

# r\_int\_day\_3\_data\_transformation

Nicholus Tint Zaw

2022-11-10

## Content

This lecture focus on the implementation of data wrangling using the **dplyer** package and will cover some exercise using the popular functions from **dplyer** package.

1. filter
2. arrange
3. select
4. mutate
5. summarise
6. group\_by

## Some useful commands for exploring dataset

```
# load dataset

df <- polls_us_election_2016 # from dslabs package

head(df) # print first 6 obs
tail(df) # print last 6 obs

names(df) # variable/column names

unique(df$grade) # inspect unique values in a specific variable/column

typeof(df$grade) # inspect the type of variable
class(df$grade) # inspect the type of variable

table(df$grade) # frequency table
```

## filter : select the sub-set of the dataset

```
filter(df, grade == "D")
filter(df, state == "Ohio")

# filter with multiple conditions
```

```
interested_grades <- c("D", "B")

filter(df, grade %in% interested_grades) # one line

filter(df, grade == "D" | grade == "B")
```

Using filter with pipe (%>%) function.

```
df %>%
  filter(state == "Illinois")
```

**arrange** : order the dataset by given variable

```
head(arrange(df, samplesize)) # ascending order - default

head(arrange(df, desc(population))) # descending order

head(arrange(df, population))
head(arrange(df, desc(population)))
```

use with pipe function,

```
df %>% arrange(population)

df %>% arrange(desc(population))
```

**select**: keep only variables require for data processing

```
head(select(df, state, samplesize, population))

head(select(df, -(c(state, population, samplesize))))
```

```
# starts_with("")

df %>%
  select(starts_with("adj"))

# ends_with("")
df %>%
  select(ends_with("date"))

# contains("")
df %>%
  select(contains("po"))

# matches("(.)\\1") - regular expression
```

```
df %>%
  select(matches("raw")) # check the result.

# what is the different between matches and contains?

df1 <- data.frame(colnm = 1:5, col1 = 24, col2 = 46)
df1 %>%
  select(contains("col"))

df1 %>%
  select(matches("col\\d+"))

# num_range("x", 1:3)
# pls check at help file - type ?dplyr::select in console.
```

Wanna study more about regular expression, check [here](#).

## mutate: adding new column to existing one

```
df %>%
  mutate(ss_small = ifelse(samplesize < 1000, 1, 0)) %>%
  select(samplesize, ss_small)

# if want to add into existing dataframe, override the existing dataframe with resulted dataframe.
df <- df %>%
  mutate(ss_small = ifelse(samplesize < 1000, 1, 0)) # check the variable names and numbers
```

## summarise: perform sumstat functions

```
summarise(df, ss_mean = mean(samplesize)) # what happened!

summarise(df, ss_mean = mean(samplesize, na.rm = TRUE)) # Missing values are contagious

# Missing values are contagious
NA + 1

10 * 100 * NA

100/NA

# sumstat table for samplesize var
df %>%
  summarise(man = mean(samplesize, na.rm = TRUE),
            sd = sd(samplesize, na.rm = TRUE),
            median = median(samplesize, na.rm = TRUE),
            min = min(samplesize, na.rm = TRUE),
            max = max(samplesize, na.rm = TRUE))
```

**group\_by:** manipulation at group level, sub-group level

```
# get the sample size mean value per state
df %>%
  group_by(state) %>%
  summarise(mean = mean(samplesize, na.rm = TRUE))

df %>%
  group_by(population) %>%
  summarise(mean = mean(samplesize, na.rm = TRUE))

df %>%
  group_by(state) %>%
  count()
```

## Exercise

- Use iris dataset and calculate sumstat by Species.
- Select the only observation from versicolor and virginica species.