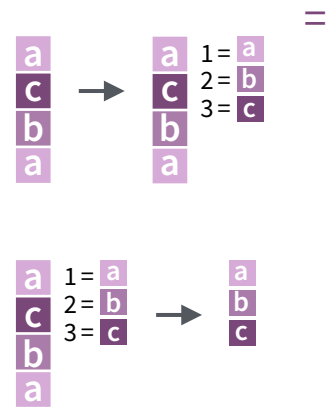
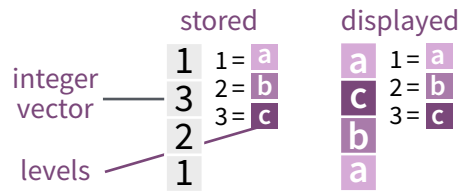


Factors with forcats : : CHEAT SHEET

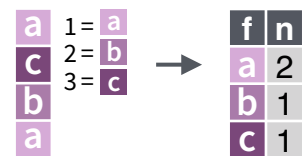
The **forcats** package provides tools for working with factors, which are R's data structure for categorical data.

Factors

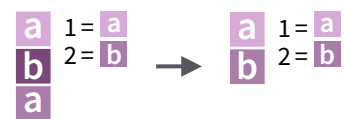
R represents categorical data with factors. A **factor** is an integer vector with a **levels** attribute that stores a set of mappings between integers and categorical values. When you view a factor, R displays not the integers, but the levels associated with them.



Inspect Factors



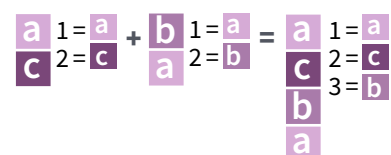
fct_count(f, sort = FALSE, prop = FALSE) Count the number of values with each level. `fct_count(f)`



fct_match(f, lvls) Check for lvls in f. `fct_match(f, "a")`

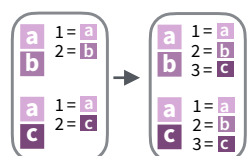
fct_unique(f) Return the unique values, removing duplicates. `fct_unique(f)`

Combine Factors



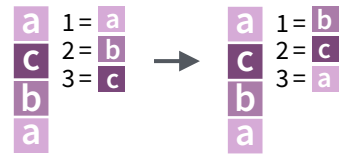
fct_c(...) Combine factors with different levels. Also **fct_cross()**.

f1 <- factor(c("a", "c"))
f2 <- factor(c("b", "a"))
fct_c(f1, f2)

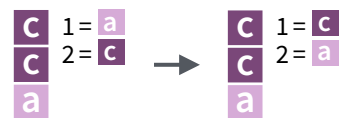


fct_unify(fs, levels = lvls_union(fs)) Standardize levels across a list of factors. `fct_unify(list(f2, f1))`

Change the order of levels



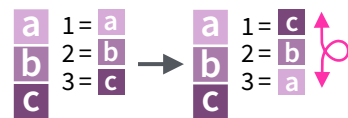
fct_relevel(.f, ..., after = 0L) Manually reorder factor levels. `fct_relevel(f, c("b", "c", "a"))`



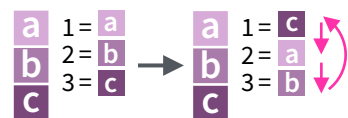
fct_infreq(f, ordered = NA) Reorder levels by the frequency in which they appear in the data (highest frequency first). Also **fct_inseq()**.
f3 <- factor(c("c", "c", "a"))
`fct_infreq(f3)`



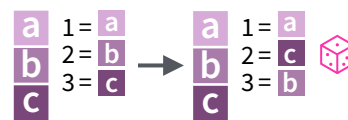
fct_inorder(f, ordered = NA) Reorder levels by order in which they appear in the data. `fct_inorder(f2)`



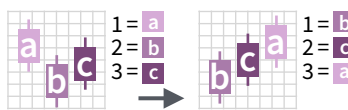
fct_rev(f) Reverse level order. **f4 <- factor(c("a", "b", "c"))**
`fct_rev(f4)`



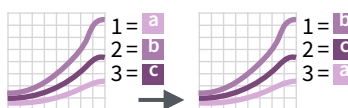
fct_shift(f) Shift levels to left or right, wrapping around end. `fct_shift(f4)`



fct_shuffle(f, n = 1L) Randomly permute order of factor levels. `fct_shuffle(f4)`

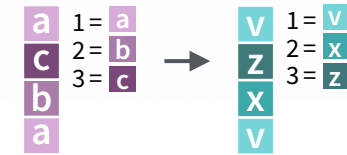


fct_reorder(.f, .x, .fun = median, ..., .desc = FALSE) Reorder levels by their relationship with another variable. `boxplot(PlantGrowth, weight ~ fct_reorder(group, weight))`

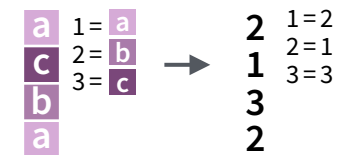


fct_reorder2(.f, .x, .y, .fun = last2, ..., .desc = TRUE) Reorder levels by their final values when plotted with two other variables. `ggplot(diamonds, aes(carat, price, color = fct_reorder2(color, carat, price))) + geom_smooth()`

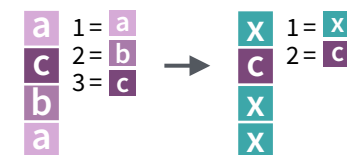
Change the value of levels



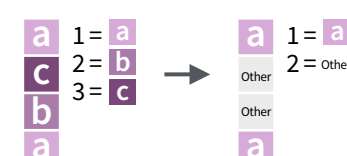
fct_recode(.f, ...) Manually change levels. Also **fct_relabel()** which obeys purrr::map syntax to apply a function or expression to each level. `fct_recode(f, v = "a", x = "b", z = "c")`
`fct_relabel(f, ~ paste0("x", .x))`



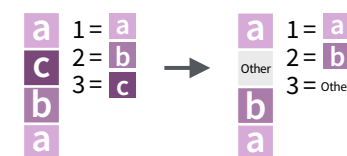
fct_anon(f, prefix = "") Anonymize levels with random integers. `fct_anon(f)`



fct_collapse(.f, ..., other_level = NULL) Collapse levels into manually defined groups. `fct_collapse(f, x = c("a", "b"))`



fct_lump_min(f, min, w = NULL, other_level = "Other") Lumps together factors that appear fewer than min times. Also **fct_lump_n()**, **fct_lump_prop()**, and **fct_lump_lowfreq()**. `fct_lump_min(f, min = 2)`



fct_other(f, keep, drop, other_level = "Other") Replace levels with "other." `fct_other(f, keep = c("a", "b"))`

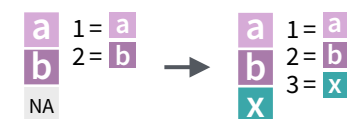
Add or drop levels



fct_drop(f, only) Drop unused levels. **f5 <- factor(c("a", "b"), c("a", "b", "x"))**
f6 <- fct_drop(f5)



fct_expand(f, ...) Add levels to a factor. `fct_expand(f6, "x")`



fct_na_value_to_level(f, level = "(Missing)") Assigns a level to NAs to ensure they appear in plots, etc. **f7 <- factor(c("a", "b", NA))**
`fct_na_value_to_level(f7, level = "(Missing)")`