#### QTM 151

Week 9 – forcats

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#### Recap

#### We learned:

- qplot: quick way to make ggplot graphs.
- ggplotly: transform ggplot objects into nice plotly viz.
- plot\_ly: create nice plotly graphs.
- dplyr methods: data wrangling
- dplyr \*\_join methods: joining data
- tidyr methods: reshape datasets

#### **Great job!!**

Do you have any questions about any of these contents?

Today we are going to talk about **forcats** (package for categorical variables)

#### This week

We will have a **quiz** posted today after 4:00 PM. Due by **Tuesday** (because of the holidays this week).

We will have a **problem set** posted tomorrow, due by the next lab.

Thank you for your answers to the matching survey. I will match you later today.

Thank you ver much for your answers to the midterm evaluations survey. I will send you the overall results in an announcement later tomorrow.

There are a few changes that I can do to improve the class. Thanks to your answers, I have a clear path to improve things.

Our GitHub page is: https://github.com/umbertomig/qtm151

# **Getting Started**

## Getting Started: loading packages

```
# Loading tidyverse
library(tidyverse)
## — Attaching packages -
                                                            tidyv
## / ggplot2 3.3.3 / purrr 0.3.4
## / tibble 3.1.0 / dplyr 1.0.5
## / tidyr 1.1.3 / stringr 1.4.0
## / readr 1.4.0
                     ✓ forcats 0.5.0
## — Conflicts
                                                       tidyverse o
## x dplyr::filter() masks stats::filter()
## x dplyr::lag() masks stats::lag()
```

### Loading data - GSS

```
# Loading the GSS Cat and GSS
 gss cat ← read csv('https://raw.githubusercontent.com/umbertomig/
##
## — Column specification
## cols(
  year = col double(),
###
     marital = col character(),
###
     age = col double(),
###
###
     race = col character(),
     rincome = col character(),
###
     partyid = col character(),
###
     relig = col character(),
###
     denom = col character(),
###
###
     tvhours = col double()
## )
```

### forcats

#### forcats

Provide tools to work with categorical data. The methods we will use in here are:

- fct\_reorder(): reorder levels in a categorical variable
- fct\_relevel(): move the position of a particular level
- fct\_infreq(): reorder levels in increasing frequency
- fct\_recode(): recode factor variable
- fct\_collapse(): collapse levels in categorical variable
- fct\_lump(): lump together small groups
- case\_when(): useful to recode data

## fct\_reorder

#### fct\_reorder

Suppose we want to study the relationship between religion and time watching tv.

This graph here doesn't help much:

```
relig_summary \( \) gss_cat %>%
  group_by(relig) %>%
  summarise(tvhours=mean(tvhours, na.rm = T))

ggplot(relig_summary, aes(x=tvhours, y=relig)) +
  geom_point()
```

#### fct\_reorder

Reordering the levels of a factor using *fct\_reorder()* helps with the interpretation.

fct\_reorder() takes three arguments:

- f, the factor whose levels you want to modify.
- x, a numeric vector that you want to use to reorder the levels.
- Optionally, fun, a function used if there are multiple values of x for each value of f. The default value is median.

```
ggplot(relig_summary, aes(x=tvhours, y=fct_reorder(relig, tvhours))) =
geom_point()
```

**Your turn:** plot the marital status by frequency (use the function counts). Then reorder it.

## fct\_relevel

### fct\_relevel

In the religions x tv hours plot, note that the "Don't know" is the largest category.

It is also not very informative. We can easily send it to the bottom:

### fct\_relevel

Can we do ascending, and place the "Don't know" in the bottom?

Yes!

**Your turn:** plot the party id by frequency (use the function counts). Then reorder it. Then place "No answer" and "Independent" at the bottom.

fct\_infreq

### fct\_infreq

You can use fct\_infreq() to order levels in increasing frequency:.

• This is the simplest type of reordering because it does not need any extra variables.

You may want to combine with fct\_rev().

```
gss_cat %>%
  mutate(marital = fct_infreq(marital)) %>%
  ggplot(aes(marital)) +
    geom_bar()
```

## fct\_infreq + fct\_rev (elegant)

And that's what happens when we combine both:

```
gss_cat %>%
mutate(marital = marital %>% fct_infreq() %>% fct_rev()) %>%
  ggplot(aes(marital)) + geom_bar()
```

**Your turn:** plot the race by frequency.

## fct\_recode

#### fct\_recode

Recode a categorical variable is always painful, regardless of the statistical software.

Luckly, the people that wrote forcats made it as easy as it can be by creating the fct\_recode function.

Check this plot. TV hours by party affiliation:

```
gss_cat %>%
  drop_na(tvhours) %>%
  group_by(partyid) %>%
  summarise(meantv=mean(tvhours)) %>%
  ggplot(aes(x=meantv, y=fct_reorder(partyid, meantv)))+
  geom_point()
```

#### fct\_recode

Note that we can combine some low-information categories together as others:

```
gss cat %>%
 drop na(tvhours) %>%
 mutate(partyidnew = fct_recode(partyid,
    "Republican, strong" = "Strong republican",
    "Republican, weak" = "Not str republican",
    "Independent, near rep" = "Ind, near rep",
    "Independent, near dem" = "Ind, near dem",
    "Democrat, weak" = "Not str democrat",
    "Democrat, strong"
                           = "Strong democrat",
    "Other"
                           = "No answer",
    "Other"
                           = "Don't know",
    "Other"
                           = "Other party")) %>%
 group by(partyidnew) %>%
  summarise(meantvhours = mean(tvhours)) %>%
 ggplot(aes(x=meantvhours,
```

## fct\_collapse

#### fct\_collapse

fct\_collapse is good to put several factor levels together.

Look at this code for party id frequency:

```
gss_cat %>%
  mutate(partyidnew = fct_collapse(partyid,
    other = c("No answer", "Don't know", "Other party"),
    rep = c("Strong republican", "Not str republican"),
    ind = c("Ind,near rep", "Independent", "Ind,near dem"),
    dem = c("Not str democrat", "Strong democrat")
    )) %>%
    count(partyidnew)
```

Your turn: collapse the gss wrkstat variable to simplify it.

# fct\_lump

### fct\_lump

fct\_lump aggregates the small-frequency levels together.

The default is the most frequent + others:

```
gss_cat %>%
  mutate(relignew = fct_lump(relig)) %>%
  count(relignew)
```

### fct\_lump

If we want the five most frequent values + others:

```
gss_cat %>%
  mutate(relig = fct_lump(relig, n=5)) %>%
  count(relig)
```

**Your turn:** count the levels in the gss\_cat denom variable. Find a nice way to display it.

# case\_when

#### case\_when

case\_when is useful for recoding variables. Look at the example below:

```
mtcars %>%
  mutate (
    gear_char =
    case_when(
        gear=3 ~ "three",
        gear=4 ~ "four",
        gear=5 ~ "five"
    )
)
```

**Your turn:** for gss\_cat, create a plot to look at how average age varies across income level (rincome)

## Questions?

## Have a great weekend!