

Factors



Getting and Cleaning Data

Factors are for **categorical variables**

Categorical variables: there are a limited number of possible values any data point can take

Example: months

- There are 12 possible months in a calendar year
- For a factor variable containing information about month, there are only 12 possible values each data point can have





<https://forcats.tidyverse.org/>

```
> ?fct|
```

- ? fct_anon
- ? fct_c
- ? fct_collapse
- ? fct_count
- ? fct_drop
- ? fct_expand
- ? fct_explicit_na

fct_anon

Replaces factor levels with arbitrary numeric identifiers. Neither the values nor the order of the levels are preserved.

Press F1 for additional help



```
# All 12 months
```

```
all_months <- c("Jan", "Feb", "Mar", "Apr", "May", "Jun",  
               "Jul", "Aug", "Sep", "Oct", "Nov", "Dec")
```

```
# Our data
```

```
some_months <- c("Mar", "Dec", "Jan", "Apr", "Jul")
```

```
# Sorting some_months -- alphabetical!
```

```
sort(some_months)
```

```
> sort(some_months)
```

```
[1] "Apr" "Dec" "Jan" "Jul" "Mar"
```

Sorts alphabetically



```
# Create a new object containing the some_months data,  
# but specifying the factors as those in all_months  
month_factored <- factor(some_months, levels = all_months)
```

```
# Compare the data before and after factorization  
month_factored  
some_months
```

```
# Now when we sort the factored dataset,  
# it is in the order we specified in all_months!
```

```
sort(month_factored)
```

```
> sort(month_factored)
```

```
[1] Jan Mar Apr Jul Dec
```

```
Levels: Jan Feb Mar Apr May Jun Jul Aug Sep Oct Nov Dec
```

Sorts in order of specified levels!



```
> months_relevel <- fct_relevel(month_factored, "Jul", "Aug",  
"Sep", "Oct", "Nov", "Dec", after = 0)  
>  
> months_relevel  
[1] Mar Dec Jan Apr Jul  
Levels: Jul Aug Sep Oct Nov Dec Jan Feb Mar Apr May Jun  
> sort(months_relevel)  
[1] Jul Dec Jan Mar Apr  
Levels: Jul Aug Sep Oct Nov Dec Jan Feb Mar Apr May Jun
```

Sorts in order of re-ordered levels



```
> months_inorder <- fct_inorder(some_months)
>
> months_inorder
[1] Mar Dec Jan Apr Jul
Levels: Mar Dec Jan Apr Jul
>
> sort(months_inorder)
[1] Mar Dec Jan Apr Jul
Levels: Mar Dec Jan Apr Jul
```

Levels match order of appearance in the
some_months dataset



Chicken Weights by Feed Type

Description

An experiment was conducted to measure and compare the effectiveness of various feed supplements on the growth rate of chickens.

Usage

```
chickwts
```

Format

A data frame with 71 observations on the following 2 variables.

`weight`

a numeric variable giving the chick weight.

`feed`

a factor giving the feed type.

Details

Newly hatched chicks were randomly allocated into six groups, and each group was given a different feed supplement. Their weights in grams after six weeks are given along with feed types.



```
# Take a look at frequency of each level  
# using tabyl() from janitor package
```

```
library(janitor)
```

```
tabyl(chickwts$feed)
```

chickwts\$feed	n	percent
casein	12	0.1690141
horsebean	10	0.1408451
linseed	12	0.1690141
meatmeal	11	0.1549296
soybean	14	0.1971831
sunflower	12	0.1690141

← Least frequent

← Most frequent

```
# Order levels by frequency
```

```
fct_infreq(chickwts$feed) %>% levels()
```

```
[1] "soybean" "casein" "linseed" "sunflower" "meatmeal" "horsebean"
```

Most frequent



Least frequent



```
# Order levels by frequency
```

```
fct_infreq(chickwts$feed) %>% levels()
```

```
[1] "soybean" "casein" "linseed" "sunflower" "meatmeal" "horsebean"
```

Most frequent



Least frequent



fct_rev()

```
# Reverse factor level order
```

```
fct_infreq(chickwts$feed) %>%
```

```
fct_rev() %>%
```

```
levels()
```

```
[1] "horsebean" "meatmeal" "sunflower" "linseed" "casein" "soybean"
```

Least frequent



Most frequent



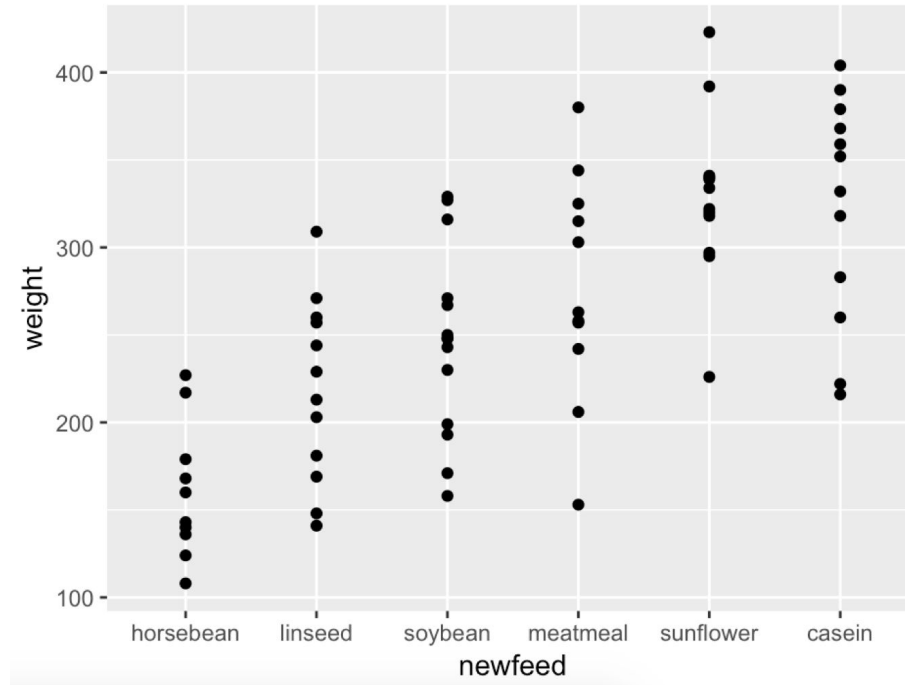
```
# Order levels by a second numeric variable
```

```
chickwts %>%
```

```
  mutate(newfeed = fct_reorder(feed, weight)) %>% # Reorder feed types by chicken weight
```

```
  ggplot(aes(x = newfeed, y = weight)) + # Plot the feed type on the X and chicken weights on the Y axes
```

```
  geom_point() # Plot this data as points
```



Feed levels ordered by
value of weight

```
# We can use mutate to create a new column
```

```
# and fct_recode() to:
```

```
# 1. group horsebean and soybean, and sunflower and linseed into single levels
```

```
# 2. rename all the other levels
```

```
chickwts %>%
```

```
  mutate(feed_recode = fct_recode(feed,  
    "seed"      = "linseed",  
    "bean"      = "horsebean",  
    "bean"      = "soybean",  
    "meal"      = "meatmeal",  
    "seed"      = "sunflower",  
    "casein"    = "casein"
```

```
  )) %>%
```

```
  tabyl(feed_recode)
```

feed_recode	n	percent
casein	12	0.1690141
bean	24	0.3380282
seed	24	0.3380282
meal	11	0.1549296

Group horsebean
and soybean into
a single level
called "bean"



```
# Convert numeric variable to factor
```

```
chickwts %>%
```

```
  mutate(weight_recode = ifelse(weight <= 200, "low", "high"),
```

```
         weight_recode = factor(weight_recode)) %>%
```

```
  tabyl(weight_recode)
```

```
weight_recode  n    percent
```

```
      high  54  0.7605634
```

```
      low   17  0.2394366
```



Summarizing: Factors



Getting and Cleaning Data