Check-in 3

Again, the check-ins will review material presented in class but will also require you to think about new concepts, integrate across topics, and search for information. Some will be complex and take time to figure out. Feel free to work in groups on this.

Save this Rmd file in your checkins directory. Please submit the assignment as a pdf file on Canvas. Insert answers within the code chunks unless directed otherwise.

Problem 1: (7 pt) Inside the code chunk below, write a **single command** (using pipes) that assigns to the object dog_data the following in this order:

1. Imports stevens_etal_2020_obed_data1.csv from the following URL:

https://decisionslab.unl.edu/data/stevens_etal_2020_obed_data1.csv

- 2. Includes only the follow variables in this order: date, class, dog_sex, cgc_test, all four cort columns (using a helper function).
- 3. Relabels cgc_test to cgc.
- 4. Excludes observations where the dog's sex is missing.
- 5. Creates a new column called mean_cort that generates a per-observation mean over all of the cort columns (ignoring NAs).
- 6. Sorts the data by dog_sex then cort1.
- 7. Gives a glimpse() of the data.

Problem 2: (3 pts) For dog data, write a single command (using pipes) that does the following:

- 1. applies the log() function to all of the cort columns
- 2. returns observations only for female dogs who have either passed or failed the CGC (omit missing data).

Note for this and the remaining questions, do not assign your output to an object.

Problem 3: (4 pts) For dog data, write a single command (using pipes) that does the following:

- 1. returns only dogs who have cort values for all four time points
- 2. creates a new column called *cgc_complete* that converts their CGC outcome to either "Complete" if they passed or failed and "Incomplete" if there is missing data, placed after the *cgc* column
- 3. creates a new column called *cort_level* that categorizes *mean_cort* as "High" if 0.4 or above, "Medium" if between 0.3 and 0.4, and "Low" if 0.3 or below, placed after *mean_cort*
- 4. sorts by mean_cort from high to low.

Problem 4: (2 pts) For dog_data, write a single command (using pipes) that calculates the mean for each of the cort columns. *Hint: because dog_data is grouped row-wise, you need to ungroup() it before anything else.*

Problem 5: (2 pts) For dog_data, write a single command (using pipes) that calculates the mean, standard deviation, and sample size of the cort1 variable aggregated by dog sex and CGC status.