

Data Wrangling with R: Day 2

Formatting factors with `forcats`

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There are two types of categorical variables

Nominal where there is no intrinsic ordering to the categories

E.g. blue, grey, black, white.

Ordinal where there is a clear order to the categories

E.g. Strongly disagree, disagree, neutral, agree, strongly agree.

Categorical variables in R

Part 1

- In R, categorical variables may be encoded in various ways.

```
cat_chr <- c("red", "white", "blue")
cat_fct <- factor(c("red", "white", "blue"))

class(cat_chr)

## [1] "character"

class(cat_fct)

## [1] "factor"
```

- Then you have categorical variables that look like a numerical variable (e.g. coded variables like say 1=male, 2=female)
- And also those that have fixed levels of numerical values (e.g. ToothGrowth\$dose: 0.5, 1.0 and 2.0)

So why encode as **factor** instead of **character**?

In some cases, characters are converted to factors (or vice-versa) in functions so they can be similar.

The main idea of a factor is that the variable has a *fixed number of levels*

Categorical variables in R

Part 2

- When a variable is encoded as a **factor** then there is an attribute with the levels

```
data <- c(2, 2, 1, 1, 3, 3, 3, 1)
```

```
factor(data)
```

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

```
## Levels: 1 2 3
```

- You can easily change the labels of the variables:

```
factor(data,  
       labels = c("I", "II", "III"))
```

```
## [1] II  II  I   I    III III III I
```

```
## Levels: I II III
```

Categorical variables in R

Part 3

- Order of the factors are determined by the input:

```
# numerical input are ordered in increasing order
```

```
factor(c(1, 3, 10))
```

```
## [1] 1 3 10
```

```
## Levels: 1 3 10
```

```
# character input are ordered alphabetically
```

```
factor(c("1", "3", "10"))
```

```
## [1] 1 3 10
```

```
## Levels: 1 10 3
```

```
# you can specify order of levels explicitly
```

```
factor(c("1", "3", "10"), levels = c("1", "3", "10"))
```

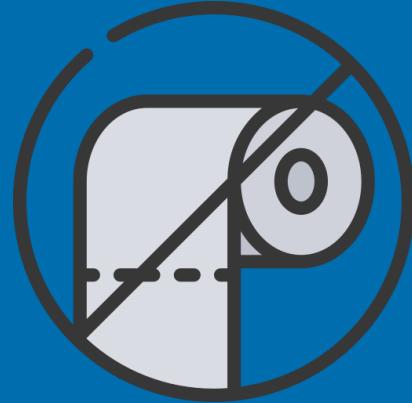
```
## [1] 1 3 10
```

Why would the order of the levels matter?

- Some downstream analysis may use it

```
data("population", package = "tidyverse")
population %>%
  filter(year == 2013) %>%
  # just choose 5 countries
  slice(c(1, 11, 21, 31, 41)) %>%
  ggplot(aes(population, country)) +
  geom_col()
```

```
population %>%
  filter(year == 2013) %>%
  slice(c(1, 11, 21, 31, 41)) %>%
  mutate(country =
    reorder(country, population)) %>%
  ggplot(aes(population, country)) +
  geom_col()
```



Cautionary tales of working with factors

Numerical factors in R

```
x <- factor(c(10, 20, 30, 10, 20))
mean(x)

## Warning in mean.default(x): argument is not numeric
## or logical: returning NA

## [1] NA
```

⚠ as.numeric function returns the internal integer values of the factor

```
mean(as.numeric(x))

## [1] 1.8
```

You probably want to use:

```
mean(as.numeric(levels(x)[x]))
## [1] 18
```

```
mean(as.numeric(as.character(x))) .
## [1] 18
```

Defining levels explicitly Part 1

- If the variable contain values that are not in the levels of the factors, then those values will become a missing value

```
factor(c("Yes", "No", "Maybe"), levels = c("Yes", "No"))

## [1] Yes  No   <NA>
## Levels: Yes No
```

- This can be useful at times, but it's a good idea to check the values before it is transformed as NA

```
factor(c("Yes", "No", "No", "Yess"), levels = c("Yes", "No"))

## [1] Yes  No   No   <NA>
## Levels: Yes No
```

Defining levels explicitly

Part 2

- You can have levels that are not observed

```
f <- factor(c("Yes", "Yes", "Yes", "No"), levels = c("Yes", "Maybe", "No"))
f

## [1] Yes Yes Yes No
## Levels: Yes Maybe No
```

- This can be useful at times downstream, e.g.

```
table(f)

## f
##   Yes Maybe   No
##     3      0      1
```

Combining factors as vectors

```
f1 <- factor(c("F", "M", "F"))
f2 <- factor(c("F", "F"))
```

- What do you think the output will be for below?

```
c(f1, f2)
```

```
## [1] 1 2 1 1 1
```

- Was that expected?
- The c function strips the class when you combine factors

```
unclass(f1)
```

```
## [1] 1 2 1
## attr(,"levels")
## [1] "F" "M"
```

Combining factors in a data frame

```
df1 <- data.frame(f = factor(c("a", "b")))
df2 <- data.frame(f = factor(c("c", "b")))
```

- What do you think the output below will be?

```
rbind(df1, df2)
```

```
##      f
## 1  a
## 2  b
## 3  c
## 4  b
```

```
rbind(df1, df2)$f
```

```
## [1] a b c b
## Levels: a b c
```

Working with factors with forcats

Formatting factors

- The `forcats` package is part of `tidyverse`
- Like the `stringr` package the main functions in `forcats` **prefix with `fct_` or `lvl_`** and the **first argument is a factor (or a character) vector**

some functions do not allow character as input, e.g. `fct_c`

- The list of available commands are:

- | | | | |
|--------------------------------|---------------------------------|-----------------------------|----------------------------|
| • <code>fct_anon</code> | • <code>fct_infreq</code> | • <code>fct_match</code> | • <code>fct_shift</code> |
| • <code>fct_c</code> | • <code>fct_inorder</code> | • <code>fct_other</code> | • <code>fct_shuffle</code> |
| • <code>fctCollapse</code> | • <code>fct_inseq</code> | • <code>fct_recode</code> | • <code>fct_unify</code> |
| • <code>fct_count</code> | • <code>fct_lump</code> | • <code>fct_relabel</code> | • <code>fct_unique</code> |
| • <code>fct_cross</code> | • <code>fct_lump_lowfreq</code> | • <code>fct_relevel</code> | • <code>lvl_expand</code> |
| • <code>fct_drop</code> | • <code>fct_lump_min</code> | • <code>fct_reorder</code> | • <code>lvl_reorder</code> |
| • <code>fct_expand</code> | • <code>fct_lump_n</code> | • <code>fct_reorder2</code> | • <code>lvl_revalue</code> |
| • <code>fct_explicit_na</code> | • <code>fct_lump_prop</code> | • <code>fct_rev</code> | • <code>lvl_union</code> |

Combining factors as vectors with `forcats`

```
f1 <- factor(c("F", "M", "F"))
f2 <- factor(c("F", "F"))

c(f1, f2)

## [1] 1 2 1 1 1

fct_c(f1, f2)

## [1] F M F F F
## Levels: F M

c1 <- c("F", "M", "F")

fct_c(c1, f2)

## Error: All elements of `...` must be factors
```

Count levels in a factor

```
data("gss_cat", package = "forcats")
table(gss_cat$race)

##
##          Other           Black          White
##          1959            3129          16395
## Not applicable
##          0
```

- `table` in Base R is useful but you may want the output as a data frame

```
fct_count(gss_cat$race, sort = TRUE, prop = TRUE)

## # A tibble: 4 x 3
##       f             n      p
##   <fct>     <int>  <dbl>
## 1 White      16395  0.763
```

Collapse levels in a factor

```
levels(gss_cat$marital)

## [1] "No answer"      "Never married"   "Separated"
## [4] "Divorced"       "Widowed"        "Married"

gss_cat$marital %>%
  fct_collapse(Single = c("Never married", "Separated", "Divorced")) %>%
  fct_relevel("No answer", after = Inf) %>% # move to last place
  fct_count()

## # A tibble: 4 x 2
##       f             n
##   <fct>     <int>
## 1 Single     9542
## 2 Widowed    1807
## 3 Married    10117
## 4 No answer   17
```

Lumping factor levels

Part 1

- Sometimes you have a lot of levels and you'd prefer to lump some of them together to the "Other" category
- What criterion do you use to lump levels together?
- There are four main criterion to lump levels using `fct_lump*` functions:
 - `fct_lump_n`: lump all levels except the `n` most frequent
 - `fct_lump_min`: lump together those less than `min` counts
 - `fct_lump_prop`: lump together those less than proportion of `prop`
 - `fct_lump_lowfreq`: lump up least frequent levels such that the Other level is still the smallest level
 - `fct_lump` lifecycle superseded, it is better to use one of the above functions instead

Lumping factor levels

Part 2

```
levels(gss_cat$relig)  
  
## [1] "No answer"  
## [2] "Don 't know"  
## [3] "Inter-nondenominational"  
## [4] "Native american"  
## [5] "Christian"  
## [6] "Orthodox-christian"  
## [7] "Moslem/islam"  
## [8] "Other eastern"  
## [9] "Hinduism"  
## [10] "Buddhism"  
## [11] "Other"  
## [12] "None"  
## [13] "Jewish"  
## [14] "Catholic"
```

```
fct_lump_n(gss_cat$relig, n = 2) %>%  
  fct_count(sort = TRUE, prop = TRUE)  
  
## # A tibble: 3 x 3  
##   f           n      p  
##   <fct>     <int> <dbl>  
## 1 Protestant 10846 0.505  
## 2 Other      5513  0.257  
## 3 Catholic   5124  0.239  
  
fct_lump_lowfreq(gss_cat$relig) %>%  
  fct_count(sort = TRUE, prop = TRUE)  
  
## # A tibble: 2 x 3  
##   f           n      p  
##   <fct>     <int> <dbl>  
## 1 Protestant 10846 0.505
```

**</> If you installed the `dwexercise` package,
run below in your R console**

```
learnr::run_tutorial("day2-exercise-02", package = "dwexercise")
```

 **If the above doesn't work for you, go [here](#).**
 **Questions or issues, let us know!**

15:00

Session Information

```
devtools::session_info()

## - Session info -----
##   setting  value
##   version  R version 4.0.1 (2020-06-06)
##   os        macOS Catalina 10.15.7
##   system   x86_64, darwin17.0
##   ui        RStudio
##   language (EN)
##   collate  en_AU.UTF-8
##   ctype    en_AU.UTF-8
##   tz       Australia/Melbourne
##   date     2020-12-01
##
## - Packages -----
##   package * version  date     lib
##   anicon      0.1.0    2020-06-21 [1]
##   assertthat   0.2.1    2019-03-21 [2]
```

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