learning

Function	Package	Description
<pre>logistic_reg() set_engine() set_mode() fit() nearest_neighbor()</pre>	tidymodels tidymodels tidymodels tidymodels tidymodels	Specifies a logistic regression model Specifies the computational engine for a model Sets the mode (e.g. classification or regression) for a model Fits the model to data Specifies a k-nearest neighbors model
<pre>tune() finalize_workflow() workflow() add_model() add_recipe()</pre>	tidymodels tidymodels tidymodels tidymodels tidymodels	Marks a parameter for tuning in a model Finalizes the workflow with specific parameters Creates a workflow object Adds a model to a workflow Adds a recipe to a workflow
<pre>tune_grid() select_best() predict()</pre>	tidymodels tidymodels stats	Tunes hyperparameters across a grid of values Selects the best tuning parameter combination based on a metric predicts outcome variable according to a specified model-e.g Predict.lm- predict.glm
<pre>initial_split() vfold_cv()</pre>	rsample rsample	splits dataset in test and training and test data randomly splits data into different folds (for cross validation)
<pre>recipe() grid_space_filling() roc_auc()</pre>	recipes dials yardstick	creates a recipe for processing data for making a search grid calculated the area under the AUC curve

1.6 Many models

Function	Package	Description
<pre>add_predictions() add_residuals() group_by() ungroup() nest()</pre>	modelr modelr dplyr dplyr tidyr	Adds model predictions to a data frame Adds residuals from a model to a data frame Groups data by one or more variables Removes grouping structure from a data frame Creates a nested data frame by collapsing rows into list-columns
unnest() select() pull() pluck()	tidyr dplyr dplyr purrr	Expands list-columns back into regular columns Selects specific columns from a data frame Extracts a single column as a vector Extracts an element from a list or vector by index or name
map()	purrr	Applies a function to each element of a list or vector
map2()	purrr	Applies a function to pairs of elements from two lists
glance()	broom	Generates a summary of model diagnostics in a tidy format

1.7 Parsing

Function	Package	Description
class()	base	Returns the class of an object
typeof()	base	Determines the type or storage mode of any obiect
mode()	base	Indicates the mode of storage
as.numeric()	base	Converts data to numeric format
as.character()	base	Converts data to character format
as.factor()	base	Converts data to factor format
gsub()	base	Find and replace
reshape()	base	Reshape datasets between wide and long formats
pivot_longer()	tidyr	Convert wide data to long format
<pre>pivot_wider()</pre>	tidyr	Convert long data to wide format.
na.omit()	base	Remove missing values
is.na()	base	Check for missing values
ymd ()	lubridate	Parse dates
hms()	lubridate	Parse times
<pre>parse_datetime()</pre>	readr	Parse datetimes
separate()	tidyr	Split one column into multiple columns

1.8 Selecting

Function	Package	Description
any_of()	dplyr	Selects columns that match any of the given
all_of()	dplyr	names in a vector Selects columns that match all the given names
starts_with()	dplyr	in a vector Selects columns whose names start with a
ends_with()	dplyr	specified prefix Selects columns whose names end with a
contains()	dplyr	specified suffix Selects columns whose names contain a specified string
matches()	dplyr	Selects columns whose names match a specified
select()	dplyr	regular expression Selects specified columns from a data frame
slice_min()	dplyr	Selects rows with the smallest values of a
slice_max()	dplyr	variable Selects rows with the largest values of a variable

1.9 Parallel computing

Function	Package	Description
tic() toc() detectCores() makeCluster() registerDoParallel	tictoc tictoc doParallel doParallel doParallel	Initialize time taking of a function Stop time taking of a function Detect cores in CPU Initialize cores Register the cluster
stopCluster() foreach() %dopar%	doParallel doParallel	Close off clusters Designate tasks for parallelization

1.10 Making maps

Function	Package	Description
<pre>distm() st_as_sf() geom_sf() st_crop() st_layers()</pre>	geosphere sf sf & ggplot2 sf sf	Calculate distance between 2 points Converting data frame to geometric object Allows mapping of geometric objects Cropping maps (x and y chords) Check layers of files
st_read() st_transform()	sf sf	Read map data Transform map projection

2 Topics

2.1 Filtering

The symbol | works as an "or" operator, meaning it will return items that satisfy either one or both of the conditions before and after the operator.

```
filter(flights, dest == "IAH" | dest == "HOU")
```

The & is an "and" operator, meaning both conditions need to hold. This is useful in conjunction with the |-operator, since otherwise you can simply just add filters after each other like this:

```
filter(!is.na(flight2), !is.na(flight1))
```

Another operator %in% returns TRUE if an item exists inside a vector:

```
flights %>%
  filter(dest %in% c("IAH", "HOU")
```

To create the opposite logic, i.e. "not in c()", you can use !:

```
flights %>%
filter(!(dest %in% c("IAH", "HOU")))
```

2.2 Selecting

Some basic selection functions from dplyr are starts_with, ends_with and contains.

```
select(flights, starts_with("dep_"), starts_with("arr_"))
```

2.2.1 Regex (regular expressions)

The function matches uses regular expressions. These can be used to match precise patterns.

```
select(flights, matches("^(dep|arr)_(time|delay)$"))
```

Some common Regex meta characters are:

- . Matches any character except newline.
- ^ Matches the beginning of a string.
- \$ Matches the end of a string.
- [] Matches any one of the characters inside the brackets.
- | Logical OR.
- * Matches 0 or more repetitions of the preceding character.
- + Matches 1 or more repetitions.
- ? Matches 0 or 1 repetition (optional match).
- {n,m} Matches between n and m repetitions.

2.3 Loops and iterations

2.3.1 Standard for-loop

```
for(i in 1:n) {
   ... do something with i...
}
```

Note that we can iterate over any type of vector, not just numbers, and we can give the iteration variable any name we want. In the example above it is i.

2.3.2 While loop

Repeat until a certain condition is met. For example

```
i <- 1
while(i < 10) {
  print(i)
  i <- i + 1
}</pre>
```

2.4 Math

The Integer Division Operator %/% performs integer division. It divides two numbers and returns the whole number part of the quotient, discarding any remainder. Example:

```
10 %/% 3 # Returns 3 (quotient without remainder)
```

The Modulo Operator %% operator returns the remainder from the division of two numbers. Example:

```
10 %% 3 # Returns 1 (remainder)
```

2.5 Creating own functions

In R, we have the opportunity to create own functions. In order to do so, we have to stick to the following syntax.

2.5.1 Basic syntax

```
function_name <-
function(input1, input2, input3){
    # In the following, function is defined/written
    example = (input1 + input2) * input3
    return(example) # Using return, we can exactly define function's return
}</pre>
```

2.6 Plotting

We use ggplot2 as the standard package for plotting, and the main function is ggplot. We supply a data frame to the first argument and an aesthetic mapping to the second argument. We add layers of plotting components using the plus sign. A simple example:

```
ggplot(df, aes(x = x_variable, y = y_variable, colour = grouping_variable)) +
  geom_point()
```

Many types of layers may contain other data sets via the data argument and/or updated aesthetic mappings via the mapping argument. Data and mappings are typically inherited from the layer above if not specified in a new layer. There are many types of functions for making further adjustments to labels, titles, axes and other properties. A more complete example may look like this:

2.6.1 Grouping

You can visually group items in plots together using the color argument.

```
ggplot(data = mpg, mapping = aes(x = displ, y = hwy, color=drv))
```

Grouping can also be used in calculations. For example, if you want to create several regression lines based on classification, you can group with group argument:

```
geom_smooth(aes(group = drv)) # Creates separate regression lines.
```

Sometimes need to group by several variables at the same time. You can use unite() to achieve this.

```
whodata %>%
  unite(country_sex, country, sex, remove = FALSE) %>%
  ggplot()+
  geom_line(aes(x=year,y=cases, group=country_sex, color=sex))
```

2.6.2 Stacking plots

You can stack several plots on top of each other. If you stack plots of the same type , it may be useful to distinguish them using color. For example:

```
ggplot(diamonds) +
  geom_freqpoly(aes(x = x, color = "x"), binwidth = 0.1) +
  geom_freqpoly(aes(x = y, color = "y"), binwidth = 0.1)
```

2.6.3 One categorical and one continuous variable

To explore the relationship between a continuous variable and one categorical variable, some plotting options are:

- geom_violin
- geom_freqpoly With colour-argument
- geom_histogram Withfacet_wrap(vars())
- If the data set is large, you may want to use geom_lv from the library lvplot.

Example:

```
ggplot(diamonds, aes(x=price,colour=cut))+
  geom_freqpoly(bins=50)
```

2.6.4 Two categorical variables

To explore the relationship between two categorical variables, geom_tile is often useful.

```
diamonds %>% count(cut,color) %>% ggplot(aes(x=cut,y=color))+
  geom_tile(aes(fill=n))
```

If you want to normalize proportions within a given categorical variable, here's an example of that:

```
diamonds %>%
  count(color, cut) %>%
  group_by(color) %>%
  mutate(
    prop=n/sum(n)
) %>%
  ggplot(aes(x=color,y=cut))+
  geom_tile(aes(fill=prop))
```

2.6.5 Two continuous variables

There are many options to explore the relationship between two categorical variables, such as:

- geom_point
- geom_smooth
- geom_boxplot where a continuous variable is divided into bins.

```
ggplot(diamonds, aes(x = cut_width(price, 2000, boundary = 0), y = carat)) +
geom_boxplot(varwidth = TRUE)
```

Tip: choosing binned plots versus such as scatterplots depends on the nature of data analysis. For example, binned plots can "hide" outliers that are unusual combinations of x and y, but where x or y individually are not extreme values.

2.6.6 Discrete variables

A small tip about plotting discrete variables: If there is overplotting (e.g. multiple "dots" overlapping in dotplots; which happens often with discrete variables), then use jitter functions.

```
ggplot(data, aes(x = category, y = value)) +
geom_jitter(width = 0.2, height = 0.1)
```

2.6.7 Statistics

Most geoms come in pairs with complementary statistics arguments that are almost always used in concert. These functions can be used to retrieve the data that is used to generate the plot. For example, for these geoms:

```
geom_smooth()
geom_dotplot()
geom_point()
geom_bar()
```

We have these stat functions respectively:

```
stat_smooth()
stat_bindot()
stat_qq()
stat_count()
```

The corresponding stat functions can be found by reading the documentation with for example <code>?geom_smooth.</code>

2.7 Reading data

Sometimes data contains the delimiter. In that case, use quote argument to escape:

```
read_csv("x,y\n1,'a,b'", quote="'")
```

2.8 Parsing data

There are various parsing functions from $lubdridate\ library$, such as ymd(). These functions can be used to convert objects into time and date formats.

2.8.1 Dates and time

Examples:

```
d1 <- "January 1, 2010"
d2 <- "2015-Mar-07"
d3 <- "06-Jun-2017"
d4 <- c("August 19 (2015)", "July 1 (2015)")
d5 <- "12/30/14" # Dec 30, 2014
t1 <- "1705"
t2 <- "11:15:10.12 PM"
```

Can be parsed respectively:

```
mdy(d1)  #> [1] "2010-01-01"

ymd(d2)  #> [1] "2015-03-07"

dmy(d3)  #> [1] "2017-06-06"

mdy(gsub("\\(", "", gsub("\\)", "", d4)))  #> [1] "2015-08-19" "2015-07-01"

mdy(d5)  #> [1] "2014-12-30"

hm(t1)  #> [1] "17:05:00"

hms(t2)  #> [1] "23:15:10.12"
```

2.8.2 Pivoting

The library tidyr provides various ways to pivot data. An example of a pivot is from

```
preg <- tribble(
    ~pregnant, ~male, ~female,
    "yes", NA, 10,
    "no", 20, 12
)</pre>
```

To a longer data format:

```
preg_tidy2 <- preg %>%
  pivot_longer(c(male, female), names_to = "sex", values_to = "count", values_drop_na = TRUE)
```

2.8.3 Splitting columns

You can split a column into multiple columns using separate() and extract() from tidyr:

```
tibble(x = c("a,b,c", "d,e,f,g", "h,i,j")) %>%
separate(x, c("one", "two", "three"))
```

In the above case, the second item has four "values" $^{"d,e,f,g"}$. In this case, the g is dropped. There are various arguments in the functions to determine how to deal with extra or missing items like this.

2.8.4 Basic find and replace

You can perform a basic find-and-replace operation on a vector like this:

y[y == "find"] <- "replace"</pre>