Herding cats with dplyr

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dplyr

80% of the work involved with data analysis involves cleaning and shaping the data until it's in the state you need. Bracket subsetting is handy, but it can be cumbersome and difficult to read, especially for complicated operations. Enter dplyr!

dplyr is a package for making data manipulation easier. (It does a lot more too, but this is what we'll focus on).

Unlike the subsetting commands we've already worked on, dplyr is designed to be highly expressive, and highly readable. It's structured around a set of verbs, or grammar of data manipulation. The core functions we'll talk about are below:

- select
- arrange
- filter
- group_by
- mutate
- summarise/summarize

Getting the package

```
install.packages("dplyr")
```

Again, you may be asked to choose a CRAN mirror; RStudio is a good choice.

Unlike the rmarkdown package, we'll be using dplyr in the console. In order to do that, we need to *load* the package into our environment so we can access functions from dplyr. We do this with the library command:

```
library("dplyr")
```

You only need to install a package once per computer, but you need to load it every time you open a new R session and want to use that package.

Subsetting Data

The first two dplyr commands we'll use help us to subset our data by rows and columns.

select

The first command we'll use is **select**, which allows us to choose columns from our dataset. Let's use our **cats** dataset and select only the **coat** column; we did this previously with

```
cats[, "coat"]

## [1] tabby maltese brown black calico tabby brown brown
## [9] black maltese
## Levels: black brown calico maltese tabby
```

With dplyr, we don't need to enclose our column names in quotes

```
select(cats, coat)
```

```
##
         coat
## 1
        tabby
## 2
      maltese
## 3
        brown
## 4
        black
## 5
       calico
## 6
        tabby
## 7
        brown
## 8
        brown
        black
## 10 maltese
```

Notice how the output differs slightly; all the main dplyr verbs behave consistently in that their inputs and outputs are both data.frames, rather than returning a simple vector as the bracket-indexing method did. All of the main "verbs" we'll talk about will return a data.frame as their result.

We can select more columns by giving select additional arguments, and our output data.frame will have columns according to the order of our arguments

select(cats, coat, cat id) ## coat cat id ## 1 tabby 321 ## 2 maltese 250 ## 3 brown 219 ## 4 black 182 ## 5 calico 107 234 ## 6 tabby ## 7 brown 196 ## 8 brown 311 ## 9 130 black ## 10 maltese 349

filter

So where select allowed us to select *columns*, filter operated on *rows*. Say we want to see the all the cats with black coats; we saw earlier how to use that using bracket-indexing:

```
cats[cats$coat == "black", ]
##
                street coat
                                       age weight fixed wander dist roamer
                                 sex
## 4
        140 Robin Way black female 7.172
                                            8.053
                                                       1
                                                                0.030
                                                                           no
## 9 16982 Kennedy Rd black female 2.851
                                                       0
                                                                0.065
                                            3.291
                                                                           no
##
     cat id
## 4
        182
## 9
        130
In dplyr, this looks like
filter(cats, coat == "black")
##
                                       age weight fixed wander dist roamer
                street coat
                                 sex
        140 Robin Way black female 7.172
                                            8.053
                                                       1
                                                                0.030
                                                                           no
## 2 16982 Kennedy Rd black female 2.851
                                            3.291
                                                       0
                                                                0.065
                                                                           no
##
     cat_id
## 1
        182
## 2
        130
```

Notice we don't have to use the \$ operator to tell filter where the coat column is; it's smart enough to assume we want the coat column from the data.frame we passed in.

arrange

Maybe you have a set of observations in your data that you want to organize by their value. arrange allows us to change the order of rows in our dataset based on their values.

arrange(cats, coat) ## age weight fixed wander dist coat street sex ## 1 140 Robin Way black female 7.172 8.053 1 0.030 0 ## 2 16982 Kennedy Rd black female 2.851 3.291 0.065 ## 3 201 Hollywood Ave brown female 4.601 3.947 1 0.076 ## 4 5.626 1 115 Via Santa Maria brown male 6.917 0.097 ## 5 303 Harding Ave brown male 3.713 3.982 1 0.033 ## 6 135 Charles St calico male 4.660 6.193 1 0.085 ## 7 242 Harding Ave maltese female 8.234 12.368 1 0.033 ## 8 16528 Marchmont Dr maltese female 4.594 6.994 0 0.059 ## 9 tabby female 3.003 0 Los Robles Way 3.993 0.040 ## 10 130 Vista Del Campo tabby female 3.796 3.860 1 0.085 ## roamer cat id ## 1 182 no ## 2 130 no ## 3 219 no ## 4 yes 196 ## 5 no 311 ## 6 107 yes ## 7 250 no ## 8 349 no ## 9 321 nο ## 10 234 no # you can include additional columns to help sort the data arrange(cats, coat, sex) ## age weight fixed wander dist street coat sex ## 1 140 Robin Way black female 7.172 8.053 1 0.030 ## 2 16982 Kennedy Rd black female 2.851 3.291 0 0.065 ## 3 3.947 1 201 Hollywood Ave brown female 4.601 0.076 ## 4 115 Via Santa Maria brown male 6.917 5.626 1 0.097 ## 5 303 Harding Ave brown male 3.713 3.982 1 0.033 ## 6 1 135 Charles St calico male 4.660 6.193 0.085 ## 7 242 Harding Ave maltese female 8.234 12.368 1 0.033 ## 8 16528 Marchmont Dr maltese female 4.594 6.994 0 0.059 ## 9 0 Los Robles Way tabby female 3.003 3.993 0.040 ## 10 130 Vista Del Campo tabby female 3.796 3.860 1 0.085 ## roamer cat id ## 1 no 182 ## 2 130 no ## 3 219 no

4

5

196

311

yes

no

```
## 6 yes 107
## 7 no 250
## 8 no 349
## 9 no 321
## 10 no 234
```

mutate

10

349

no

One common task in working with data is updating/cleaning some of the values in columns. mutate allows us to do this relatively easily. Let's say I don't want a lot of decimal places in one of my measurements. I can use mutate to update my existing variable:

```
mutate(cats, weight = round(weight, 2))
##
                                               age weight fixed wander dist
                    street
                               coat
                                        sex
## 1
                                                               0
            Los Robles Way
                              tabby female 3.003
                                                     3.99
                                                                        0.040
## 2
          242 Harding Ave maltese female 8.234
                                                    12.37
                                                               1
                                                                        0.033
                              brown female 4.601
## 3
        201 Hollywood Ave
                                                     3.95
                                                               1
                                                                        0.076
## 4
             140 Robin Way
                                                                        0.030
                              black female 7.172
                                                     8.05
                                                               1
## 5
            135 Charles St
                             calico
                                       male 4.660
                                                     6.19
                                                               1
                                                                        0.085
## 6
      130 Vista Del Campo
                              tabby female 3.796
                                                     3.86
                                                               1
                                                                        0.085
## 7
      115 Via Santa Maria
                                       male 6.917
                                                               1
                                                                        0.097
                              brown
                                                     5.63
## 8
          303 Harding Ave
                              brown
                                       male 3.713
                                                     3.98
                                                               1
                                                                        0.033
## 9
          16982 Kennedy Rd
                              black female 2.851
                                                     3.29
                                                               0
                                                                        0.065
## 10
       16528 Marchmont Dr maltese female 4.594
                                                     6.99
                                                                        0.059
##
      roamer cat id
## 1
          no
                 321
## 2
                 250
          no
## 3
                 219
          no
## 4
                 182
          no
## 5
         ves
                 107
## 6
                 234
          no
## 7
                 196
         yes
## 8
                 311
          no
## 9
                 130
          no
```

Another common task is generating a new column based on values that are already in the dataset you are working on. mutate helps us do this, and tacks a new column to the end of our data frame.

```
# let's say you want to add two variables together
mutate(cats, new_variable = age + weight)
```

```
## street coat sex age weight fixed wander_dist
## 1 Los Robles Way tabby female 3.003 3.993 0 0.040
## 2 242 Harding Ave maltese female 8.234 12.368 1 0.033
```

```
## 3
        201 Hollywood Ave
                             brown female 4.601
                                                   3.947
                                                                      0.076
                                                             1
## 4
                                                             1
             140 Robin Way
                             black female 7.172
                                                  8.053
                                                                      0.030
## 5
           135 Charles St calico
                                      male 4.660
                                                   6.193
                                                             1
                                                                      0.085
## 6
      130 Vista Del Campo
                             tabby female 3.796
                                                   3.860
                                                             1
                                                                      0.085
      115 Via Santa Maria
## 7
                             brown
                                      male 6.917
                                                   5.626
                                                             1
                                                                      0.097
## 8
                                      male 3.713
                                                             1
          303 Harding Ave
                             brown
                                                   3.982
                                                                      0.033
## 9
                             black female 2.851
                                                  3.291
                                                             0
                                                                      0.065
         16982 Kennedy Rd
## 10
       16528 Marchmont Dr maltese female 4.594 6.994
                                                             0
                                                                      0.059
##
      roamer cat id new variable
## 1
                 321
                            6.996
          no
## 2
          no
                 250
                           20.602
## 3
                 219
                            8.548
          no
## 4
                 182
                           15.225
          no
## 5
                 107
                           10.853
         yes
## 6
                 234
                            7.656
          no
## 7
                           12.543
                 196
         yes
## 8
                            7.695
          no
                 311
## 9
                 130
                            6.142
          no
## 10
                 349
                           11.588
          no
```

you can include as many new variables as you want, separated by a comma
mutate(cats, new_var_1 = age + weight, new_var_2 = age * weight)

```
##
                    street
                              coat
                                       sex
                                             age weight fixed wander dist
## 1
                             tabby female 3.003
                                                             0
           Los Robles Way
                                                 3.993
                                                                     0.040
## 2
          242 Harding Ave maltese female 8.234 12.368
                                                             1
                                                                     0.033
## 3
        201 Hollywood Ave
                             brown female 4.601
                                                  3.947
                                                             1
                                                                     0.076
## 4
            140 Robin Way
                             black female 7.172
                                                  8.053
                                                             1
                                                                     0.030
## 5
           135 Charles St calico
                                     male 4.660
                                                  6.193
                                                             1
                                                                     0.085
## 6
      130 Vista Del Campo
                             tabby female 3.796
                                                  3.860
                                                             1
                                                                     0.085
## 7
      115 Via Santa Maria
                             brown
                                     male 6.917
                                                  5.626
                                                             1
                                                                     0.097
## 8
          303 Harding Ave
                             brown
                                     male 3.713 3.982
                                                             1
                                                                     0.033
## 9
         16982 Kennedy Rd
                             black female 2.851
                                                  3.291
                                                             0
                                                                     0.065
       16528 Marchmont Dr maltese female 4.594
                                                             0
                                                                     0.059
## 10
                                                  6.994
##
      roamer cat id new var 1 new var 2
## 1
                 321
                         6.996
                                11.990979
          no
## 2
          no
                 250
                        20.602 101.838112
## 3
          no
                219
                         8.548
                                18.160147
## 4
                182
                        15.225
                                57.756116
          no
## 5
                107
                        10.853
                                28.859380
         yes
## 6
                234
                         7.656
                                14.652560
          no
## 7
                        12.543
                196
                                38.915042
         yes
## 8
                311
                         7.695
                                14.785166
          no
## 9
                130
                         6.142
                                 9.382641
          no
## 10
                        11.588
                                32.130436
          no
                349
```

Pipes

You'll often find yourself needing to use multiple functions in a row to organize some data that you're working on. This can sometimes lead to dense code that is difficult to read.

```
# for example
sort(round(sqrt(cats$age * 2), 3))
```

```
## [1] 2.388 2.451 2.725 2.755 3.031 3.033 3.053 3.719 3.787 4.058
```

In the code above, I have multiple steps to get my result, but you have to read what's going on from the inside out. This can be cumbersome, especially if you need to understand how one function's output influences the next operation.

Using Pipes

dplyr includes a special operator designed to make code flow and appear more readable.

It's written as %>%, and you can call it the "pipe" operator.

Our example above can be re-written as:

```
cats$age * 2 %>%
  sqrt() %>%
  round(3) %>%
  sort()
```

```
## [1] 4.246242 11.642876 6.505814 10.141208 6.589240 5.367544 9.780638 ## [8] 5.250182 4.031314 6.495916
```

Instead of being nested within a bunch of commands, you can see read the code as a series of statements: 1. Multiply cats\$age by 2, then 2. Take the square-root of these values, then 3. Round the result to the 3rd digit, then 4. Sort the values in ascending order

I encourage you to think of the %>% as short-hand for "then", when reading code that uses it!

"Pipe" operators are found in other languages; they get their name from the idea that your code can be thought of as a "pipeline".

Let's look at another example.

```
round(1.23456789, 3)
```

```
## [1] 1.235
```

We can use a pipe operator to acheive the same thing.

```
1.23456789 %>% round(3)
```

```
## [1] 1.235
```

The pipe takes care of making sure the output of the expression on the left-hand-side (a simple numeric, in this case) is inserted as the first argument of the expressing on the right-hand-side.

We can also pipe into other argument positions by using a period as a placeholder.

```
3 %>% round(1.23456789, .)
```

```
## [1] 1.235
```

These are contrived examples, and I don't suggest using pipes for simple operations like rounding. The pipes really become useful when chaining together multiple operations in sequence, as we'll do with our dplyr functions.

Combining Select & Filter

The pipe is really helpful when combined with the data-manipulation of dplyr. Remember how we used filter to select only the black cats? What if we only want to see the ID's of those cats, rather than all the info about them? We've already seen we can use select to pick out certain columns. We can use that to select the cat_id column from our filtered dataset like so

```
# reading from the inside out
select(filter(cats, coat == "black"), cat_id)

## cat_id
## 1  182
## 2  130
```

That might not look too bad now, but what if we wanted to do another operation on that output? We'd add another layer of nesting, and having to read that line from the inside-out can quickly become annoying. We can use the pipe operator to clean that up.

```
# reading from left to right
filter(cats, coat == "black") %>% select(cat_id)

## cat_id
## 1 182
## 2 130
```

We could even add another pipe to feed cats into filter; it isn't necessary, but it makes it even easier to see what we're operating on in this chain of commands. We'll combine this with some line breaks to really make this easy to read:

```
cats %>%
  filter(coat == "black") %>%
  select(cat_id)

## cat_id
## 1 182
## 2 130
```

summarize

While mutate creates new columns, it's often useful to summarize multiple rows into a single value. Say we want to find the mean weight of all these cats; enter summarize! Like mutate, the arguments to summarize (after the data.frame we want to operate on) are expressions. We can combine summarize with the mean function to get a mean weight for our collection of cats like so:

```
cats %>% summarize(mean_weight = mean(weight))

## mean_weight
## 1 5.8307
```

Notice how we have only a single value returned, but it's still in a data.frame format. This is subtle, but important; all these basic dplyr verbs take in data.frames and also return data.frames. This consistency helps make long chains of dplyr operations possible.

group_by

A very common data analysis task is to do operations like we did above, but to do them on a group-by-group basis. To do this with dplyr, we'll use the group_by function.

Let's look at the mean weights of our cats, grouping up by coat. This will give us the mean weight of the black cats, mean weight of the calico cats, etc. We can do this by inserting a group_by function into our earlier expression for computing mean weight:

```
cats %>%
  group_by(coat) %>%
  summarize(mean_weight = mean(weight))
```

```
## # A tibble: 5 x 2
##
     coat
             mean weight
##
                    <dbl>
     <fct>
## 1 black
                     5.67
## 2 brown
                     4.52
## 3 calico
                     6.19
## 4 maltese
                     9.68
## 5 tabby
                     3.93
```

Ta-da!

We can also use mutate on a per-group basis. Let's make a new column which centers our weights around zero; this can be done by subtracting the group's mean weight from each cat's weight:

```
cats %>%
  group_by(coat) %>%
  mutate(centered_weight = weight - mean(weight))
```

```
## # A tibble: 10 x 10
```

```
## # Groups:
                coat [5]
##
      street coat
                    sex
                            age weight fixed wander dist roamer cat id
##
      <fct>
             <fct> <fct> <dbl>
                                  <dbl> <int>
                                                     <dbl> <fct>
                                                                    <int>
##
    1 Los R~ tabby fema~
                           3.00
                                   3.99
                                            0
                                                     0.04 no
                                                                      321
    2 242 H~ malt~ fema~
                           8.23
                                 12.4
                                            1
                                                     0.033 no
                                                                      250
##
##
    3 201 H~ brown fema~
                           4.60
                                   3.95
                                            1
                                                     0.076 no
                                                                      219
                                                                      182
##
    4 140 R~ black fema~
                           7.17
                                   8.05
                                            1
                                                     0.03
                                                           no
    5 135 C~ cali~ male
                           4.66
                                            1
                                                     0.085 yes
                                                                      107
##
                                   6.19
##
    6 130 V~ tabby fema~
                           3.80
                                   3.86
                                            1
                                                     0.085 no
                                                                      234
    7 115 V~ brown male
                           6.92
                                            1
                                                     0.097 yes
                                                                      196
##
                                   5.63
    8 303 H~ brown male
                           3.71
                                   3.98
                                            1
                                                     0.033 no
                                                                      311
   9 16982~ black fema~
                           2.85
                                            0
                                                     0.065 no
                                   3.29
                                                                      130
## 10 16528~ malt~ fema~
                           4.59
                                   6.99
                                                     0.059 no
                                                                      349
## # ... with 1 more variable: centered_weight <dbl>
```

Going further

This is an introductory look at dplyr, just enough to make you dangerous. As you continue your R journey I suggest looking into the other awesome things you can do with this package!