

Data Cleaning

Data Cleaning

In general, data cleaning is a process of investigating your data for inaccuracies, or recoding it in a way that makes it more manageable.

□ MOST IMPORTANT RULE - LOOK □ AT YOUR DATA! □

Dealing with Missing Data

Missing data types

One of the most important aspects of data cleaning is missing values.

Types of “missing” data:

- **NA** - general missing data
- **NaN** - stands for “**Not a Number**”, happens when you do $0/0$.
- **Inf** and **-Inf** - Infinity, happens when you take a positive number (or negative number) by 0.

Finding Missing data

Each missing data type has a function that returns **TRUE** if the data is missing:

- `NA - is.na`
- `NaN - is.nan`
- `Inf and -Inf - is.infinite`

Useful checking functions

- `is.na` - is TRUE if the data is FALSE otherwise
- `!` - negation (NOT)
 - if `is.na(x)` is TRUE, then `!is.na(x)` is FALSE
- `any` will be TRUE if ANY are true
 - `any(is.na(x))` - do we have any NA's in x?

```
A = c(1, 2, 4, NA)
```

```
B = c(1, 2, 3, 4)
```

```
any(is.na(A)) # are there any NAs - YES/TRUE
```

```
[1] TRUE
```

```
any(is.na(B)) # are there any NAs- NO/FALSE
```

```
[1] FALSE
```

naniar

Sometimes you need to look at lots of data though... the [naniar package](#) is a good option.

The `pct_complete()` function shows the percentage that is complete for a given data object.

```
#install.packages("naniar")  
library(naniar)  
x = c(0, NA, 2, 3, 4, -0.5, 0.2)  
naniar::pct_complete(x)
```

```
[1] 85.71429
```

Air quality data

The `airquality` dataset comes with R about air quality in New York in 1973.

`?airquality` # use this to find out more about the data

```
airqual <- tibble(airquality)
```

```
airqual
```

```
# A tibble: 153 × 6
```

	Ozone	Solar.R	Wind	Temp	Month	Day
	<int>	<int>	<dbl>	<int>	<int>	<int>
1	41	190	7.4	67	5	1
2	36	118	8	72	5	2
3	12	149	12.6	74	5	3
4	18	313	11.5	62	5	4
5	NA	NA	14.3	56	5	5
6	28	NA	14.9	66	5	6
7	23	299	8.6	65	5	7
8	19	99	13.8	59	5	8
9	8	19	20.1	61	5	9
10	NA	194	8.6	69	5	10

```
# ... with 143 more rows
```


nanian: `pct_complete()`

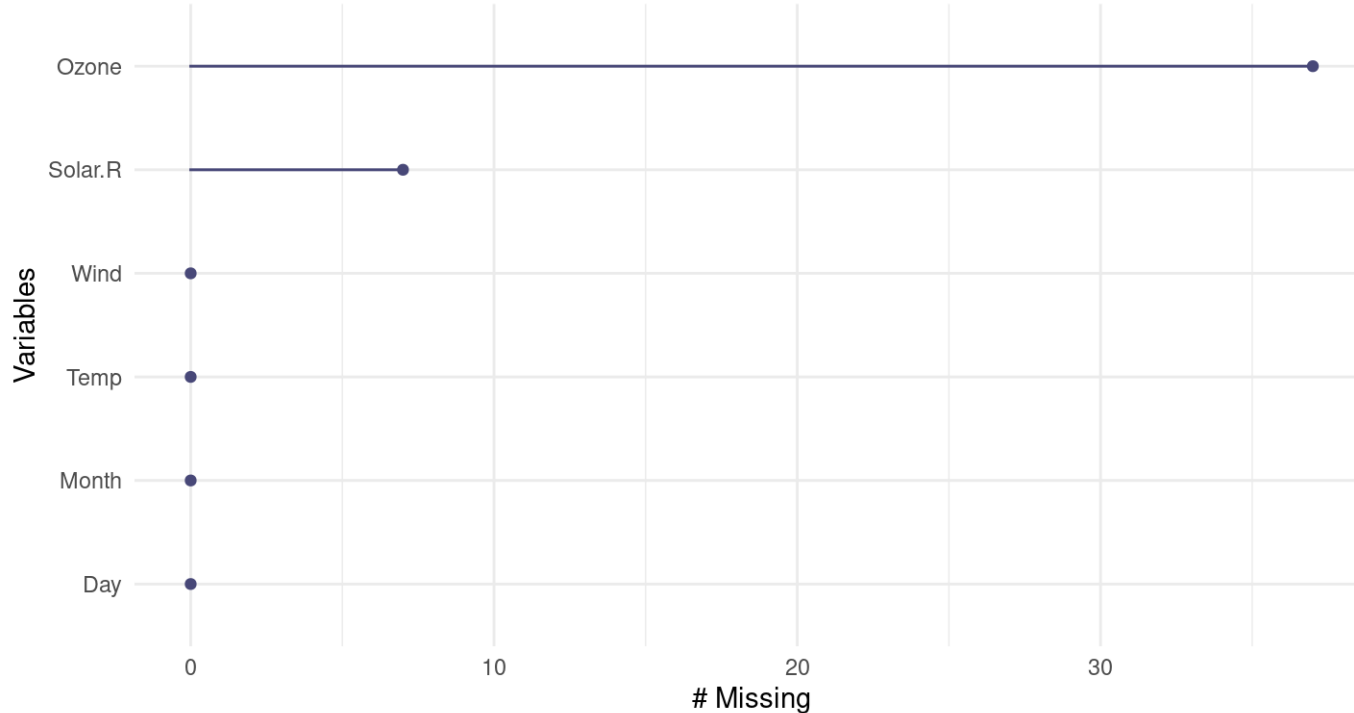
```
pct_complete(airquality)
```

```
[1] 95.20697
```

naniar plots

The `gg_miss_var()` function creates a nice plot about the number of missing values for each variable.

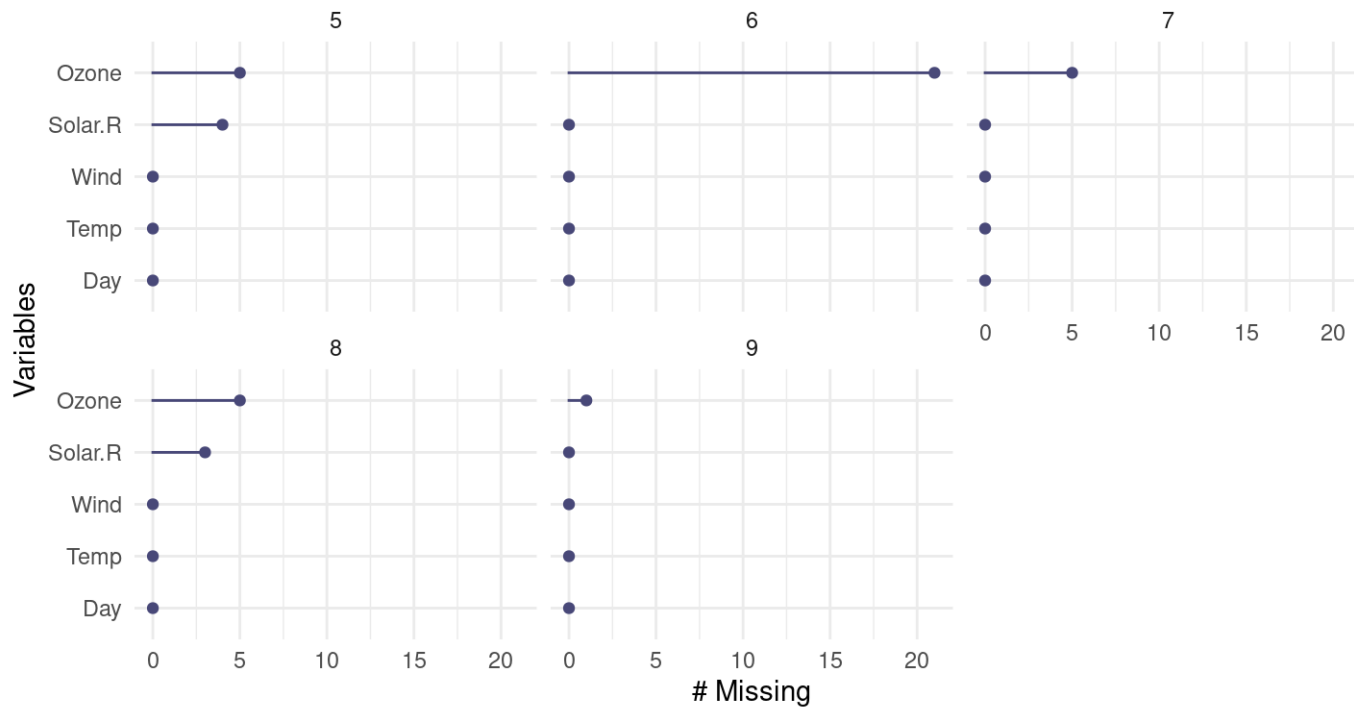
```
naniar::gg_miss_var(airqual)
```



naniar plots

We can use the `facet` argument to make more plots about a specific variable.

```
naniar::gg_miss_var(airqual, facet = Month)
```



Missing Data Issues

Recall that mathematical operations with **NA** often result in **NAs**.

```
sum(c(1, 2, 3, NA))
```

```
[1] NA
```

```
mean(c(2, 4, NA))
```

```
[1] NA
```

```
median(c(1, 2, 3, NA))
```

```
[1] NA
```

Missing Data Issues

This is also true for logicals. This is a good thing. The NA data could be > 2 or not, we don't know, so R says there is no **TRUE** or **FALSE**, so that is missing.

```
x = c(0, NA, 2, 3, 4, -0.5, 0.2)
```

```
x > 2
```

```
[1] FALSE    NA FALSE  TRUE  TRUE FALSE FALSE
```

filter() and missing data

Be careful with missing data using subsetting:

`filter()` removes missing values by default. To keep them need to add `is.na()`:

```
x # looks like the 1st and 3rd element should be TRUE
```

```
[1] 0.0 NA 2.0 3.0 4.0 -0.5 0.2
```

```
x %in% c(0, 2) # uh oh - not good!
```

```
[1] TRUE FALSE TRUE FALSE FALSE FALSE FALSE
```

```
x %in% c(0, 2) | is.na(x) # do this
```

```
[1] TRUE TRUE TRUE FALSE FALSE FALSE FALSE
```

filter() and missing data

df

```
# A tibble: 6 × 2
```

	Dog	Cat
	<dbl>	<dbl>
1	0	NA
2	NA	8
3	2	6
4	3	NA
5	1	2
6	1	NA

```
df %>% filter(Dog < 3)
```

```
# A tibble: 4 × 2
```

	Dog	Cat
	<dbl>	<dbl>
1	0	NA
2	2	6
3	1	2
4	1	NA

to remove rows with NAs for one variable use **drop_na()**

Avoid using filter for NA values. Instead use drop_na()

```
df %>% drop_na(Dog)
```

```
# A tibble: 5 × 2
```

	Dog	Cat
	<dbl>	<dbl>
1	0	NA
2	2	6
3	3	NA
4	1	2
5	1	NA

!NA does not work as you might expect because you can't tell if something is not actually NA- R doesn't ever assume to know what the value of NA` is

```
NA == NA
```

```
[1] NA
```


tidyr::drop_na

This function will drop rows with **any** missing data in **any** column when used on a df.

df

```
# A tibble: 6 × 2
```

	Dog	Cat
	<dbl>	<dbl>
1	0	NA
2	NA	8
3	2	6
4	3	NA
5	1	2
6	1	NA

```
drop_na(df)
```

```
# A tibble: 2 × 2
```

	Dog	Cat
	<dbl>	<dbl>
1	2	6
2	1	2

Drop columns with any missing values

```
df<-df %>% mutate(test =c(1,2,3,4,5,6))  
miss_var_which(df)
```

```
[1] "Dog" "Cat"
```

```
df %>% select(!miss_var_which(df))
```

```
# A tibble: 6 × 1
```

	test
	<dbl>
1	1
2	2
3	3
4	4
5	5
6	6

Removing columns with threshold of percent missing row values

```
is.na(df)
```

```
      Dog   Cat test
[1,] FALSE  TRUE FALSE
[2,]  TRUE FALSE FALSE
[3,] FALSE FALSE FALSE
[4,] FALSE  TRUE FALSE
[5,] FALSE FALSE FALSE
[6,] FALSE  TRUE FALSE
```

```
colMeans(is.na(df))
```

```
      Dog      Cat      test
0.1666667 0.5000000 0.0000000
```

```
df %>% select(which(colMeans(is.na(df)) < 0.2))
```

```
# A tibble: 6 × 2
```

```
  Dog test
<dbl> <dbl>
```

Change a value to be **NA**

The `na_if()` function of `dplyr` can be helpful for this. Let's say we think that all 0 values should be **NA**.

`na_if(vector to change, value to replace with NA)`

```
df %>% select(Dog) %>% na_if(0)
```

```
# A tibble: 6 × 1
```

```
  Dog  
  <dbl>  
1   NA  
2   NA  
3    2  
4    3  
5    1  
6    1
```

```
df %>% mutate(Dog = na_if(Dog, 0))
```

```
# A tibble: 6 × 3
```

```
  Dog   Cat test  
  <dbl> <dbl> <dbl>
```

Think about **NA**

Sometimes removing **NA** values leads to distorted math - be careful! Think about what your **NA** means for your data (are you sure ?).

Is an **NA** for values so low they could not be reported? Or is it this and also if there was a different issue?

Think about **NA**

If it is something more like a zero then you might want it included in your data like a zero.

Example: - survey reports NA if student has never tried cigarettes - survey reports 0 if student has tried cigarettes but did not smoke that week

You might want to keep the NA values so that you know the original sample size.

Word of caution

Calculating percentages will give you a different result depending on your choice to include NA values.

```
red_blue
```

```
# A tibble: 3 × 2
  color col_count
  <chr>     <int>
1 blue         3
2 red          3
3 <NA>         3
```

```
red_blue %>% mutate(percent =
  col_count/sum(pull(red_blue, col_count)))
```

```
# A tibble: 3 × 3
  color col_count percent
  <chr>     <int>     <dbl>
1 blue         3    0.333
2 red          3    0.333
3 <NA>         3    0.333
```

Word of caution

```
red_blue %>% mutate(percent =  
  col_count/sum(pull(drop_na(red_blue), col_count)))
```

```
# A tibble: 3 × 3
```

	color	col_count	percent
	<chr>	<int>	<dbl>
1	blue	3	0.5
2	red	3	0.5
3	<NA>	3	0.5

Should you be dividing by 9 or 6? It depends on your data

Pay attention to your data and your NAs!

Check values

Check the values for your variables, are they what you expect?

`count()` is a great option because it gives tells you:

1. The unique values
2. the amount of these values

Check if rare values make sense

```
bike <- jhur::read_bike()
```

```
bike %>% count(subType)
```

```
# A tibble: 4 × 2
```

	subType	n
	<chr>	<int>
1	STCLN	1
2	STRALY	3
3	STRPRD	1623
4	<NA>	4

Lab Part 1

[lab part 1](#)

[Website](#)

Recoding Variables

Example of Recoding

Say we have some data about samples in a diet study:

```
data_diet
```

```
# A tibble: 12 × 4
```

	Diet	Gender	Weight_start	Weight_change
	<chr>	<chr>	<int>	<int>
1	A	Male	179	4
2	B	m	249	3
3	B	Other	200	9
4	A	F	210	19
5	B	Female	236	13
6	B	M	232	-7
7	A	f	243	15
8	B	O	231	10
9	B	Man	197	-3
10	A	f	202	18
11	B	F	189	-8
12	B	O	169	14

Oh dear...

This needs lots of recoding.

```
data_diet %>%  
  count(Gender, Diet)
```

```
# A tibble: 10 × 3
```

	Gender	Diet	n
	<chr>	<chr>	<int>
1	f	A	2
2	F	A	1
3	F	B	1
4	Female	B	1
5	m	B	1
6	M	B	1
7	Male	A	1
8	Man	B	1
9	0	B	2
10	Other	B	1

dp1yr can help!

Using Excel to find all of the different ways **gender** has been coded, would be a matter of filtering and changing all by hand or using if statements. This can be hectic!

In **dp1yr** you can use the **recode** function (need **mutate** here too!):

General Format - this is not code!

```
{data_input} %>%  
  mutate({variable_to_fix} = {Variable_fixing, {old_value} = {new_value},  
                                         {another_old_value} = {new_value})
```

```
data_diet %>%  
  mutate(Gender = recode(Gender, M = "Male",  
                           m = "Male",  
                           Man = "Male",  
                           O = "Other",  
                           f = "Female",  
                           F = "Female")) %>%  
  count(Gender, Diet)
```

Or you can use **case_when()**.

The `case_when()` function of `dplyr` can help us to do this as well.

General Format - this is not code!

```
{data_input} %>%  
  mutate({variable_to_fix} = case_when{Variable_fixing}condition  
                                             ~ {value_for_cond}))
```

Note that automatically values not reassigned explicitly by `case_when` will be **NA**.

Use of `case_when()`

```
data_diet %>%  
  mutate(Gender = case_when(Gender == "M" ~ "Male"))
```

```
# A tibble: 12 × 4
```

	Diet	Gender	Weight_start	Weight_change
	<chr>	<chr>	<int>	<int>
1	A	<NA>	179	4
2	B	<NA>	249	3
3	B	<NA>	200	9
4	A	<NA>	210	19
5	B	<NA>	236	13
6	B	Male	232	-7
7	A	<NA>	243	15
8	B	<NA>	231	10
9	B	<NA>	197	-3
10	A	<NA>	202	18
11	B	<NA>	189	-8
12	B	<NA>	169	14

More complicated case_when()

```
data_diet %>%  
  mutate(Gender = case_when(  
    Gender %in% c("M", "male", "Man", "m", "Male") ~ "Male",  
    Gender %in% c("F", "Female", "f", "female") ~ "Female",  
    Gender %in% c("O", "Other") ~ "Other"))
```

```
# A tibble: 12 × 4
```

	Diet	Gender	Weight_start	Weight_change
	<chr>	<chr>	<int>	<int>
1	A	Male	179	4
2	B	Male	249	3
3	B	Other	200	9
4	A	Female	210	19
5	B	Female	236	13
6	B	Male	232	-7
7	A	Female	243	15
8	B	Other	231	10
9	B	Male	197	-3
10	A	Female	202	18
11	B	Female	189	-8
12	B	Other	169	14

Another reason for `case_when()`

`case_when` can do very sophisticated comparisons

```
data_diet <- data_diet %>%  
  mutate(Effect = case_when(Weight_change > 0 ~ "Increase",  
                             Weight_change == 0 ~ "Same",  
                             Weight_change < 0 ~ "Decrease"))
```

```
head(data_diet)
```

```
# A tibble: 6 × 5
```

	Diet	Gender	Weight_start	Weight_change	Effect
	<chr>	<chr>	<int>	<int>	<chr>
1	A	Male	179	4	Increase
2	B	m	249	3	Increase
3	B	Other	200	9	Increase
4	A	F	210	19	Increase
5	B	Female	236	13	Increase
6	B	M	232	-7	Decrease

```
# A tibble: 3 × 3
```

	Diet	Effect	n
	<chr>	<chr>	<int>
1	A	Increase	4

What if our data looked like this?

```
diet_comb
```

```
# A tibble: 3 × 2  
  change      n  
  <chr>    <int>  
1 A_Increase    4  
2 B_Decrease    3  
3 B_Increase    5
```

Separating columns based on a separator

- From `tidyr`, you can split a data set into multiple columns:

```
diet_comb %>%  
  separate(change, into = c("Diet", "Change"))
```

```
# A tibble: 3 × 3  
  Diet Change      n  
  <chr> <chr>   <int>  
1 A     Increase  4  
2 B     Decrease  3  
3 B     Increase  5
```

Separating columns based on a separator

You can specify the separator with `sep`.

```
diet_comb %>%  
  separate(change, into = c("Diet", "Change"), sep = " ")
```

```
# A tibble: 3 × 3  
  Diet    Change      n  
  <chr>  <chr>    <int>  
1 A_diet Increase     4  
2 B_diet Decrease     3  
3 B_diet Increase     5
```

Uniting columns based on a separator

- From `tidyr`, you can unite:

```
df = tibble(id = rep(1:5, 3), visit = rep(1:3, each = 5))  
head(df, 4)
```

```
# A tibble: 4 × 2
```

	id	visit
	<int>	<int>
1	1	1
2	2	1
3	3	1
4	4	1

```
df_united <- df %>% unite(col = "unique_id", id, visit, sep = "_")  
head(df_united, 4)
```

```
# A tibble: 4 × 1
```

	unique_id
	<chr>
1	1_1
2	2_1
3	3_1
4	4_1

Strings functions

Splitting/Find/Replace and Regular Expressions

- R can do much more than find exact matches for a whole string!

The `stringr` package

The `stringr` package:

- Modifying or finding **part** or all of a character string
- We will not cover `grep` or `gsub` - base R functions
 - are used on forums for answers
- Almost all functions start with `str_*`

stringr

`str_detect`, and `str_replace` search for matches to argument pattern within each element of a character vector (not data frame or tibble!).

- `str_detect` - returns `TRUE` if pattern is found
- `str_replace` - replaces pattern with replacement

Download Salary FY2014 Data

From <https://data.baltimorecity.gov/City-Government/Baltimore-City-Employee-Salaries-FY2015/nsfe-bg53>, from <https://data.baltimorecity.gov/api/views/nsfe-bg53/rows.csv>

Read the CSV into R **Sal**:

```
Sal = jhur::read_salaries() # or
```

```
head(Sal)
```

```
# A tibble: 6 × 7
```

	name	JobTitle	AgencyID	Agency	HireDate	AnnualSalary	GrossPay
	<chr>	<chr>	<chr>	<chr>	<chr>	<chr>	<chr>
1	Aaron, Patricia G	Facilitie...	A03031	OED-E...	10/24/1...	\$55314.00	\$53626....
2	Aaron, Petra L	ASSISTANT...	A29045	State...	09/25/2...	\$74000.00	\$73000....
3	Abaine, Yohannes T	EPIDEMIOLOG...	A65026	HLTH-...	07/23/2...	\$64500.00	\$64403....
4	Abbene, Anthony M	POLICE OF...	A99005	Polic...	07/24/2...	\$46309.00	\$59620....
5	Abbey, Emmanuel	CONTRACT ...	A40001	M-R I...	05/01/2...	\$60060.00	\$54059....
6	Abbott-Cole, Michelle	CONTRACT ...	A90005	TRANS...	11/28/2...	\$42702.00	\$20250....

'Find'str_detect() function: finding values: stringr

```
Sal %>% filter(str_detect(name, "Rawlings"))
```

```
# A tibble: 3 × 7
```

	name	JobTitle	AgencyID	Agency	HireDate	AnnualSalary	GrossPay
	<chr>	<chr>	<chr>	<chr>	<chr>	<chr>	<chr>
1	Rawlings,Kellye A	EMERGEN...	A40302	M-R I...	01/06/2...	\$48940.00	\$73356....
2	Rawlings,Paula M	COMMUNI...	A04015	R&P-R...	12/10/2...	\$19802.00	\$10443....
3	Rawlings-Blake,Stepha...	MAYOR	A01001	Mayor...	12/07/1...	\$167449.00	\$165249...

Showing difference in `str_replace` and `str_replace_all`

`str_replace` replaces only the first instance.

```
head(pull(Sal, JobTitle))
```

```
[1] "Facilities/Office Services II" "ASSISTANT STATE'S ATTORNEY"  
[3] "EPIDEMIOLOGIST"              "POLICE OFFICER"  
[5] "CONTRACT SERV SPEC II"       "CONTRACT SERV SPEC II"
```

```
head(str_replace(pull(Sal, JobTitle), "II", "2"))
```

```
[1] "Facilities/Office Services 2" "ASSISTANT STATE'S ATTORNEY"  
[3] "EPIDEMIOLOGIST"              "POLICE OFFICER"  
[5] "CONTRACT SERV SPEC 2"        "CONTRACT SERV SPEC 2"
```

`str_replace` replaces all instances.

```
head(str_replace_all(pull(Sal, name), "a", "j"), 2)
```

```
[1] "Ajron,Pjtricij G" "Ajron,Petrj L"
```

Lab Part 2

[lab part 2](#)

[Website](#)

Extra Slides

String Splitting

- `str_split(string, pattern)` - splits strings up - returns list!

```
library(stringr)
x <- c("I really like writing R code")
df = tibble(x = c("I really", "like writing", "R code programs"))
y <- unlist(str_split(x, " "))
y
```

```
[1] "I"          "really"    "like"     "writing"  "R"         "code"
```

```
length(y)
```

```
[1] 6
```


A bit on Regular Expressions

- <http://www.regular-expressions.info/reference.html>
- They can use to match a large number of strings in one statement
- `.` matches any single character
- `*` means repeat as many (even if 0) more times the last character
- `?` makes the last thing optional
- `^` matches start of vector `^a` - starts with "a"
- `$` matches end of vector `b$` - ends with "b"

Let's look at modifiers for `stringr`

`?modifiers`

- `fixed` - match everything exactly
- `ignore_case` is an option to not have to use `tolower`

Using a fixed expression

One example case is when you want to split on a period “.”. In regular expressions `.` means **ANY** character, so we need to specify that we want R to interpret “.” as simply a period.

```
str_split("I.like.strings", ".")
```

```
[[1]]  
[1] "" "" "" "" "" "" "" "" "" "" "" "" "" "" ""
```

```
str_split("I.like.strings", fixed("."))
```

```
[[1]]  
[1] "I"      "like"   "strings"
```

```
str_split("I.like.strings", "\\.")
```

```
[[1]]  
[1] "I"      "like"   "strings"
```

Pasting strings with `paste` and `paste0`

Paste can be very useful for joining vectors together:

```
paste("Visit", 1:5, sep = "_")
```

```
[1] "Visit_1" "Visit_2" "Visit_3" "Visit_4" "Visit_5"
```

```
paste("Visit", 1:5, sep = "_", collapse = "_")
```

```
[1] "Visit_1_Visit_2_Visit_3_Visit_4_Visit_5"
```

and paste0 can be even simpler see ?paste0

```
paste0("Visit", 1:5) # no space!
```

```
[1] "Visit1" "Visit2" "Visit3" "Visit4" "Visit5"
```

!- # Before Cleaning - Subsetting with Brackets ->

->

-> -> ->

Using Regular Expressions

- Look for any name that starts with:
 - Payne at the beginning,
 - Leonard and then an S
 - Spence then capital C

```
head(str_subset( Sal$name, "^Payne.*"), 3)
```

```
[1] "Payne El,Boaz L"      "Payne El,Jackie"  
[3] "Payne Johnson,Nickole A"
```

```
head(str_subset( Sal$name, "Leonard.?S"))
```

```
[1] "Payne,Leonard S"      "Szumlanski,Leonard S"
```

```
head(str_subset( Sal$name, "Spence.*C.*"))
```

```
[1] "Spencer,Charles A"  "Spencer,Clarence W" "Spencer,Michael C"
```

Comparison of **stringr** to base R -
not covered

Splitting Strings

Substringing

stringr

- `str_split(string, pattern)` - splits strings up - returns list!

Splitting String:

In `stringr`, `str_split` splits a vector on a string into a `list`

```
x <- c("I really", "like writing", "R code programs")  
y <- stringr::str_split(x, pattern = " ") # returns a list  
y
```

```
[[1]]  
[1] "I"      "really"
```

```
[[2]]  
[1] "like"   "writing"
```

```
[[3]]  
[1] "R"      "code"   "programs"
```

str_extract

str_extract extracts matched strings - `\\d` searches for DIGITS/numbers

```
head(Sal$AgencyID)
```

```
[1] "A03031" "A29045" "A65026" "A99005" "A40001" "A90005"
```

```
head(str_extract(Sal$AgencyID, "\\d"))
```

```
[1] "0" "2" "6" "9" "4" "9"
```

'Find' functions: stringr compared to base R

Base R does not use these functions. Here is a "translator" of the `stringr` function to base R functions

- `str_detect` - similar to `grep1` (return logical)
- `grep(value = FALSE)` is similar to `which(str_detect())`
- `str_subset` - similar to `grep(value = TRUE)` - return value of matched
- `str_replace` - similar to `sub` - replace one time
- `str_replace_all` - similar to `gsub` - replace many times

Important Comparisons

Base R:

- Argument order is (pattern, x)
- Uses option (fixed = TRUE)

stringr

- Argument order is (string, pattern) aka (x, pattern)
- Uses function fixed(pattern)

'Find' functions: Finding Indices

These are the indices where the pattern match occurs:

```
grep("Rawlings", Sal$Name)
```

```
Warning: Unknown or uninitialised column: `Name`.
```

```
integer(0)
```

```
which(grepl("Rawlings", Sal$Name))
```

```
Warning: Unknown or uninitialised column: `Name`.
```

```
integer(0)
```

```
which(str_detect(Sal$Name, "Rawlings"))
```

```
Warning: Unknown or uninitialised column: `Name`.
```

```
integer(0)
```

'Find' functions: Finding Logicals

These are the indices where the pattern match occurs:

```
head(grepl("Rawlings", Sal$Name))
```

```
Warning: Unknown or uninitialised column: `Name`.
```

```
logical(0)
```

```
head(str_detect(Sal$Name, "Rawlings"))
```

```
Warning: Unknown or uninitialised column: `Name`.
```

```
logical(0)
```

'Find' functions: finding values, base R

```
grep("Rawlings", Sal$Name, value=TRUE)
```

Warning: Unknown or uninitialised column: `Name`.

```
character(0)
```

```
Sal[grep("Rawlings", Sal$Name),]
```

Warning: Unknown or uninitialised column: `Name`.

```
# A tibble: 0 × 7
```

```
# ... with 7 variables: name <chr>, JobTitle <chr>, AgencyID <chr>, Agency <chr>,
```

```
#   HireDate <chr>, AnnualSalary <chr>, GrossPay <chr>
```

Showing difference in `str_extract`

`str_extract` extracts just the matched string

```
ss = str_extract(Sal$Name, "Rawling")
```

```
Warning: Unknown or uninitialised column: `Name`.
```

```
head(ss)
```

```
character(0)
```

```
ss[ !is.na(ss)]
```

```
character(0)
```


Showing difference in `str_extract` and `str_extract_all`

`str_extract_all` extracts all the matched strings

```
head(str_extract(Sal$AgencyID, "\\d"))
```

```
[1] "0" "2" "6" "9" "4" "9"
```

```
head(str_extract_all(Sal$AgencyID, "\\d"), 2)
```

```
[[1]]
```

```
[1] "0" "3" "0" "3" "1"
```

```
[[2]]
```

```
[1] "2" "9" "0" "4" "5"
```

Using Regular Expressions

- Look for any name that starts with:
 - Payne at the beginning,
 - Leonard and then an S
 - Spence then capital C

```
head(grep("^Payne.*", x = Sal$name, value = TRUE), 3)
```

```
[1] "Payne El,Boaz L"      "Payne El,Jackie"  
[3] "Payne Johnson,Nickole A"
```

```
head(grep("Leonard.?S", x = Sal$name, value = TRUE))
```

```
[1] "Payne,Leonard S"      "Szumlanski,Leonard S"
```

```
head(grep("Spence.*C.*", x = Sal$name, value = TRUE))
```

```
[1] "Spencer,Charles A"  "Spencer,Clarence W" "Spencer,Michael C"
```

Using Regular Expressions: `stringr`

```
head(str_subset( Sal$name, "^Payne.*"), 3)
```

```
[1] "Payne El,Boaz L"      "Payne El,Jackie"  
[3] "Payne Johnson,Nickole A"
```

```
head(str_subset( Sal$name, "Leonard.?S"))
```

```
[1] "Payne,Leonard S"      "Szumlanski,Leonard S"
```

```
head(str_subset( Sal$name, "Spence.*C.*"))
```

```
[1] "Spencer,Charles A"  "Spencer,Clarence W" "Spencer,Michael C"
```

Replace

Let's say we wanted to sort the data set by Annual Salary:

```
class(Sal$AnnualSalary)
```

```
[1] "character"
```

```
sort(c("1", "2", "10")) # not sort correctly (order simply ranks the data)
```

```
[1] "1" "10" "2"
```

```
order(c("1", "2", "10"))
```

```
[1] 1 3 2
```

Replace

So we must change the annual pay into a numeric:

```
head(Sal$AnnualSalary, 4)
```

```
[1] "$55314.00" "$74000.00" "$64500.00" "$46309.00"
```

```
head(as.numeric(Sal$AnnualSalary), 4)
```

```
Warning in head(as.numeric(Sal$AnnualSalary), 4): NAs introduced by coercion
```

```
[1] NA NA NA NA
```

R didn't like the \$ so it thought turned them all to NA.

`sub()` and `gsub()` can do the replacing part in base R.

Replacing and subbing

Now we can replace the \$ with nothing (used `fixed=TRUE` because \$ means ending):

```
Sal$AnnualSalary <- as.numeric(gsub(pattern = "$", replacement="",  
                                   Sal$AnnualSalary, fixed=TRUE))  
Sal <- Sal[order(Sal$AnnualSalary, decreasing=TRUE), ]  
Sal[1:5, c("name", "AnnualSalary", "JobTitle")]
```

A tibble: 5 × 3

	name <chr>	AnnualSalary <dbl>	JobTitle <chr>
1	Mosby, Marilyn J	238772	STATE'S ATTORNEY
2	Batts, Anthony W	211785	Police Commissioner
3	Wen, Leana	200000	Executive Director III
4	Raymond, Henry J	192500	Executive Director III
5	Swift, Michael	187200	CONTRACT SERV SPEC II

Replacing and subbing: `stringr`

We can do the same thing (with 2 piping operations!) in dplyr

```
dplyr_sal = Sal
dplyr_sal = dplyr_sal %>% mutate(
  AnnualSalary = AnnualSalary %>%
    str_replace(
      fixed("$"),
      "" ) %>%
    as.numeric) %>%
  arrange(desc(AnnualSalary))
check_Sal = Sal
rownames(check_Sal) = NULL
all.equal(check_Sal, dplyr_sal)
```

```
[1] TRUE
```

Website

[Website](#)

Extra slides

Creating Two-way Tables

A two-way table. If you pass in 2 vectors, `table` creates a 2-dimensional table.

```
tab <- table(c(0, 1, 2, 3, 2, 3, 3, 2, 2, 3),  
             c(0, 1, 2, 3, 2, 3, 3, 4, 4, 3),  
             useNA = "always")
```

tab

	0	1	2	3	4	<NA>
0	1	0	0	0	0	0
1	0	1	0	0	0	0
2	0	0	2	0	2	0
3	0	0	0	4	0	0
<NA>	0	0	0	0	0	0

Creating Two-way Tables

```
tab_df = tibble(x = c(0, 1, 2, 3, 2, 3, 3, 2, 2, 3),  
                 y = c(0, 1, 2, 3, 2, 3, 3, 4, 4, 3))  
tab_df %>% count(x, y)
```

```
# A tibble: 5 × 3
```

	x	y	n
	<dbl>	<dbl>	<int>
1	0	0	1
2	1	1	1
3	2	2	2
4	2	4	2
5	3	3	4

Creating Two-way Tables

```
tab_df %>%  
  count(x, y) %>%  
  group_by(x) %>% mutate(pct_x = n / sum(n))
```

```
# A tibble: 5 × 4
```

```
# Groups:   x [4]
```

	x	y	n	pct_x
	<dbl>	<dbl>	<int>	<dbl>
1	0	0	1	1
2	1	1	1	1
3	2	2	2	0.5
4	2	4	2	0.5
5	3	3	4	1

Creating Two-way Tables

```
library(scales)
tab_df %>%
  count(x, y) %>%
  group_by(x) %>% mutate(pct_x = percent(n / sum(n)))
```

```
# A tibble: 5 × 4
```

```
# Groups:   x [4]
```

	x	y	n	pct_x
	<dbl>	<dbl>	<int>	<chr>
1	0	0	1	100%
2	1	1	1	100%
3	2	2	2	50%
4	2	4	2	50%
5	3	3	4	100%