

Biodiversity in National Parks

Created By: Alec F. for Codecademy Introduction to
Data Analysis Course

Data Analyzed

	category	scientific_name	common_names
0	Mammal	Clethrionomys gapperi gapperi	Gapper's Red-Backed Vole
1	Mammal	Bos bison	American Bison, Bison
2	Mammal	Bos taurus	Aurochs, Aurochs, Domestic Cattle
3	Mammal	Ovis aries	Domestic Sheep, Mouflon, Red Shee
4	Mammal	Cervus elaphus	Wapiti Or Elk

- Contained information for all species in the Parks
- Includes identifying factors of species such as Scientific and common name
- Key data: indicates if species has conservation status

Key Takeaways

- 5541 Unique species
- Mammals and Birds have highest percentage of conservation status with 17% and 15%
- Mammals are significantly more likely to be endangered than reptiles, but not significantly more likely to be endangered than birds

Significance Calculation

- To determine significance of conservation status:
 - Chi-Squared test conducted
 - Contingency test contained number of protected and non-protected species (left and right columns respectively)
 - First tested Mammals and Birds (Blue)
 - Next tested Mammals and Reptiles (Red)
 - P-val of 0.68 and 0.03 respectively
 - Since 0.03 is less than 0.05, it is significant

```
contingency = [[30, 146],  
               [75, 413]]  
  
print contingency  
  
chi2, pval, dof, expected =  
chi2_contingency(contingency)  
  
contingency2 = [[30, 146],  
                [5, 73]]
```

Foot and Mouth Disease

- Using a sample size calculator and data of the number of sheep with foot and mouth disease in parks
- Determined 870 is the sample size per variant
- Using this and the data on observations by park, determined number of weeks to determine significant data:
 - Yellowstone: 1.7 weeks
 - Bryce: 3.5 weeks

Baseline
conversion
rate:

15

%

Statistical
significance:

85%

90%

95%

Minimum
detectable
effect:

33.33

%

Sample
size:

870