
Grouping with other verbs

Introduction	
Learning objectives	
Packages	
Datasets	
Arranging by group	
<code>arrange()</code> can group automatically	
Filtering by group	
Filtering with nested groupings	
Mutating by group	
Mutating with nested groupings	
Wrap up	

Introduction

Data wrangling often involves applying the same operations separately to different groups within the data. This pattern, sometimes called “split-apply-combine”, is easily accomplished in {dplyr} by chaining the `group_by()` verb with other wrangling verbs like `filter()`, `mutate()`, and `arrange()` (all of which you have seen before!).

In this lesson, you’ll become confident with these kinds of grouped manipulations.

Let’s get started.

Learning objectives

1. You can use `group_by()` with `arrange()`, `filter()`, and `mutate()` to conduct grouped operations on a data frame.

Packages

This lesson will require the {tidyverse} suite of packages and the {here} package:

```
if(!require(pacman)) install.packages("pacman")
pacman::p_load(tidyverse, here)
```

Datasets

In this lesson, we will again use data from the COVID-19 serological survey conducted in Yaounde, Cameroon. Below, we import the data, create a small data frame subset, `yao` and an even smaller subset, `yao_sex_weight`.

```
yao <-  
  read_csv(here::here('data/yaounde_data.csv')) %>%  
  select(sex, age, age_category, weight_kg, occupation, igg_result,  
         igm_result)
```

yao

```
## # A tibble: 5 × 7  
##   sex      age age_category weight_kg occupation  
##   <chr> <dbl> <chr>          <dbl> <chr>  
## 1 Female  45 45 - 64           95 Informal worker  
## 2 Male    55 45 - 64           96 Salaried worker  
## 3 Male    23 15 - 29           74 Student  
## 4 Female  20 15 - 29           70 Student  
## 5 Female  55 45 - 64           67 Trader--Farmer  
## # ... with 2 more variables: igg_result <chr>,  
## #   igm_result <chr>
```

```
yao_sex_weight <-  
  yao %>%  
  select(sex, weight_kg)
```

yao_sex_weight

```
## # A tibble: 5 × 2  
##   sex      weight_kg  
##   <chr>      <dbl>  
## 1 Female      95  
## 2 Male       96  
## 3 Male       74  
## 4 Female      70  
## 5 Female      67
```

For practice questions, we will also use the sarcopenia data set that you have seen previously:

```
sarcopenia <- read_csv(here::here('data/sarcopenia_elderly.csv'))
```

sarcopenia

```
## # A tibble: 5 × 9
##   number   age age_group sex_male_1_female_0 marital_status
##   <dbl> <dbl> <chr>          <dbl> <chr>
## 1     7  60.8 Sixties             0 married
## 2     8  72.3 Seventies          1 married
## 3     9  62.6 Sixties             0 married
## 4    12   72  Seventies             0 widow
## 5    13  60.1 Sixties             0 married
## # ... with 4 more variables: height_meters <dbl>,
## #   weight_kg <dbl>, grip_strength_kg <dbl>, ...
```

Arranging by group

The `arrange()` function orders the rows of a data frame by the values of selected columns. This function is only sensitive to groupings when we set its argument `.by_group` to `TRUE`. To illustrate this, consider the `yao_sex_weight` data frame:

```
yao_sex_weight
```

```
## # A tibble: 5 × 2
##   sex      weight_kg
##   <chr>      <dbl>
## 1 Female      95
## 2 Male       96
## 3 Male       74
## 4 Female      70
## 5 Female      67
```

We can arrange this data frame by weight like so:

```
yao_sex_weight %>%
  arrange(weight_kg)
```

```
## # A tibble: 5 × 2
##   sex      weight_kg
##   <chr>      <dbl>
## 1 Female      14
## 2 Male       15
## 3 Male       15
## 4 Male       15
## 5 Female      15
```

As expected, lower weights have been brought to the top of the data frame.

If we first group the data, we might expect a different output:

```
yao_sex_weight %>%  
  group_by(sex) %>%  
  arrange(weight_kg)
```

```
## # A tibble: 5 × 2  
## # Groups:   sex [2]  
##   sex    weight_kg  
##   <chr>      <dbl>  
## 1 Female        14  
## 2 Male          15  
## 3 Male          15  
## 4 Male          15  
## 5 Female        15
```

But as you see, the arrangement is still the same.

Only when we set the `.by_group` argument to `TRUE` do we get something different:

```
yao_sex_weight %>%  
  group_by(sex) %>%  
  arrange(weight_kg, .by_group = TRUE)
```

```
## # A tibble: 5 × 2  
## # Groups:   sex [1]  
##   sex    weight_kg  
##   <chr>      <dbl>  
## 1 Female        14  
## 2 Female        15  
## 3 Female        16  
## 4 Female        16  
## 5 Female        18
```

Now, the data is *first* sorted by sex (all women first), and then by weight.

`arrange()` can group automatically

In reality we do not need `group_by()` to arrange by group; we can simply put multiple variables in the `arrange()` function for the same effect.

So this simple `arrange()` statement:

```
yao_sex_weight %>%  
  arrange(sex, weight_kg)
```

```
## # A tibble: 5 × 2  
##   sex    weight_kg
```

```
##      <chr>      <dbl>
## 1 Female      14
## 2 Female      15
## 3 Female      16
## 4 Female      16
## 5 Female      18
```

is equivalent to the more complex `group_by()`, `arrange()` statement used before:

```
yao_sex_weight %>%
  group_by(sex) %>%
  arrange(weight_kg, .by_group = TRUE)
```

The code `arrange(sex, weight_kg)` tells R to arrange the rows *first* by sex, and then by weight.

Obviously, this syntax, with just `arrange()`, and no `group_by()` is simpler, so you can stick to it.

`desc()` **for descending order**

Recall that to arrange *in descending order*, we can wrap the target variable in `desc()`. So, for example, to sort by sex and weight, but with the heaviest people on top, we can run:

```
yao_sex_weight %>%
  arrange(sex, desc(weight_kg))
```

```
## # A tibble: 5 × 2
##   sex      weight_kg
##   <chr>      <dbl>
## 1 Female      162
## 2 Female      161
## 3 Female      158
## 4 Female      135
## 5 Female      129
```

PRACTICE



With an `arrange()` call, sort the `sarcopenia` data first by sex and then by grip strength. (If done correctly, the first row should be of a woman with a grip strength of 1.3 kg). To make the arrangement clear, you should first `select()` the sex and grip strength variables.

```
# Complete the code with your answer:
Q_grip_strength_arranged <-
  sarcopenia %>%
  select(_____) %>%
  arrange(_____)
```

PRACTICE



The `sarcopenia` dataset contains a column, `age_group`, which stores age groups as a string (the age groups are “Sixties”, “Seventies” and “Eighties”). Convert this variable to a factor with the levels in the right order (first “Sixties” then “Seventies” and so on). (Hint: Look back on the `case_when()` lesson if you do not see how to relevel a factor.)

Then, with a nested `arrange()` call, arrange the data first by the newly-created `age_group` factor variable (younger individuals first) and then by `height_meters`, with shorter individuals first.

```
# Complete the code with your answer:  
Q_age_group_height <-  
  sarcopenia
```

Filtering by group

The `filter()` function keeps or drops rows based on a condition. If `filter()` is applied to grouped data, the filtering operation is carried out separately for each group.

To illustrate this, consider again the `yao_sex_weight` data frame:

```
yao_sex_weight
```

```
## # A tibble: 5 × 2  
##   sex      weight_kg  
##   <chr>      <dbl>  
## 1 Female      95  
## 2 Male       96  
## 3 Male       74  
## 4 Female     70  
## 5 Female     67
```

If we want to filter the data for the heaviest person, we could run:

```
yao_sex_weight %>%  
  filter(weight_kg == max(weight_kg))
```

```
## # A tibble: 1 × 2  
##   sex      weight_kg  
##   <chr>      <dbl>  
## 1 Female     162
```


But if we want to get heaviest person per sex group (the heaviest man *and* the heaviest woman), we can use `group_by(sex)` then `filter()`:

```
yao_sex_weight %>%
  group_by(sex) %>%
  filter(weight_kg == max(weight_kg))
```

```
## # A tibble: 2 × 2
## # Groups:   sex [2]
##   sex    weight_kg
##   <chr>      <dbl>
## 1 Male         128
## 2 Female        162
```

Great! The code above can be translated as “For each sex group, keep the row with the maximum `weight_kg` value”.

Filtering with nested groupings

`filter()` will work fine with any number of nested groupings.

For example, if we want to see the heaviest man and heaviest woman *per age group* we could run the following on the `yao` data frame:

```
yao %>%
  group_by(sex, age_category) %>%
  filter(weight_kg == max(weight_kg))
```

This code groups by sex *and* age category, and then finds the heaviest person in each sub-category.

(Why do we have 10 rows in the output? Well, 2 sex groups x 5 groups age groups = 10 unique groupings.)

The output is a bit scattered though, so we can chain this with the `arrange()` function, to arrange by sex and age group.

```
yao %>%
  group_by(sex, age_category) %>%
  filter(weight_kg == max(weight_kg)) %>%
  arrange(sex, age_category)
```

Now the data is easier to read. All women come first, then men. But we see notice a weird arrangement of the age groups! Those aged 5 to 14 should come *first* in the arrangement. Of course, we’ve learned how to fix this—the `factor()` function, and its `levels` argument:

```
yao %>%
  mutate(age_category = factor(
    age_category,
    levels = c("5 - 14", "15 - 29", "30 - 44", "45 - 64", "65 +")
  )) %>%
  group_by(sex, age_category) %>%
  filter(weight_kg == max(weight_kg)) %>%
  arrange(sex, age_category)
```

Now we have a nice and well-arranged output!



Group the `sarcopenia` data frame by age group and sex, then filter for the highest skeletal muscle index in each (nested) group.

```
# Complete the code with your answer:
Q_max_skeletal_muscle_index <-
  sarcopenia
```

Mutating by group

`mutate()` is used to modify columns or to create new ones. With grouped data, `mutate()` operates over each group independently.

Let's first consider a regular `mutate()` call, not a grouped one. Imagine that you wanted to add a column that ranks respondents by weight. This can be done with the `rank()` function inside a `mutate()` call:

```
yao_sex_weight %>%
  mutate(weight_rank = rank(weight_kg))
```

```
## # A tibble: 5 × 3
##   sex      weight_kg weight_rank
##   <chr>      <dbl>      <dbl>
## 1 Female      95        901
## 2 Male       96        908
## 3 Male       74        640.
## 4 Female     70        564.
## 5 Female     67        502.
```

The output shows that the first row is the 901st lightest individual. But it would be more intuitive to rank in descending order with the heaviest person first. We can do this with the `desc()` function:

```
yao_sex_weight %>%
  mutate(weight_rank = rank(desc(weight_kg)))
```

```
## # A tibble: 5 × 3
##   sex    weight_kg weight_rank
##   <chr>      <dbl>      <dbl>
## 1 Female         95         71
## 2 Male          96         64
## 3 Male          74        332.
## 4 Female         70        408.
## 5 Female         67        470.
```

The output shows that the person in the first row is the 71st heaviest individual.

Now, let's try to write a grouped `mutate()` call. Imagine we want to add this weight rank column *per sex group* in the data frame. That is, we want to know each person's weight rank in their sex category. In this case, we can chain `group_by(sex)` with `mutate()`:

```
yao_sex_weight %>%
  group_by(sex) %>%
  mutate(weight_rank = rank(desc(weight_kg)))
```

```
## # A tibble: 5 × 3
## # Groups:   sex [2]
##   sex    weight_kg weight_rank
##   <chr>      <dbl>      <dbl>
## 1 Female         95         53.5
## 2 Male          96         13.5
## 3 Male          74         148
## 4 Female         70        220.
## 5 Female         67        250.
```

Now we see that the person in the first row is the 53rd heaviest *woman*. (The .5 indicates that this rank is a tie with someone else in the data.)

We could also arrange the data to make things clearer:

```
yao_sex_weight %>%
  group_by(sex) %>%
  mutate(weight_rank = rank(desc(weight_kg))) %>%
  arrange(sex, weight_rank)
```

```
## # A tibble: 5 × 3
## # Groups:   sex [1]
##   sex    weight_kg weight_rank
##   <chr>      <dbl>      <dbl>
## 1 Female         162          1
## 2 Female         161          2
```

```
## 3 Female      158      3
## 4 Female      135      4
## 5 Female      129      5
```

Mutating with nested groupings

Of course, as with the other verbs we have seen, `mutate()` also works with nested groups.

For example, below we create the nested grouping of age *and* sex with the `yao` data frame, then add a rank column with `mutate()`:

```
yao %>%
  group_by(sex, age_category) %>%
  mutate(weight_rank = rank(desc(weight_kg)))

## # A tibble: 5 × 8
## # Groups:   sex, age_category [4]
##   sex      age age_category weight_kg occupation
##   <chr> <dbl> <chr>          <dbl> <chr>
## 1 Female    45 45 - 64             95 Informal worker
## 2 Male      55 45 - 64             96 Salaried worker
## 3 Male      23 15 - 29             74 Student
## 4 Female    20 15 - 29             70 Student
## 5 Female    55 45 - 64             67 Trader--Farmer
## # ... with 3 more variables: igg_result <chr>,
## #   igm_result <chr>, weight_rank <dbl>
```

The output shows that the person in the first row is 20th heaviest *woman in the 45 to 64 age group*.



With the `sarcopenia` data, group by `age_group`, then in a new variable called `grip_strength_rank`, compute the per-age-group rank of each individual's grip strength. (To compute the rank, use `mutate()` and the `rank()` function with its default ties method.)

```
# Complete the code with your answer:
Q_rank_grip_strength <-
  sarcopenia
```

WATCH OUT



Remember to ungroup data before further analysis

As has been mentioned before, it is important ungroup your data before doing further analysis.

Consider this last example, where we computed the weight rank of individuals per age and sex group:

```
yao %>%
  group_by(sex, age_category) %>%
  mutate(weight_rank = rank(desc(weight_kg)))

## # A tibble: 5 × 8
## # Groups:   sex, age_category [4]
##   sex      age age_category weight_kg occupation
##   <chr> <dbl> <chr>          <dbl> <chr>
## 1 Female    45 45 - 64             95 Informal worker
## 2 Male      55 45 - 64             96 Salaried worker
## 3 Male      23 15 - 29             74 Student
## 4 Female    20 15 - 29             70 Student
## 5 Female    55 45 - 64             67 Trader--Farmer
## # ... with 3 more variables: igg_result <chr>,
## #   igm_result <chr>, weight_rank <dbl>
```

WATCH OUT



If, in the process of analysis, you stored this output as a new data frame:

```
yao_modified <-
  yao %>%
  group_by(sex, age_category) %>%
  mutate(weight_rank = rank(desc(weight_kg)))
```

And then, later on, you picked up the data frame and tried some other analysis, for example, filtering to get the oldest person in the data:

```
yao_modified %>%
  filter(age == max(age))

## # A tibble: 5 × 8
## # Groups:   sex, age_category [5]
##   sex      age age_category weight_kg occupation
##   <chr> <dbl> <chr>          <dbl> <chr>
## 1 Male      65 45 - 64             93 Retired
## 2 Male      78 65 +              95 Retired--Informal
wor...
## 3 Male      14 5 - 14             44 Student
## 4 Female    44 30 - 44             67 Home-maker
## 5 Female    79 65 +             40 Retired
```

```
## # ... with 3 more variables: igg_result <chr>,  
## #   igm_result <chr>, weight_rank <dbl>
```

You might be confused by the output! Why are there 55 rows of “oldest people”?

This would be because you forgot to ungroup the data before storing it for further analysis. Let's do this properly now

WATCH OUT



```
yao_modified <-  
yao %>%  
  group_by(sex, age_category) %>%  
  mutate(weight_rank = rank(desc(weight_kg))) %>%  
  ungroup()
```

Now we can correctly obtain the oldest person/people in the data set:

```
yao_modified %>%  
  filter(age == max(age))
```

```
## # A tibble: 2 × 8  
##   sex      age age_category weight_kg occupation igg_result  
##   <chr> <dbl> <chr>          <dbl> <chr>      <chr>  
## 1 Female    79 65 +             40 Retired   Negative  
## 2 Female    79 65 +             81 Home-maker Negative  
## # ... with 2 more variables: igm_result <chr>,  
## #   weight_rank <dbl>
```

Wrap up

`group_by()` is a marvelous tool for arranging, mutating, filtering based on the groups within a single or multiple variables.

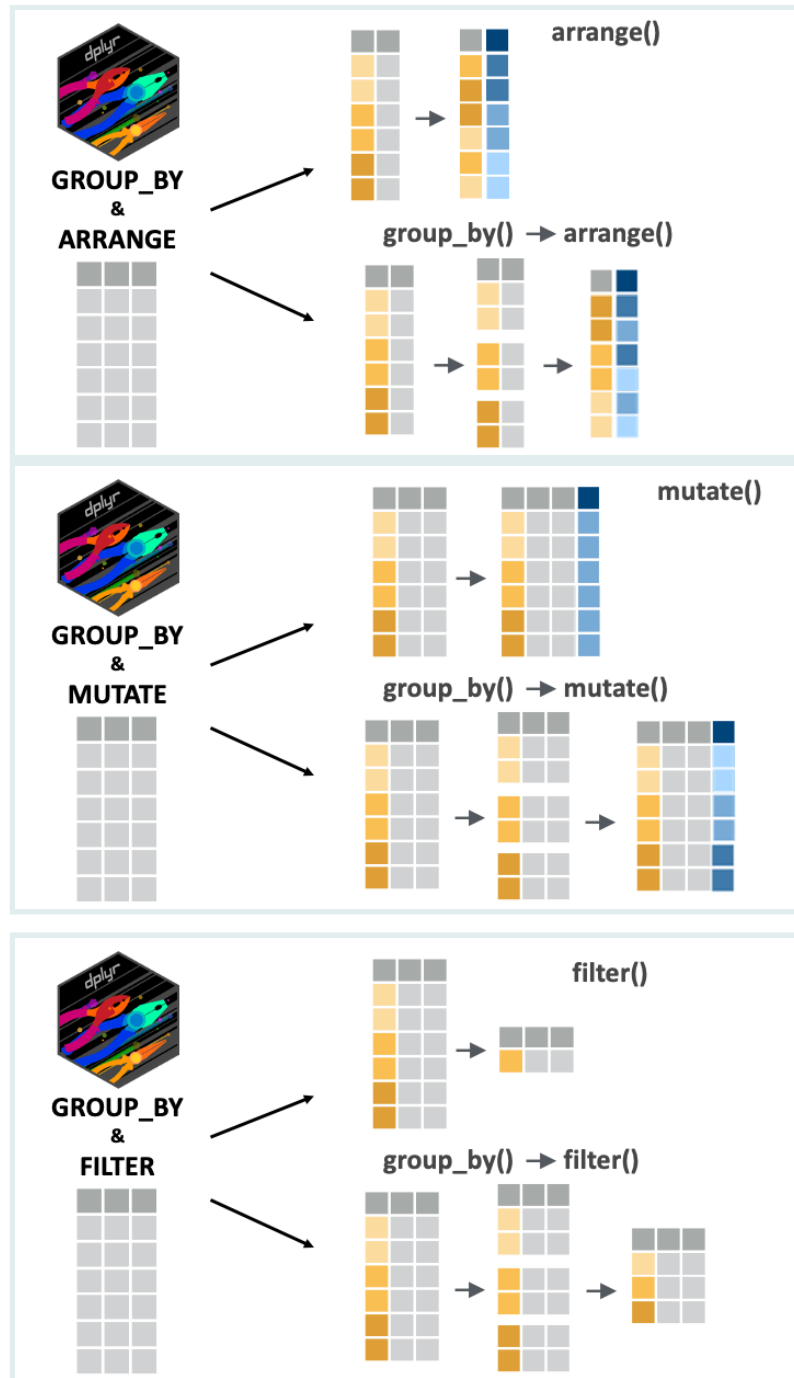


Fig: filter() and group_by()

There are numerous ways of combining these verbs to manipulate your data. We invite you to take some time and to try these verbs out in different combinations!

See you next time!

Contributors

The following team members contributed to this lesson:



LAURE VANCAUWENBERGHE

Data analyst, the GRAPH Network

A firm believer in science for good, striving to ally programming, health and education



KENE DAVID NWOSU

Data analyst, the GRAPH Network

Passionate about world improvement

References

Some material in this lesson was adapted from the following sources:

- Horst, A. (2022). *Dplyr-learnr*. <https://github.com/allisonhorst/dplyr-learnr> (Original work published 2020)
- *Group by one or more variables*. (n.d.). Retrieved 21 February 2022, from https://dplyr.tidyverse.org/reference/group_by.html
- *Create, modify, and delete columns – Mutate*. (n.d.). Retrieved 21 February 2022, from <https://dplyr.tidyverse.org/reference/mutate.html>
- *Subset rows using column values – Filter*. (n.d.). Retrieved 21 February 2022, from <https://dplyr.tidyverse.org/reference/filter.html>
- *Arrange rows by column values – Arrange*. (n.d.). Retrieved 21 February 2022, from <https://dplyr.tidyverse.org/reference/arrange.html>

Artwork was adapted from:

- [Horst, A. \(2022\). *R & stats illustrations by Allison Horst*. <https://github.com/allisonhorst/stats-illustrations> \(Original work published 2018\)](https://github.com/allisonhorst/stats-illustrations)

title: “Grouping operations on other verbs” credits: “This document serves as an accompaniment for a lesson found on <https://thegraphcourses.org>.

The GRAPH Courses is a project of the Global Research and Analyses for Public

Health (GRAPH) Network, with the support of the World Health Organization (WHO) and other partners” date: “November 2022” author: “The GRAPH Courses team” –