

# Biodiversity Project

## 1 Introduction

The goal of this project is to analyze biodiversity data from the National Parks Service, particularly around various species observed in different national park locations.

This project will scope, analyze, prepare, plot data, and seek to explain the findings from the analysis.

Here are a few questions that this project has sought to answer:

- What is the distribution of conservation status for species?
- Are certain types of species more likely to be endangered?
- Are the differences between species and their conservation status significant?
- Which animal is most prevalent and what is their distribution amongst parks?

### Data sources:

Both `Observations.csv` and `Species_info.csv` was provided by <https://www.codecademy.com/>.

The data for this project is *inspired* by real data.

### 1.1 Import Python Modules

Here are the primary modules that will be used in this project:

```
[105]: import pandas as pd
import numpy as np
from matplotlib import pyplot as plt
import seaborn as sns

%matplotlib inline
```

### 1.2 Getting data to know

**species** The `species_info.csv` contains information on the different species in the National Parks. The columns in the data set include: - **category** - The category of taxonomy for each species - **scientific\_name** - The scientific name of each species - **common\_names** - The common names of each species - **conservation\_status** - The species conservation status

```
[147]: species = pd.read_csv('species_info.csv', encoding='utf-8')
print("Columns:", species.columns)
```

```
species.head().style.set_caption("Species DataFrame Head").  
↳background_gradient(cmap='viridis')
```

```
Index(['category', 'scientific_name', 'common_names', 'conservation_status'],  
      dtype='object')
```

```
[147]: <pandas.io.formats.style.Styler at 0x12f2b2cf0>
```

**observations** The `Observations.csv` contains information from recorded sightings of different species throughout the national parks in the past 7 days. The columns included are:

- **scientific\_name** - The scientific name of each species
- **park\_name** - The name of the national park
- **observations** - The number of observations in the past 7 days

```
[172]: observations=pd.read_csv('observations.csv',encoding='utf-8')  
print("Columns:",observations.columns)  
observations.head().style.set_caption("Conservations DataFrame Head").  
↳background_gradient(cmap='viridis')
```

```
Columns: Index(['scientific_name', 'park_name', 'observations'], dtype='object')
```

```
[172]: <pandas.io.formats.style.Styler at 0x1414f0230>
```

**Data Characteristics** Dimensions of the data sets, for `species` there are 5,824 rows and 4 columns while `observations` has 23,296 rows and 3 columns.

```
[150]: print(f"species shape: {species.shape}")  
print(f"observations shape: {observations.shape}")
```

```
species shape: (5824, 4)  
observations shape: (23296, 3)
```

## 2 Exploring data

Exploring the `species` data a little more in depth. Finding the number of distinct species in the data. Column `scientific_name` have 5,541 unique species. There seems to be a lot of species in the national parks!

```
[170]: print(f"number of species:{species.scientific_name.nunique()}")
```

```
number of species:5541
```

```
Unique categories
```

```
[158]: print(f"nnumber of categories:{species.category.nunique()}")  
print(f"categories:{species.category.unique()}")
```

```
nnumber of categories:7  
categories:['Mammal' 'Bird' 'Reptile' 'Amphibian' 'Fish' 'Vascular Plant'  
           'Nonvascular Plant']
```

Amount of each category

```
[161]: species.groupby("category").size()
```

```
[161]: category
      Amphibian           80
      Bird             521
      Fish             127
      Mammal           214
      Nonvascular Plant  333
      Reptile           79
      Vascular Plant    4470
      dtype: int64
```

Combining both tables together and create one table depending on their scientific name

```
[174]: combined = pd.merge(observations, species, on='scientific_name', how='left')
      combined.head()
      print("Columns:", combined.columns)
```

```
Columns: Index(['scientific_name', 'park_name', 'observations', 'category',
               'common_names', 'conservation_status'],
              dtype='object')
```

Lets find out Least observed and Most observed animal

```
[167]: least_obs_idx = combined['observations'].idxmin()
      most_obs_idx = combined['observations'].idxmax()

      least_observed = combined.loc[least_obs_idx]
      most_observed = combined.loc[most_obs_idx]

      print("Least Observed Animal:")
      print(least_observed[['common_names', 'scientific_name', 'observations',
                             ↪ 'park_name', 'category', 'conservation_status']])
      print("\nMost Observed Animal:")
      print(most_observed[['common_names', 'scientific_name', 'observations',
                             ↪ 'park_name', 'category', 'conservation_status']])
```

Least Observed Animal:

```
common_names      Golden Corydalis, Scrambled Eggs
scientific_name    Corydalis aurea
observations              9
park_name          Bryce National Park
category           Vascular Plant
conservation_status      NaN
Name: 10368, dtype: object
```

Most Observed Animal:

```
common_names      Deep-Root Clubmoss, Ground Cedar
```

```

scientific_name      Lycopodium tristachyum
observations          321
park_name            Yellowstone National Park
category             Vascular Plant
conservation_status   NaN
Name: 12447, dtype: object

```

Lets check conservation status to find out is there any endangered species out there, and what kind of conservation status is existing

```

[181]: not_nan_conservation = combined[combined['conservation_status'].notna()]

print("Number of rows with conservation_status values:", not_nan_conservation.
      ↪shape[0])

print("Unique conservation statuses:",
      ↪not_nan_conservation['conservation_status'].unique())

```

```

Number of rows with conservation_status values: 880
Unique conservation statuses: ['Species of Concern' 'Threatened' 'Endangered'
                              'In Recovery']

```

The column has 4 categories, Species of Concern, Endangered, Threatened, In Recovery, and nan values.

Finding out exact amount of each conservation status

A lot of values is NaN so i convert them to No Intervention

```

[284]: combined.fillna('No Intervention', inplace=True)
combined.groupby("conservation_status").size()

```

```

[284]: conservation_status
Endangered          80
In Recovery         24
No Intervention     24752
Species of Concern   732
Threatened          44
dtype: int64

```

```

[ ]: Building a pie chart having this data including only info that we have

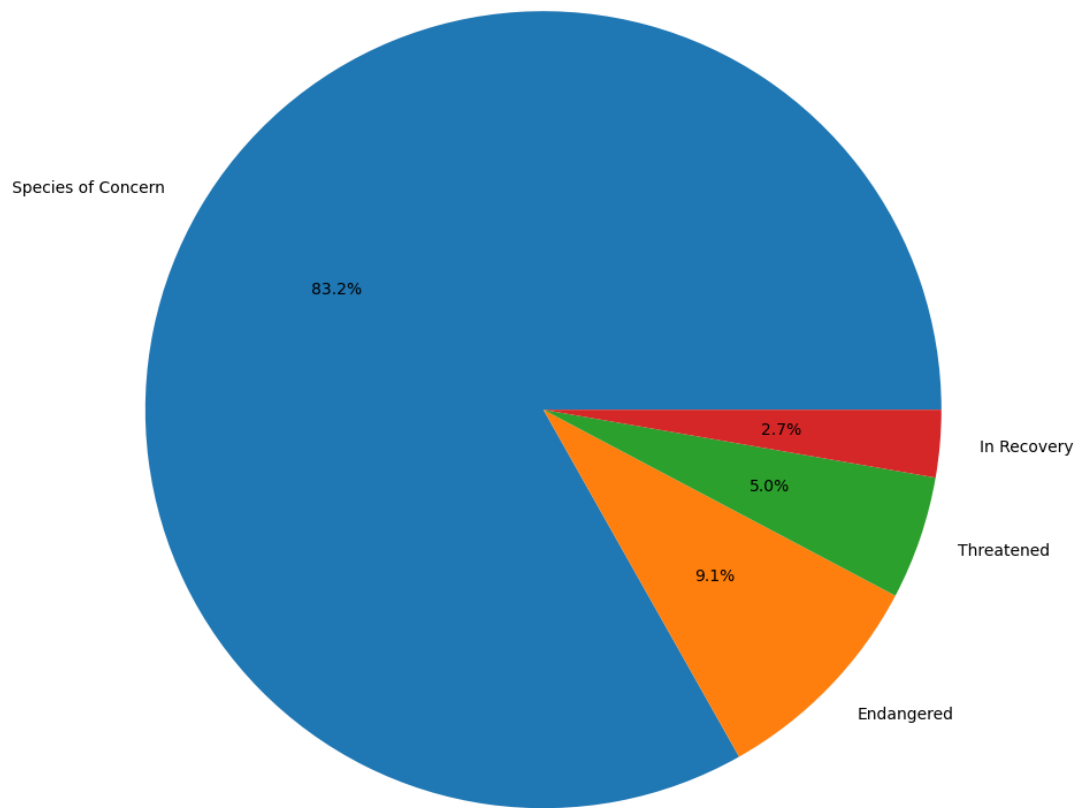
```

```

[186]: status_counts = not_nan_conservation['conservation_status'].value_counts()
plt.figure(figsize=(10, 8))
plt.pie(status_counts.values, labels=status_counts.index, autopct='%1.1f%%')
plt.title("Percentage Distribution of Conservation Status Values")
plt.axis('equal')
plt.tight_layout()
plt.show()

```

Percentage Distribution of Conservation Status Values



Finding out which category of animals in each conservation status

```
[286]: conservationCategory = combined[combined.conservation_status != "No Intervention"]\
        .groupby(["conservation_status", "category"])['scientific_name']\
        .count()\
        .unstack()
```

conservationCategory

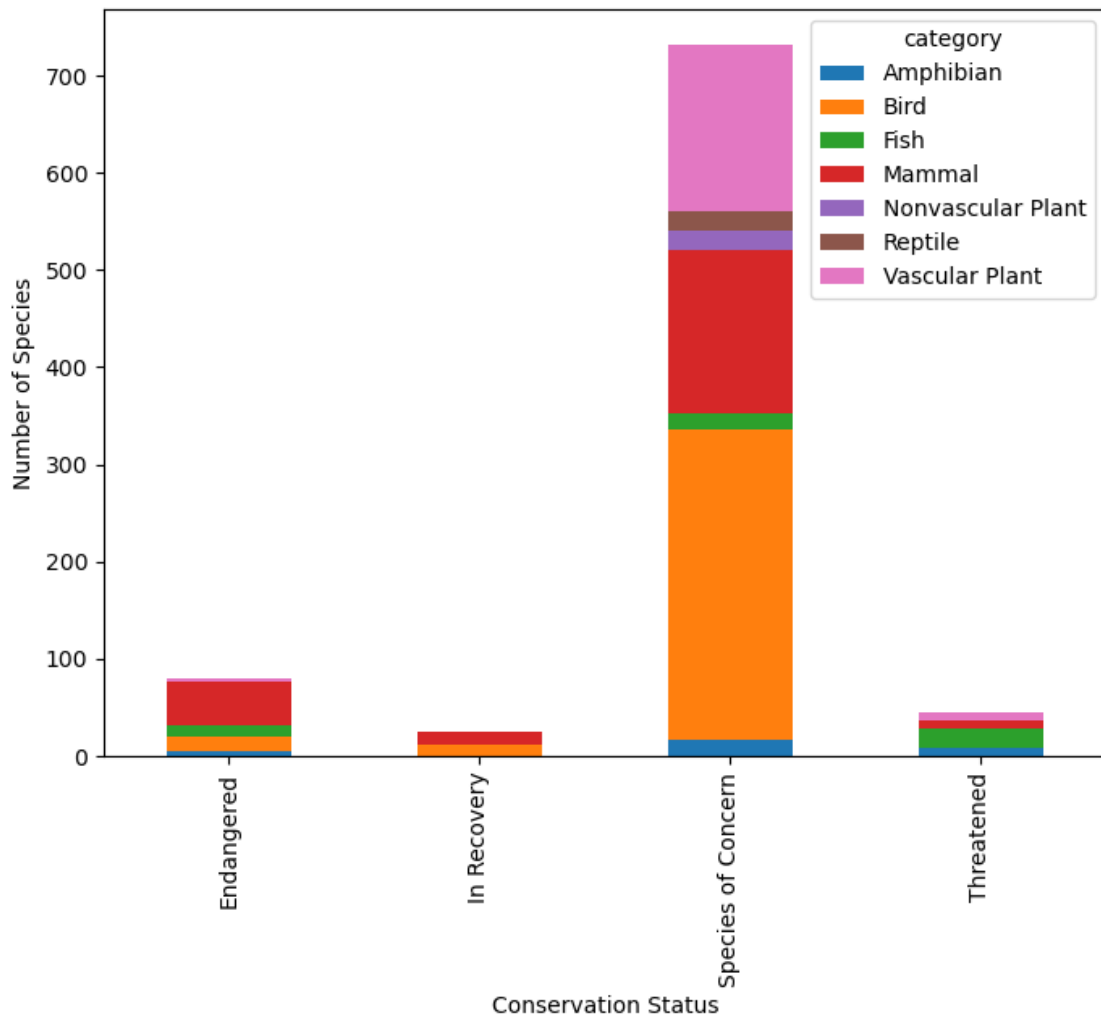
```
[286]: category      Amphibian   Bird   Fish   Mammal   Nonvascular Plant \
conservation_status
Endangered          4.0    16.0   12.0    44.0                NaN
In Recovery         NaN    12.0    NaN    12.0                NaN
Species of Concern  16.0   320.0   16.0   168.0               20.0
Threatened          8.0     NaN   20.0    8.0                NaN

category      Reptile   Vascular Plant
conservation_status
```

Endangered	NaN	4.0
In Recovery	NaN	NaN
Species of Concern	20.0	172.0
Threatened	NaN	8.0

Building stacked bar chart depending on this

```
[219]: ax = conservationCategory.plot(kind = 'bar', figsize=(8,6),stacked=True)
ax.set_xlabel("Conservation Status")
ax.set_ylabel("Number of Species")
plt.show();
```



Finding if certain types of species are more likely to be endangered. To do so I created new column called `is_protected` that will include any species that had value other than no No Intervention

```
[222]: combined['is_protected'] = combined.conservations_status != 'No Intervention'
```

Once the new column is created, group by `category` and `is_protected` to show the break down of each species type and protection status.

It's easy to see that Birds, Vascular Plants, and Mammals have a higher absolute number of species protected.

```
[228]: category_counts = combined.groupby(['category', 'is_protected'])\
        .scientific_name.nunique()\
        .reset_index()\
        .pivot(columns='is_protected',\
                index='category',\
                values='scientific_name')\
        .reset_index()\
category_counts.columns = ['category', 'not_protected', 'protected']

category_counts
```

```
[228]:
```

	category	not_protected	protected
0	Amphibian	72	7
1	Bird	413	75
2	Fish	115	11
3	Mammal	146	30
4	Nonvascular Plant	328	5
5	Reptile	73	5
6	Vascular Plant	4216	46

Calculating rate of protection

```
[233]: category_counts['percent_protected'] = category_counts.protected /\
        (category_counts.protected + category_counts.not_protected) * 100

category_counts
```

```
[233]:
```

	category	not_protected	protected	percent_protected
0	Amphibian	72	7	8.860759
1	Bird	413	75	15.368852
2	Fish	115	11	8.730159
3	Mammal	146	30	17.045455
4	Nonvascular Plant	328	5	1.501502
5	Reptile	73	5	6.410256
6	Vascular Plant	4216	46	1.079305

**Statistical Significance** This section will run some chi-squared tests to see if different species have statistically significant differences in conservation status rates. The first test will be called `contingency1` and will need to be filled with the correct numbers for mammals and birds.

The results from the chi-squared test returns many values, the second value which is 0.69 is the p-value. The standard p-value to test statistical significance is 0.05. For the value retrieved from this test, the value of 0.69 is much larger than 0.05. In the case of mammals and birds there doesn't

seem to be any significant relationship between them i.e. the variables independent.

```
[237]: from scipy.stats import chi2_contingency

contingency1 = [[30, 146],
                [75, 413]]
chi2_contingency(contingency1)
```

```
[237]: Chi2ContingencyResult(statistic=0.1617014831654557, pvalue=0.6875948096661336,
dof=1, expected_freq=array([[ 27.8313253, 148.1686747],
[ 77.1686747, 410.8313253]]))
```

The next pair, is going to test the difference between **Reptile** and **Mammal**. This time the p-value is 0.039 which is below the standard threshold of 0.05 which can be take that the difference between reptile and mammal is statistically significant. Mammals are shown to have a statistically significant higher rate of needed protection compared with Reptiles.

```
[240]: contingency2 = [[30, 146],
                      [5, 73]]
chi2_contingency(contingency2)
```

```
[240]: Chi2ContingencyResult(statistic=4.289183096203645, pvalue=0.038355590229699,
dof=1, expected_freq=array([[ 24.2519685, 151.7480315],
[ 10.7480315,  67.2519685]]))
```

## 2.1 National Parks

Lets see all unique national parks and everything related to it

```
[197]: unique_categories = combined.groupby('park_name')['category'].nunique()
print(f"All Parks:{combined.park_name.unique()}")
print(f"Total number of observations:{combined.observations.sum()}")
```

```
All Parks:['Great Smoky Mountains National Park' 'Yosemite National Park'
'Bryce National Park' 'Yellowstone National Park']
Total number of observations:3645247
```

Lets make bar chart from them:

Bar Chart: Unique Parks and Amount of different species per each park

As we can see that each park have same amount of species per park which can happen beacause the data is artificial in other case that would be really strange on real data.

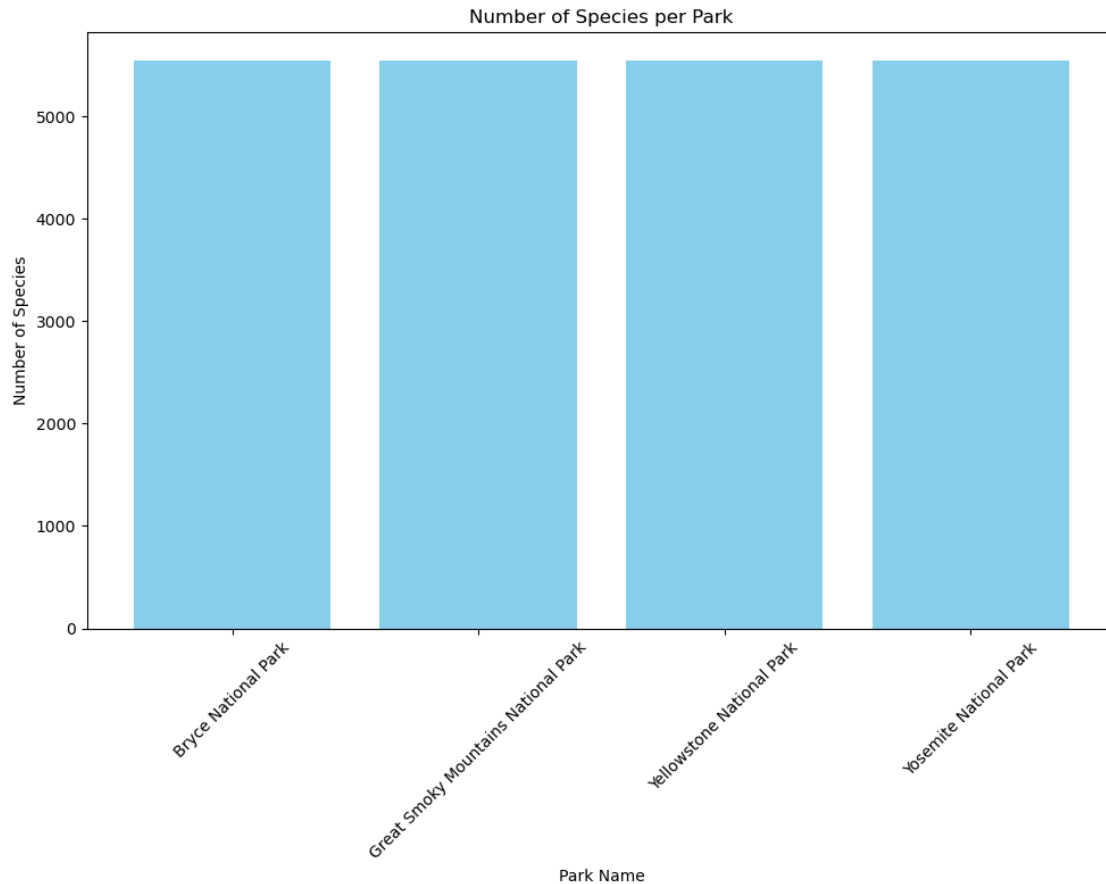
```
[82]: species_count = combined.groupby('park_name')['scientific_name'].nunique()

plt.figure(figsize=(10, 8))
plt.bar(species_count.index, species_count.values, color='skyblue')

plt.xlabel("Park Name")
plt.ylabel("Number of Species")
```



```
plt.title("Number of Species per Park")
plt.xticks(rotation=45)
plt.tight_layout()
plt.show()
```



Bar Chart: Unique Parks vs. Most Popular Category Count Same thing here but for plants to be popular is explainable

```
[86]: category_counts = combined.groupby(['park_name', 'category']).size().
      ↪reset_index(name='count')

most_popular = category_counts.loc[category_counts.
      ↪groupby('park_name')['count'].idxmax()]

plt.figure(figsize=(10, 6))
bars = plt.bar(most_popular['park_name'], most_popular['count'],
      ↪color='lightgreen')

plt.xlabel("Park Name")
```

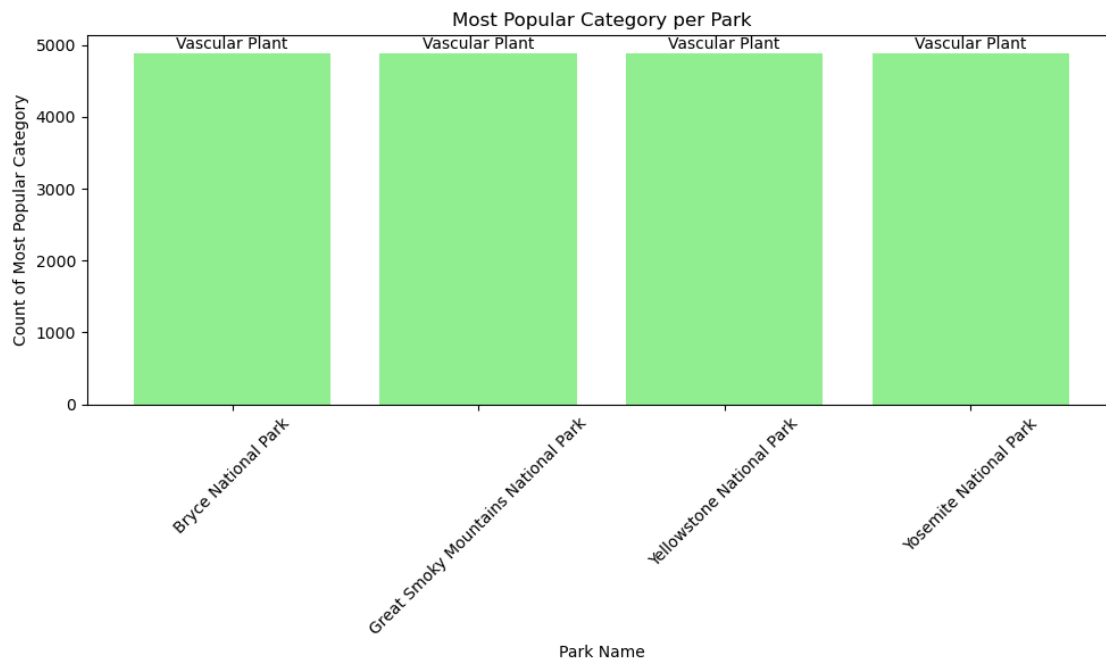
```

plt.ylabel("Count of Most Popular Category")
plt.title("Most Popular Category per Park")
plt.xticks(rotation=45)

for index, row in most_popular.iterrows():
    plt.text(row['park_name'], row['count'] + 5, row['category'], ha='center', va='bottom')

plt.tight_layout()
plt.show()

```



So making our observations we can say that most popular category per park is **Vascular plant** and each park has over 5000 **species**

**Species in Parks** The next set of analysis will come from data from the conservationists as they have been recording sightings of different species at several national parks for the past 7 days. The first step is to look at the the common names from **species** to get an idea of the most prevalent animals in the dataset. The data will be need to be split up into individual names.

```

[268]: from itertools import chain
import string

def remove_punctuations(text):
    for punctuation in string.punctuation:

```

```

        text = text.replace(punctuation, '')
    return text

common_Names = combined[combined.category == "Mammal"]\
    .common_names\
    .apply(remove_punctuations)\
    .str.split().tolist()

common_Names[:6]

```

```

[268]: [['American', 'Mink'],
        ['Northern', 'Short', 'Tailed', 'Shrew', 'Northern', 'ShortTailed', 'Shrew'],
        ['WhiteTailed', 'Deer'],
        ['WhiteTailed', 'Deer', 'WhiteTailed', 'Deer'],
        ['Panther', 'Mountain', 'Lion'],
        ['Cougar', 'Mountain', 'Lion', 'Puma']]

```

Clean duplicates

```

[271]: cleanRows = []

for item in common_Names:
    item = list(dict.fromkeys(item))
    cleanRows.append(item)

cleanRows[:6]

```

```

[271]: [['American', 'Mink'],
        ['Northern', 'Short', 'Tailed', 'Shrew', 'ShortTailed'],
        ['WhiteTailed', 'Deer'],
        ['WhiteTailed', 'Deer'],
        ['Panther', 'Mountain', 'Lion'],
        ['Cougar', 'Mountain', 'Lion', 'Puma']]

```

Putting everything into one list, and the counting most popular

```

[274]: res = list(chain.from_iterable(i if isinstance(i, list) else [i] for i in
    ↪ cleanRows))
res
res[:6]

```

```

[274]: ['American', 'Mink', 'Northern', 'Short', 'Tailed', 'Shrew']

```

```

[276]: words_counted = []

for i in res:
    x = res.count(i)
    words_counted.append((i,x))

```

```
pd.DataFrame(set(words_counted), columns=['Word', 'Count']).
↳sort_values("Count", ascending = False).head(10)
```

```
[276]:
```

	Word	Count
21	Bat	144
32	Myotis	108
98	Shrew	104
54	American	96
184	Mouse	72
80	Mountain	68
74	Common	64
144	Gray	64
119	Chipmunk	60
33	Brown	56

From this analysis, it seems that most popular is Bats, Myotis and Shrews, so Bat occurred 144 times, Myotis 108 times while Shrew came up 104 times.

In the data, there are several different scientific names for different types of bats. The next task is to figure out which rows of species are referring to bats. A new column made up of boolean values will be created to check if is\_bat is True.

```
[293]: combined['is_bat'] = combined.common_names.str.contains(r"\bBat\b", regex =
↳True)

species.head(10)
```

```
[293]:
```

	category	scientific_name	\
0	Mammal	Clethrionomys gapperi	gapperi
1	Mammal	Bos bison	
2	Mammal	Bos taurus	
3	Mammal	Ovis aries	
4	Mammal	Cervus elaphus	
5	Mammal	Odocoileus virginianus	
6	Mammal	Sus scrofa	
7	Mammal	Canis latrans	
8	Mammal	Canis lupus	
9	Mammal	Canis rufus	

	common_names	conservation_status	\
0	Gapper's Red-Backed Vole	NaN	
1	American Bison, Bison	NaN	
2	Aurochs, Aurochs, Domestic Cattle (Feral), Dom...	NaN	
3	Domestic Sheep, Mouflon, Red Sheep, Sheep (Feral)	NaN	
4	Wapiti Or Elk	NaN	
5	White-Tailed Deer	NaN	
6	Feral Hog, Wild Pig	NaN	

7	Coyote	Species of Concern
8	Gray Wolf	Endangered
9	Red Wolf	Endangered

	is_bat
0	False
1	False
2	False
3	False
4	False
5	False
6	False
7	False
8	False
9	False

Here is a subset of the data where is\_bat is true, returning see the rows that matched. There seems to be a lot of species of bats and a mix of protected vs. non-protected species.

```
[295]: combined[combined.is_bat]
```

```
[295]:
```

	scientific_name	park_name \
286	Lasiurus blossevillii	Bryce National Park
331	Corynorhinus rafinesquii	Yosemite National Park
450	Nycticeius humeralis	Yellowstone National Park
670	Lasiurus blossevillii	Great Smoky Mountains National Park
827	Lasiurus borealis	Yosemite National Park
...	...	...
25460	Eptesicus fuscus	Bryce National Park
25461	Eptesicus fuscus	Bryce National Park
25476	Myotis leibii	Yellowstone National Park
25530	Lasionycteris noctivagans	Bryce National Park
25531	Lasionycteris noctivagans	Bryce National Park

	observations	category \
286	113	Mammal
331	188	Mammal
450	219	Mammal
670	70	Mammal
827	134	Mammal
...	...	...
25460	72	Mammal
25461	72	Mammal
25476	233	Mammal
25530	128	Mammal
25531	128	Mammal

common_names	conservation_status \
--------------	-----------------------

286	Western Red Bat	Species of Concern
331	Rafinesque's Big-Eared Bat	No Intervention
450	Evening Bat	No Intervention
670	Western Red Bat	Species of Concern
827	Eastern Red Bat, Red Bat	No Intervention
...	...	...
25460	Big Brown Bat	Species of Concern
25461	Big Brown Bat, Big Brown Bat	Species of Concern
25476	Eastern Small-Footed Bat, Eastern Small-Footed...	Species of Concern
25530	Silver-Haired Bat	Species of Concern
25531	Silver-Haired Bat, Silver-Haired Bat	Species of Concern

	is_protected	is_bat
286	True	True
331	False	True
450	False	True
670	True	True
827	False	True
...	...	...
25460	True	True
25461	True	True
25476	True	True
25530	True	True
25531	True	True

[144 rows x 8 columns]

Let's see how many total bat observations(across all species) were made at each national park.

The total number of bats observed in each park over the past 7 days are in the table below. Yellowstone National Park seems to have the largest with 8,362 observations and the Great Smoky Mountains National Park having the lowest with 2,411.

```
[298]: combined[combined.is_bat].groupby('park_name').observations.sum().reset_index()
```

```
[298]:
```

	park_name	observations
0	Bryce National Park	3433
1	Great Smoky Mountains National Park	2411
2	Yellowstone National Park	8362
3	Yosemite National Park	4786

Now let's see each park broken down by protected bats vs. non-protected bat sightings. It seems that every park except for the Great Smoky Mountains National Park has more sightings of protected bats than not. This could be considered a great sign for bats.

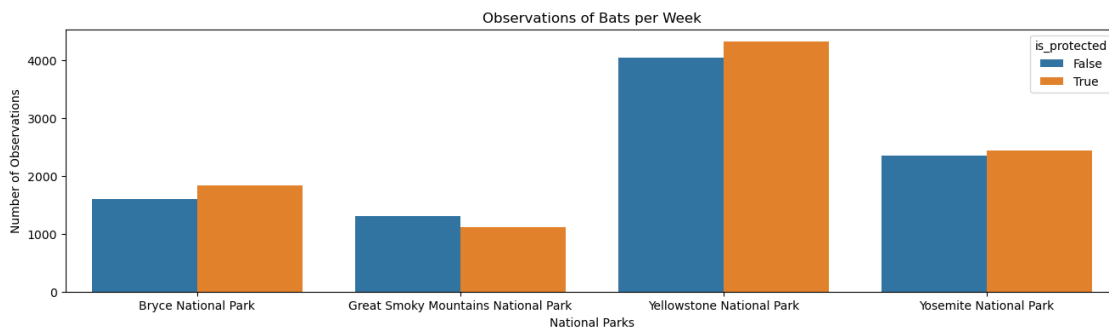
```
[303]: obs_by_park = combined[combined.is_bat].groupby(['park_name', 'is_protected']).
        ↪ observations.sum().reset_index()
obs_by_park
```

```
[303]:
```

	park_name	is_protected	observations
0	Bryce National Park	False	1596
1	Bryce National Park	True	1837
2	Great Smoky Mountains National Park	False	1299
3	Great Smoky Mountains National Park	True	1112
4	Yellowstone National Park	False	4044
5	Yellowstone National Park	True	4318
6	Yosemite National Park	False	2345
7	Yosemite National Park	True	2441

Creating a bar chart based on this data

```
[306]: plt.figure(figsize=(16, 4))
sns.barplot(x=obs_by_park.park_name, y= obs_by_park.observations,
            hue=obs_by_park.is_protected)
plt.xlabel('National Parks')
plt.ylabel('Number of Observations')
plt.title('Observations of Bats per Week')
plt.show()
```



## 2.2 Conclusions

The project was able to make several data visualizations and inferences about the various species in four of the National Parks that comprised this data set.

This project was also able to answer some of the questions first posed in the beginning:

- What is the distribution of conservation status for species?
  - The vast majority of species were not part of conservation.(24752 vs 880)
- Are certain types of species more likely to be endangered?
  - Mammals and Birds had the highest percentage of being in protection.
- Are the differences between species and their conservation status significant?
  - While mammals and Birds did not have significant difference in conservation percentage, mammals and reptiles exhibited a statistically significant difference.
- Which animal is most prevalent and what is their distribution amongst parks?
  - the study found that bats occurred the most number of times and they were most likely to be found in Yellowstone National Park.

- Found that number of species per park and most popular category is the same for all parks which proves that data is most likely to be artificial because its hard to imagine in real life

## 2.3 Further Research

This dataset only included observations from the last 7 days which prohibits analyze changes over time. It would be curious to see how the conservation status for various species changes over time. Another piece that is missing is the Area of each park, it can be assumed that Yellowstone National Park might be much larger than the other parks which would mean that it would exhibit more observations and greater biodiversity. Lastly, if precise locations were recorded, the spatial distribution of the species could also be observed and test if these observations are spatially clustered.

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