

# Bobcats, Coyotes & Foxes

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# 1 Introduction

This paper is for a midterm project dedicated towards replicating the results from a paper on a morphometric modeling approach to distinguishing among bobcat, coyote, and gray fox scats. This midterm project requires determining the best approach for addressing research questions as shown in Section 1.2.

Understanding the distribution and behavior of various species is fundamental to ecology, as it helps ecologists understand the relationship between organisms and their environments. There is a lot ecological significance behind tracking such species, as researchers can see precisely when individual animals depart from one location and arrive at another. Such insights reveal a individual animal's seasonal movements, feeding locations and more. Such information is valuable because the more ecologists know about animals' seasonal usage of habitats, the more they can protect the areas the animals need to survive.

Given that adult Gray Foxes have a greater average body length than adult Bobcats, and a smaller average body length than adult Coyotes, we hypothesize that Gray Fox scat will exhibit a significantly larger average diameter than Bobcat scat, but a smaller average diameter than Coyote scat.

## 1.1 Background Information

Conservation ecologists monitor species populations to assess the health of the ecosystem and just how effective conservation strategies are. While there are several methods for estimating population sizes, such as mark-recapture and camera-trap surveys, an alternative approach to this is analyzing biological remnants such as scat, as performed by Dr. Reid.

In a 2015 study by Dr. Rachel Reid, Reid had investigated whether morphological, biogeochemical and contextual traits distinguish between bobcat, coyote and fox scat. Morphological traits are the most cost-effective for field identification, while biogeochemical traits require laboratory analysis.

## 1.2 The Research Questions

The purpose of this paper is to analyze the data to evaluate whether specific morphological or biogeochemical traits can reliably differentiate species. We also research biological and ecological factors that explain the observed patterns. The research questions are listed as follows:

- Which (if any) morphological and biogeochemical traits distinguish between originating species of the scat samples?
- Why do you think those traits differ across species?

## 1.3 The Diet, Habitat & Physical Characteristics of Bobcats

Bobcats are mostly carnivorous. Their diet consists of a variety of animals, such as rabbits, rodents, wood rats birds, insects and many more. They also will consume plant material such as grass. Bobcats will also hunt pets or small livestock such as chicken if they're not kept in a secure enclosure. Bobcats can be found in diverse habitats throughout California. Suitable Bobcat habitat includes vegetation types, brushy stages of low and mid-elevation conifer, forests and desert environments. Interestingly, Bobcats prefer areas with dense bush cover. When it comes to physical characteristics, Bobcats are medium-sized cats with muscular bodies. They weight between 12 and 25 pounds. Bobcats have a round face with ruffs of fur on the side of the head. They have pointed ears, and appear to be approximately one quarter of the size of a mountain lion.

## 1.4 The Diet, Habitat & Physical Characteristics of Coyotes

The diet of a coyote consists mainly only mice, rats, ground squirrels, gophers, rabbits and carrion. They also eat insects and birds. In rural areas of California, they prey heavily on sheep, cattle and poultry. In urban and suburban areas, domestic cats, dogs and other pets can be food items. Coyotes can be found in the hotter drier regions of California. They can also be found in mountainous or humid areas in the state. Some physical characteristics of the coyote are that they are medium sized animals that belong to the Dog family. They can weigh between 22 to 25 pounds on average, with males being the larger sex. They also have large erect ears, slender muzzle and a bushy tail. Lastly, they have a distinctive voice and are proficient predators.

## 1.5 The Diet, Habitat & Physical Characteristics of Gray Foxes

The diet of a gray fox consists mainly of small rodents, birds and berries. They will also eat insects, eggs, acorns and fungi. Gray foxes can be found in populating coastal or mountain forests at lower elevations. They rarely dig their own dens, but will instead rest in crevices, under boulders and in hollow logs. Some physical characteristics of Gray Foxes are that it has short legs, a silvery-gray coat with patches of yellow, brown, rust, or white on the throat and the belly. Black tipped guard hairs form a dark line down its back to the tip of the tail.

## 1.6 Population Estimation Methods for Conservation

Next, we look at the different population estimation methods for conservation. We also look at their strengths and weaknesses.

### 1.6.1 Mark & Recapture Benefits and Drawbacks

For mobile organisms, we use a method called Mark & Recapture. This method involves marking a sample of captured animals and then releasing them back into the environment to allow them to mix with the rest of the population. A common issue with mark-recapture methods is that the process of capturing and marking the animals changes their behavior. This is otherwise known as a trap response.

### 1.6.2 Quadrats Benefits and Drawbacks

A quadrant is a frame used in ecology to isolate a standard unit of area for the study of the distribution of an item over a large area. Quadrats typically occupy an area of  $0.25\text{ m}^2$ , and are traditionally square. Some benefits it is quick, inexpensive and portable. Some of the disadvantages are it is not very accurate and the sample can seem unintentionally biased.

### 1.6.3 Transects Benefits and Drawbacks

A transect is a straight line that cuts through a natural landscape so that standardized observations and measurements can be made. Some advantages of using transects are it is quick, inexpensive and portable. A disadvantage of transects is that it is often used in inappropriate situations, so care must be taken when deciding whether or not to use a transect.

## 1.7 Dataset Description

With regards to the dataset used for performing data pre-processing, statistical and visualization techniques, it consists of 110 observations and 19 features. The features are of two different data types, such as `categorical`, `int64` and `float64`. This is a small dataset, and some of the features have null values, which are `scat_taper_mm`, `scat_taper_index`, `scat_mass_grams`, `d13C`, `d15N` and `CN Ratio`.

## 2 Methods

In this section, we discuss how we cleaned the dataset, including handling of missing values, data type conversions, and any transformations that we applied. We will also justify any decisions made about outlier treatment (removal, transformation or retention). Moreover, we look at the different statistical and visualization techniques we've used in this paper and why such approaches were appropriate.

### 2.1 Data Type Conversion

Looking at the dataset, we needed to understand what the features were about. We needed to know what the units were, what the features mean, as well as the description of the features. We renamed the columns to indicate some clarity as to what the feature was about, and we also renamed some numerical features to indicate what the unit of that feature was. After using `.info()` to look at the data type of each feature, we could see that some of the features were incorrectly represented as `int64` or `object`. For features that have binary values, or features that were missrepresented as `object` we converted them into `category` using `.astype()`.

### 2.2 Imputing or Dropping?

When we looked at the size of the dataset, we saw that it was 19 features in total and about 110 observations. In the context of machine learning applications, such a dataset is too small to drop any observations that have any missing values. Instead, we chose to impute. When it came to performing imputations, we needed to decide between the mean and median method if imputation for feature. The reasoning behind each method of imputation is as follows:

- **Mean:** we use the mean for numerical features that are normally distributed
- **Median:** we use the median for numerical features that are skewed and contain outliers

In order to figure out whether a feature is normally distributed or skewed, we need to use data visualization. For situation, we can plot a kernel density estimation plot for each of the features as well as box plot.

### 2.3 Visualization Tehcniques

When it came to performing visualization techniques, we had to set a response variable, which was `scat_species`. Given that the response variable was categorical, this means we performed our visualizations according to the following rules:

- **Categorical vs Numerical:** for plotting a categorical feature against numerical features, we explored using either box plots or violin plots
- **Categorical vs Categorical:** for plotting a categorical feature against categorical features, we explored using heat maps

The visualizations were performed as we compared the response variable against features from both morphological and biogeochemical traits.

### 2.4 Statistical Testing Tehcniques

When it came to performing statistical testing, we used our previously set response variable which was categorical against a number of categorical and numerical features, this means we performing our statistical testing according to the following rules:

- **Kruskal-Wallis:** for comparing a categorical variable with multiple values against a numerical variable, it answers the question of whether atleast one group defined by the categorical variable differs from at-least one other.
- **$\chi^2$  Test of Independence:** it is a hypothesis test for comparing two categorical variables and measures whether there is a 'dependence' between them
- **Dunn Testing:** used as a post-hoc analysis after a Kruskal-Wallis test to identify which specific groups are significantly different from each other.

The statistical tests were performed as we compared the response variables against features from both morphological and biogeochemical traits. The justification for all statistical tests and visualizations were all dependent on the type of variable we were dealing against the response variable, and this will be explained in further detail in Section 3.

### 2.5 Family-Wise Error Rate

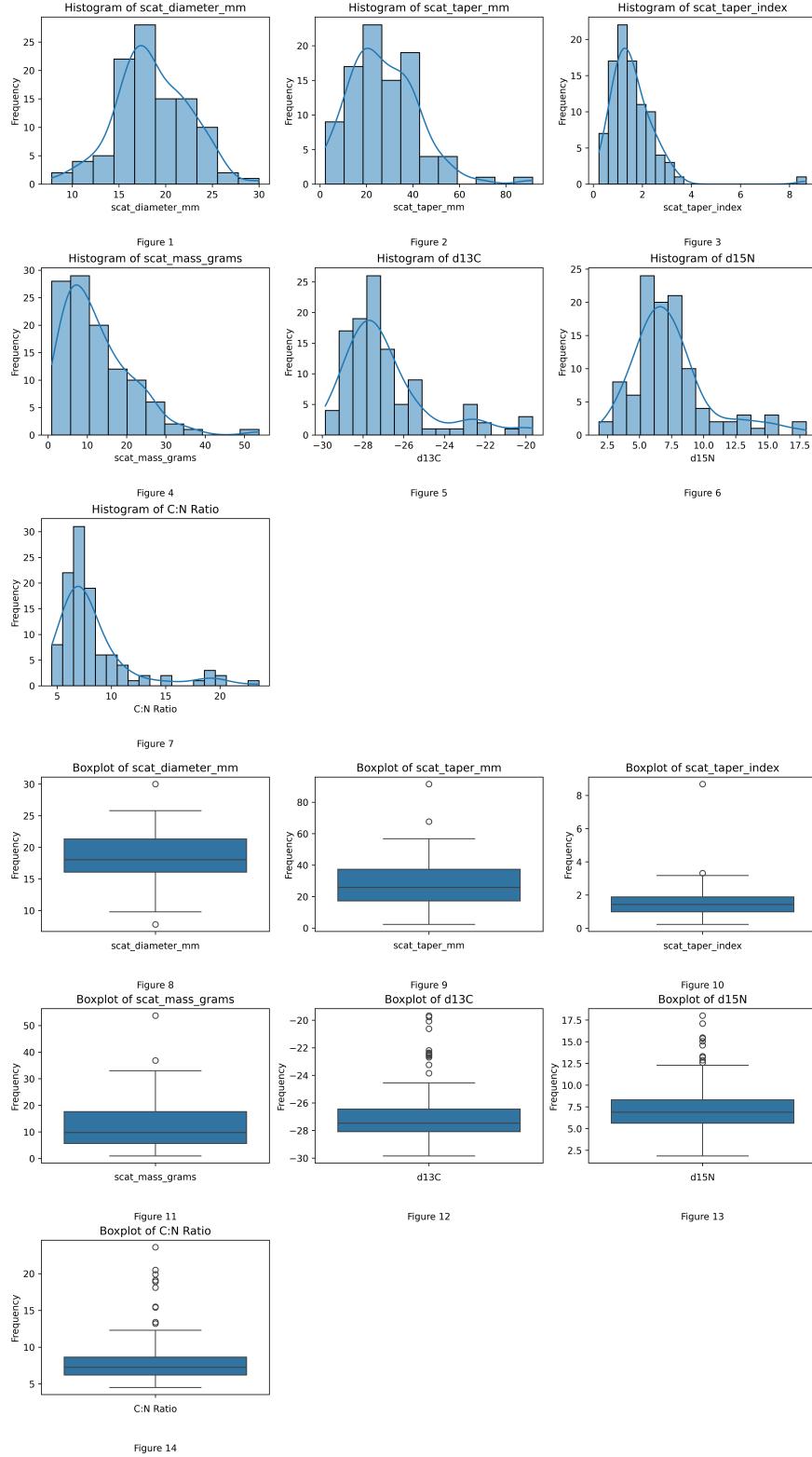
When it came to determining the family-wise error rate method for each of the tests, we decided to go with bonferroni correction. It is a multiple comparision adjustment method that reduces the significance level to control for false positives when performing multiple hypothesis tests. The justification behind using it is that we would like to reduce the likelihood of false positives when performing multiple comparisions.

### 3 Results

In this section, we discuss the different visualization techniques and statistical testing methods that were performed to evaluate the relationships between the morphological and biogeochemical traits and species. We also determine which traits can be used to distinguish atleast one species from the rest.

#### 3.1 Histograms & Boxplots

In this page, we display all the histograms and box plots of all features that have missing values so that we can justify the method of imputation.



## 3.2 The Feature Description Table

In this section, we created a table that categorizes each variable in the data set as morphological, biogeochemical, contextual or not a trait.

Feature Names	Feature Types	Feature Names	Feature Types
scat_species	Not a Trait	scat_age	Morphological
scat_month	Contextual	scat_number	Morphological
scat_year	Contextual	scat_length_cm	Morphological
scat_site	Contextual	scat_diameter_mm	Morphological
scat_location	Contextual	scat_taper_mm	Morphological
scat_taper_index	Morphological	scat_mass_grams	Morphological
d13C	Biogeochemical	scat_ropey	Morphological
d15N	Biogeochemical	scat_segmented	Morphological
C:N Ratio	Biogeochemical	scat_flat	Morphological
scat_scrape	Morphological		

Table 1: A Categorized Table of Features

## 3.3 Imputations of Missing Value

In this section, we look at the kernel density estimation plots and the box plots to justify our reasons for the method of imputation between mean or median for each of the features that have missing values.

### 3.3.1 Method of Imputation for scat\_diameter\_mm

For this feature, if we were to look at Figure 1, we can see that the histogram has a unimodal peak, and the peak is at the center of the histogram. This also means that there is no skew at all. In Figure 8, we can see a box plot that has very few outliers and shows no skew at all too. As a result, we can say that the feature is normally distributed, hence it should have imputations using the mode.

### 3.3.2 Method of Imputation for scat\_taper\_mm, scat\_taper\_index, scat\_mass\_grams, d13C, d15N & C:N Ratio

For these following features, if we were to look at figures 2 to 7, we can see that each of them have a histogram with a single peak, making it unimodal, and those peaks are shifted to the left, which means that the histograms are skewed to the right. Furthermore, when we look at the boxplots from figures 9 to 14, we can evidently see the skew, and we can also see the huge number of outliers in figures 12 to 14. As a result, we can impute the missing values of these features with the median.

### 3.4 Numerical Morphological Features Box Plots, Kruskal-Wallis & Dunn Testing

In this section, we look at all of the box plots, and we justify the use of such a visualization, as well as using Kruskal-Wallis and Dunn Testing as post-hoc. We justify the use of box plots because we are comparing two variables which are categorical and numerical, hence a box plot visualization would be the best.

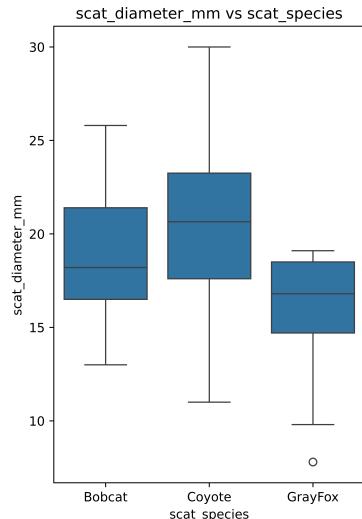


Figure 15

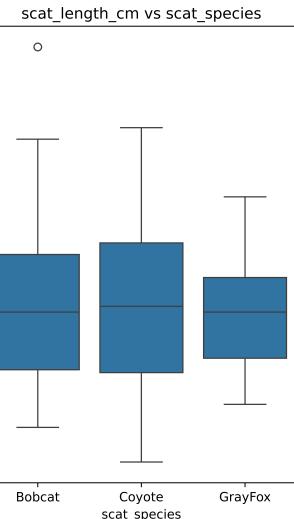


Figure 16

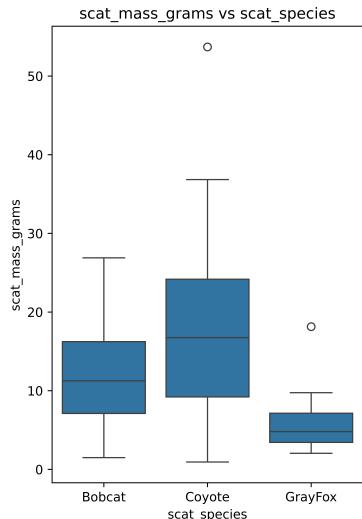


Figure 17

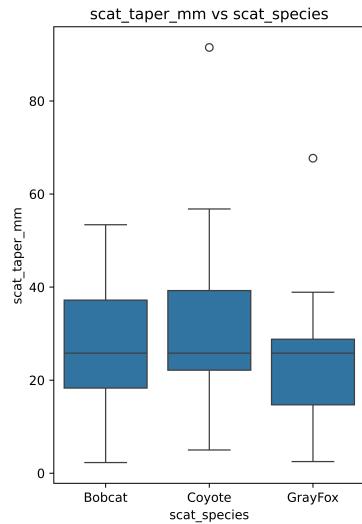


Figure 18

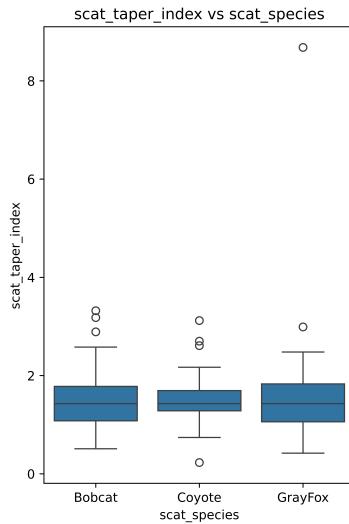


Figure 19

### 3.4.1 scat\_diameter\_mm vs scat\_species

When we look at figure 15, and compare the three box plots of Bobcat, Coyotes & Foxes, we can evidently say that the medians between Gray Fox is different compared to Bobcat & Coyote, which suggests that the central tendency of the `scat_diameter_mm` is different across all three species. Furthermore, judging by the IQR spread of Bobcats & Coyotes, we can say that they both have more variability than Gray Foxes. It is worth noting that none of the boxplots have any outliers, which means that the data within those species is consistent. For Bobcats, we can say that the box plot has a positive skew, and for Coyotes, we can say that the box plot has no skew, and for Gray Foxes, the box plot has a negative skew.

With regards to statistical testing, it would be the most appropriate to use Kruskal-Wallis because `scat_species` consists of three values versus a numerical variable, which is `scat_diameter_mm`. After performing the Kruskal-Wallis test, we found that the test significance was true, which had meant that we rejected the null hypothesis and concluded that there is a statistically significant difference between atleast one of the groups. When performing the Kruskal-Wallis test, it is important to note that we had applied Bonferroni Correction to reduce the chances of any Type I error.

After performing the statistical testing, we moved on to performing a post-hoc test, where we used Dunn's test to find out which specific groups are different, and the results of the test are shown below:

Comparison	Bobcat	Coyote	GrayFox
Bobcat	1.000	0.208	0.001
Coyote	0.208	1.000	0.000
GrayFox	0.001	0.000	1.000

Table 2: Dunn's Test p-values for `scat_diameter_mm` between Species

Looking at the results of the table above, we can say that the Bobcats & Coyotes don't show a significant difference in scat diameter. Bobcats & Gray Foxes show a significant difference in scat diameter and Coyotes & Gray Foxes also show a significant difference in diameter. In conclusion, we can say that the scat diameter of Gray Foxes is different from both Bobcats & Coyotes, but there is no significant difference between Bobcats & Coyotes.

### 3.4.2 scat\_length\_cm vs scat\_species

When we look at figure 16, and compare the three box plots of Bobcat, Coyotes & Gray Foxes, we can evidently say that the medians between all three are very similar, which suggests that the central tendency of the `scat_length_cm` has little to no difference in the central values of the numerical variable across all categories. The IQR of Gray Fox is smaller than Bobcats & Coyotes, which suggests that Gray Foxes have less variability than the other two. It is also worth noting that there are very few outliers in the box plots, which means that the data is consistent. It also appears that all box plots have no skew, making them symmetrical.

With regards to statistical testing, it would be the most appropriate to use Kruskal-Wallis because `scat_species` consists of three values versus a numerical variable, which is `scat_length_cm`. After performing the Kruskal-Wallis test, we found that the test significance was false. Which had meant that we had accepted the null hypothesis and concluded that there is no statistically significant difference between atleast one of the groups. When performing the Kruskal-Wallis test, it is important to note that we had applied bonferroni correction to reduce the chances of any Type I error.

### 3.4.3 scat\_mass\_grams vs scat\_species

When we look at figure 17, and compare the three box plots of Bobcat, Coyotes & Gray Foxes, we can see that the medians of the box plots of Bobcat & Coyotes are similar, however the median of the Gray Fox is different from the rest. The IQR of the Bobcats & Coyotes are larger than the IQR of Gray Foxes, which means that it has more variability. All three box plots have very few outliers, which means that the data is consistent across all three. It would also be fair to assume that there is no skew in all three box plots, as they appear to be symmetrical.

With regards to statistical testing, it would be the most appropriate to use Kruskal-Wallis because `scat_species` consists of three values versus a numerical variable, which is `scat_mass_grams`. After performing the Kruskal-Wallis test, we found that the significance test was true. Which had meant that we had rejected the null hypothesis and concluded that there is a statistically significant difference between atleast one of the groups. When performing the Kruskal-Wallis test, it is important to note that we had applied Bonferroni Correction to reduce the chances of any Type I error.

After performing the statistical testing, we moved on to performing a post-hoc test, where we used Dunn's test to find out which specific groups are different, and the results of the test are shown below:

Comparison	Bobcat	Coyote	GrayFox
Bobcat	1.000	0.078	0.000
Coyote	0.078	1.000	0.000
GrayFox	0.000	0.000	1.000

Table 3: Dunn's Test p-values for scat\_mass\_grams between species

Looking at the results of the table above, we can say that there is no significant difference in scat mass between Bobcats & Coyotes. We can also say that there is a significant difference between Bobcats & Gray Foxes, and there is a significant difference between Coyotes & Gray Foxes. There is no difference in scat mass between Bobcats & Coyotes. As a result, it is the Gray Foxes that has a significant difference in scat mass with other species.

#### 3.4.4 scat\_taper\_mm vs scat\_species

When we look at figure 18, and compare the three box plots of Bobcat, Coyotes & Gray Foxes, we can see that the medians of the box plots of all three are similar. The IQR of all three box plots are similar, which suggests that all three categories have similar variability. All three box plots also have very few outliers, which means that the data is consistent across all three. While the Bobcat box plot has no skew, we can deduce that the Coyote box plot has a positive skew, and the Gray Fox box plot has a negative skew.

With regards to statistical testing, it would be the most appropriate to use Kruskal-Wallis because `scat_species` consists of three values versus a numerical variable, which is `scat_taper_mm`. After performing the Kruskal-Wallis test, we found that the significance test was false. Which had meant that we had accepted the null hypothesis and concluded that there no statistically significant difference between atleast one of the groups. When performing the Kruskal-Wallis test, it is important to note that we had applied Bonferroni Correction to reduce the chances of any Type I error.

#### 3.4.5 scat\_taper\_index vs scat\_species

When we look at figure 19, and compare the three box plots of Bobcat, Coyotes & Gray Foxes, we can see that the medians of the box plots of all three are similar. The IQR of all three box plots are similar, although the IQR of the Coyote box plots seems a little smaller, however, this overall indicates that all three categories have similar variability. All three box plots have few outliers, which means that the data is consistent across all three categories. It can also be seen that all three box plots have no skew.

With regards to statistical testing, it would be the most appropriate to use Kruskal-Wallis because `scat_species` consists of three values versus a numerical variable, which is `scat_taper_index`. After performing the Kruskal-Wallis test, we found that the significance test was false. Which had meant that we had accepted the null hypothesis and concluded that there no statistically significant difference between atleast one of the groups. When performing the Kruskal-Wallis test, it is important to note that we had applied Bonferroni Correction to reduce the chances of any Type I error.

### 3.5 Numerical Biogeochemical Features Box Plots, Kruskal-Wallis & Dunn Testing

In this section, we look at all of the violin plots, and we justify the use of such a visualization, as well as using Kruskal-Wallis and Dunn Testing as post-hoc. We justify the use of violin plots because we are comparing two variables which are categorical and numerical, hence a violin plot visualization would be the best.

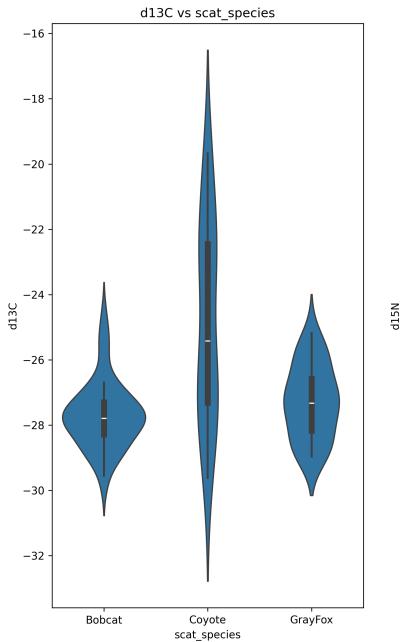


Figure 24

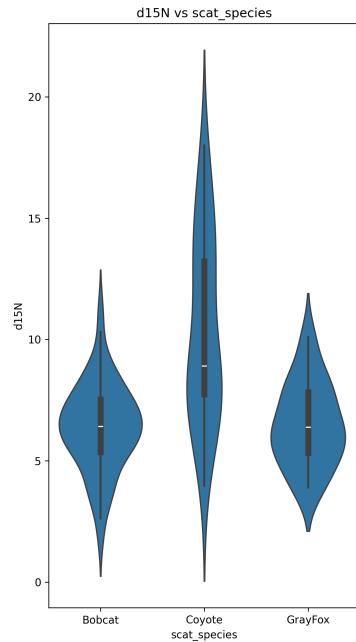


Figure 25

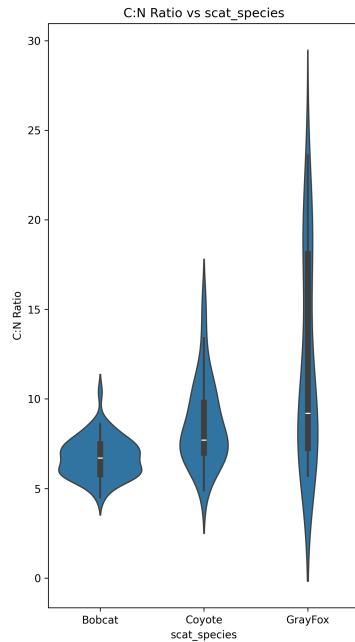


Figure 26

#### 3.5.1 d13C vs scat\_species

When we look at figure 24, and compare the three violin plots of Bobcat, Coyotes & Gray Foxes, the medians of all three violin plots are different, however the median of the Coyote violin plot is very different from the rest. When it comes to comparing the IQR of all three violin plots, we can deduct that the Coyote violin plot has the most spread, compared the IQR of both Bobcat & Gray Fox, which appear to be similar. It can be seen that the violin plot of Bobcat, has a left skew, the Coyote violin plot has no skew, and the Gray Fox violin plot has no skew either.

With regards to statistical testing, it would be the most appropriate to use Kruskal-Wallis because `scat_species` consists of three values versus a numerical variable, which is `d13C`. After performing the Kruskal-Wallis test, we found that the significance test as true. Which has meant that we have rejected the null hypothesis and concluded that there is a statistically significant difference between atleast one of the groups. When performing the Kruskal-Wallis tests, it is important to note that we had applied the Bonferroni Correction to reduce the chances of any Type I error.

After performing the statistical testing, we moved on to performing a post-hoc test, where we used Dunn's test to find out which specific groups are different, and the results of the test are shown below:

Comparison	Bobcat	Coyote	GrayFox
Bobcat	1.000	0.000	0.160
Coyote	0.000	1.000	0.010
GrayFox	0.160	0.010	1.000

Table 4: Dunn's Test p-values for d13C between species

Looking at the results of the table above, we can confirm that there is a significant difference between Bobcats & Coyotes, as well as between Bobcats & Gray Foxes, however, there is no significant difference between Coyotes & Gray Foxes. As a result, there is significant difference between Bobcats & Coyotes and Coyotes & Gray Foxes.

#### 3.5.2 d15N vs scat\_species

When we look at figure 25, and compare the three violin plots of Bobcat, Coyotes & Gray Foxes, the medians of Bobcats & Gray Foxes are the same, however the median of the Coyote is different from others. When it comes to comparing the IQR of all three violin plots, we can deduct that the IQR of the Bobcat & Gray Fox are similar, but there is a larger spread in the violin plot of Coyotes because it has a larger IQR. It can be seen that there is no skew in both Bobcat & Gray Fox violin plots, however, there is right skew in the Coyote violin plot.

With regards to statistical testing, it would be the most appropriate to use Kruskal-Wallis because `scat_species` consists of three values versus a numerical variable, which is `d15N`. After performing the Kruskal-Wallis test, we found that

the significance test as true. Which has meant that we have rejected the null hypothesis and concluded that there is a statistically significant difference between atleast one of the groups. When performing the Kruskal-Wallis tests, it is important to note that we had applied the Bonferroni Correction to reduce the chances of any Type I error.

After performing the statistical testing, we moved on to performing a post-hoc test, where we used Dunn's test to find out which specific groups are different, and the results of the test are shown below:

Comparison	Bobcat	Coyote	GrayFox
Bobcat	1.000	0.000	0.989
Coyote	0.000	1.000	0.000
GrayFox	0.989	0.000	1.000

Table 5: Dunn's Test p-values for d15N between species

Looking at the results of the table above, we can confirm that there is a significant difference between Bobcats & Coyotes. There is no signifancet difference between Bobcats & Gray Foxes, however, there is significant difference between Coyotes & Gray Foxes. As a result, we can summarize that Coyotes, have a significant difference between Coyotes & Gray Foxes.

### 3.5.3 C:N Ratio vs scat\_species

When we look at figure 26, and compare the three violin plots of Bobcat, Coyotes & Gray Foxes, the medians of Coyotes & Gray Foxes are all different, however the median of Bobcats are very different compared to the other two. When it comes to comparing the IQR of all three violin plots, we can dedcut that the IQR of the Gray Foxes are the highest, which means it has a larger spread compared to the IQR of the other two violin plots. It can be seen that there is no skew in the Bobcat violin plot, however, there is a positive skew in violin plots of both Coyotes & Gray Foxes.

With regards to statistical testing, it would be the most appropriate to use Kruskal-Wallis because `scat_species` consists of three values versus a numerical variable, which is C:N Ratio. After performing the Kruskal-Wallis test, we found that the significance test as true. Which had meant that we have rejected the null hypothesis and concluded that there is a statistically significant difference between atleast one of the groups. When performing the Kruskal-Wallis tests, it is important to note that we had applied the Bonferroni Correction to reduce the chances of any Type I error.

After performing the statistical testing, we moved on to performing a post-hoc test, where we used Dunn's test to find out which specific groups are different, and the results of the test are shown below:

Comparison	Bobcat	Coyote	GrayFox
Bobcat	1.000	0.000	0.000
Coyote	0.000	1.000	0.182
GrayFox	0.000	0.182	1.000

Table 6: Dunn's Test p-values for C:N Ratio between species

Looking at the results of the table above, we can confirm that there is a significant difference in terms of C:N Ratio between Bobcats & Coyotes, as well as Bobcats & Gray Foxes. However, there is no significant difference between Coyotes & Gray Foxes. As a result, we can summarize that Bobcats have a significant difference with Coyotes & Gray Foxes.

### 3.6 Categorical Morphological Features Heat Maps, Chi-Squared Test & Pairwise Chi-Squared Tests

In this section, we look at all of the heat maps, and we justify the use of such a visualization, as well as using Chi-Squared Test and Chi-Squared Test as post-hoc. We justify the use of heatmaps because we are comparing two variables which are categorical, hence a heatmap visualization would be the best.

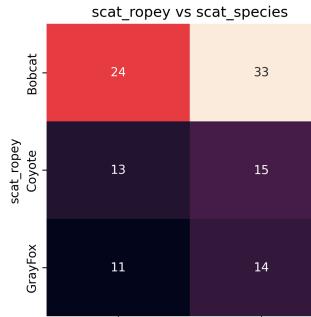


Figure 20 scat\_scrape vs scat\_species

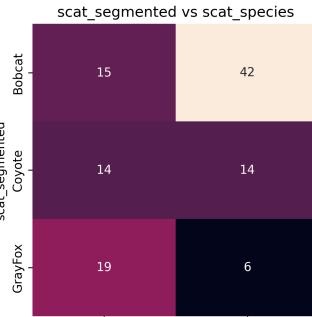


Figure 21

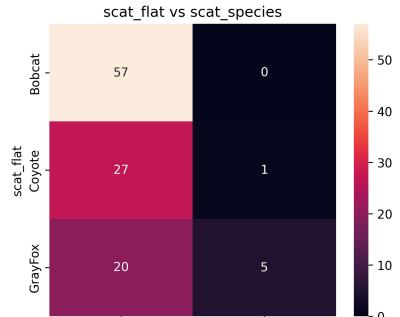


Figure 22

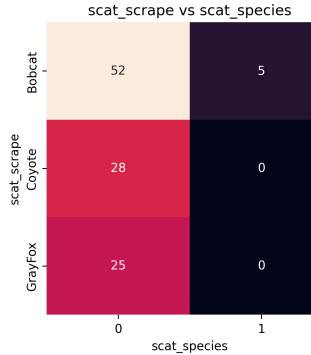


Figure 23

#### 3.6.1 scat\_ropey vs scat\_species

When we look at figure 20, it can be deduced that when it comes to the scat being ropey, it can be deduced that the number of scats being ropey or not is evenly divided across each species. It can also be deduced that there are a high number of bob cats who have and don't have scat that is ropey compared to other species.

With regards to statistical testing, it would be the most appropriate to use chi-squared because `scat_species` consists of three values versus a categorical variable `scat_ropey`. After performing the chi-squared test, we found that the significance test is false. Which had meant that we have accepted the null hypothesis in which there is no association at all between the two categorical variables.

#### 3.6.2 scat\_segmented vs scat\_species

When we look at figure 21, we can see that bobcats have a high number of scat that is segmented compared to bobcats who don't, while it is evenly divided when it comes to Coyotes. When it comes to Gray Foxes, it can be deduced that there are more Gray Foxes who don't have segmented scat. Overall, the heatmap also shows that there is a high number of bobcats who have segmented scat and gray foxes are the least number of species that have a segmented scat.

With regards to statistical testing, it would be the most appropriate to use chi-squared because `scat_species` consists of three values versus a categorical variable `scat_segmented`. After performing the chi-squared test, we found that the significance test is true. Which had meant that we have rejected the null hypothesis and that there is an association between the two categorical variables.

After performing the statistical testing, we moved on to performing a post-hoc test, where we used pair-wise chi-squared tests to find out which specific groups have an association, and the results of the pairwise tests are shown below:

Comparison	p-value
Coyote vs Bobcat	0.0547
Coyote vs GrayFox	0.0958
Bobcat vs GrayFox	0.0001

Table 7: Chi-squared test results for species comparisons

Looking at the results of the table above, we can see that there is no significant association between segmented scat of Bobcat & Coyotes. Also, we see that there is no significant association between segmented scat of Coyotes & Gray Foxes. However, there is a significant association between segmented scat of Bobcats & Gray Foxes.

### 3.6.3 scat\_flat vs scat\_species

When we look at figure 22, it can be seen that a majority of Bobcats, Coyotes & Gray Foxes don't have flat scat. The highest number of species that don't have a flat scat are Bobcats, and the lowest being Gray Foxes.

With regards to statistical testing, it would be the most appropriate to use chi-squared because `scat_species` consists of three values versus a categorical variable `scat_flat`. After performing the chi-squared test, we found that the significance test is true. Which had meant that we have rejected the null hypothesis and that there is an association between the two categorical variables.

After performing the statistical testing, we moved on to performing a post-hoc test, where we used pair-wise chi-squared tests to find out which specific groups have an association, and the results of the pairwise tests are shown below:

Comparison	p-value
Coyote vs Bobcat	0.7150
Coyote vs GrayFox	0.1470
Bobcat vs GrayFox	0.0029

Table 8: Chi-squared test results for species comparisons

Looking at the results of the table above, we can see that there is no significant association of flat scats between Coyotes & Bobcats, and the same goes for Coyotes & Gray Foxes. There is, however, significant association between Bobcats & Gray Foxes.

### 3.6.4 scat\_scrape vs scat\_species

When we look at figure 23, it can be again seen that a majority of Bobcats, Coyotes & Gray Foxes don't have scrape scat. The highest number of species that don't have a scrape scat are Bobcats, and the lowest being Gray Foxes.

With regards to statistical testing, it would be the most appropriate to use chi-squared because `scat_species` consists of three values versus a categorical variable `scat_scrape`. After performing the chi-squared test, we found that the significance test is false. Which had meant that we have accepted the null hypothesis in which there is no association at all between the two categorical variables.

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