

## CodeAcademy BioDiversity

# Are some species more likely to be endangered than others?"

#### **Project Overview**

- The Project had 2 sets of data as 2 csv files.
- 1) In the form of species info which contained information regarding species which contained columns like name, category, common name and conservation status.
- 2) The other file was observations.csv which contained data regarding type of species and their observation location and frequency.

- The first part of the research involved counting the number of unique species and unique categories of species
  along with different status of conservation. Which Were Endangered, in recovery, species of concern, Threatened
  and nan.
- We used the fillna function to replance nan with a new category called no intervention.
- We found the following data
- Conservation Status : Number of species

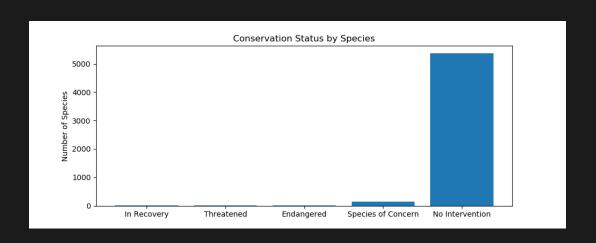
<ul> <li>Endangered</li> </ul>	15
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In Recovery 4

No Intervention 5363

Species of Concern 151

• Threatened 10



• We then wanted to ascertain how much percentage of the species in each category was protected, we did this by grouping category and scientific name to get the total number of species where the conservation status was no intervention and the total number where the conservation status was not no intervention which was placed in the protected column, we then divided protected/protected+nonprotected to get the percentage of protected species in each category. We got the following result

category	not_protected	protected	percent_protected
Amphibian	72	7	0.088608
Bird	413	75	0.153689
Fish	115	11	0.087302
Mammal	146	30	0.170455
Nonvascular Plant	328	5	0.015015
Reptile	73	5	0.064103
Vascular Plant	4216	46	0.010793

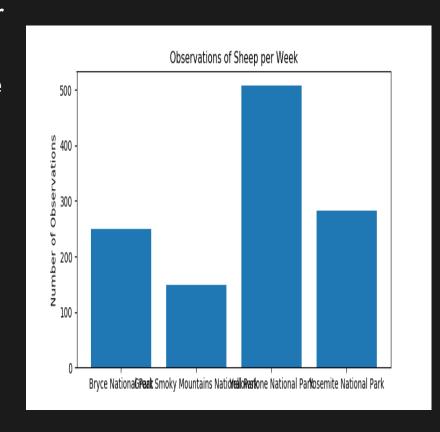
We then used the chi2 test the null hypothesis that the difference was due to chance

We used sample data from to categories Mammals and reptiles,

We found that the pvalue results from the chi2 test were signigicant

0.0383555902297 was the pvalue which was below 0.05 and therefore our null hypothesis was rejected and we could conclude certain species were more likely to be in endangered

- In the second part the national park send us another data set observations. The Data Set used scientific names of the species so we had to we had to create new column which would check if the animal was type of sheep from the original species\_info.csv,
   Once we tested weather a particular species was sheep using the lambda function, We then merged it with the observation dataframe which contained location and number observations
- We merged the two dataframes and grouped it by parkname to determine the sum of observations in each park



#### The foot and mouth Disease

#### Baseline

 The baseline was set to 15 % because as per the exercise last year 15 % of the sheep had foot and mouth disease at the Bryce national park. They want to check for reduction

### Minimum Detectable effect

This year it was 10
% at Yellowstone
that is a drop of 5/15
, 33 % which is
significant, In order
to confirm their
findings they wanted
determine the
sample size to see if
their program was
working

#### Statistical significance

We are chosing 90
 % as the statistical
 significance as this
 is how sure of the
 result that we
 derieve from the
 sample size.

The sample size was 510 as per the calculator and the 3 parameters mentioned.

#### Conclusion

- Based on the result we have concluded the following
- There is a significant difference between the levels of endangerment between different types of species. In our case it was reptiles vs mammals where the Pvalues were below 0.05, Certain species of Mamals and birds seemed to be most in danger.
- I would recommend the national parks to be careful about poaching and provide more protection for birds and mamals.
- It could also be weather conditions that might not suit species and therefore shifting might be in order to protect them.