

biodiversity

July 23, 2025

```
[4]: from matplotlib import pyplot as plt
import pandas as pd
```

```
[6]: species = pd.read_csv('species_info.csv')
```

```
[7]: species.head()
```

```
[7]:
```

	category	scientific_name	\
0	Mammal	Clethrionomys gapperi	gapperi
1	Mammal	Bos	bison
2	Mammal	Bos	taurus
3	Mammal	Ovis	aries
4	Mammal	Cervus	elaphus

	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

```
[8]: species.scientific_name.nunique()
```

```
[8]: 5541
```

```
[9]: species.category.unique()
```

```
[9]: array(['Mammal', 'Bird', 'Reptile', 'Amphibian', 'Fish', 'Vascular Plant',
        'Nonvascular Plant'], dtype=object)
```

```
[10]: species.conservation_status.unique()
```

```
[10]: array([nan, 'Species of Concern', 'Endangered', 'Threatened',
        'In Recovery'], dtype=object)
```

```
[11]: species.groupby('conservation_status').scientific_name.nunique().reset_index()
```

```
[11]: conservation_status  scientific_name
0      Endangered          15
1    In Recovery           4
2 Species of Concern     151
3    Threatened          10
```

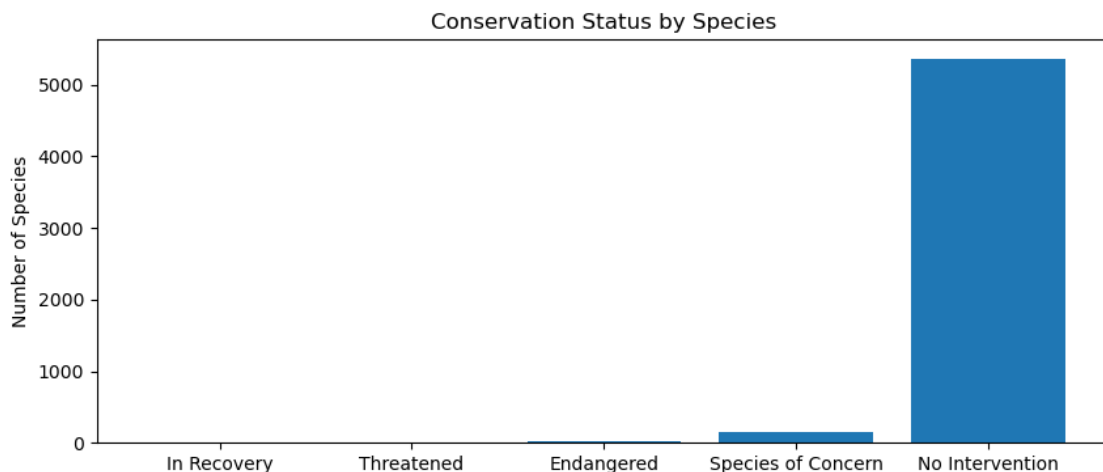
```
[12]: species.fillna('No Intervention', inplace=True)
```

```
[13]: species.groupby('conservation_status').scientific_name.nunique().reset_index()
```

```
[13]: conservation_status  scientific_name
0      Endangered          15
1    In Recovery           4
2    No Intervention     5363
3 Species of Concern     151
4    Threatened          10
```

```
[14]: protection_counts = species.groupby('conservation_status')\
      .scientific_name.nunique().reset_index()\
      .sort_values(by='scientific_name')
```

```
[15]: plt.figure(figsize=(10, 4))
ax = plt.subplot()
plt.bar(range(len(protection_counts)),
        protection_counts.scientific_name.values)
ax.set_xticks(range(len(protection_counts)))
ax.set_xticklabels(protection_counts.conservation_status.values)
plt.ylabel('Number of Species')
plt.title('Conservation Status by Species')
plt.show()
```



```
[16]: species['is_protected'] = species.conservations_status != 'No Intervention'
```

```
[17]: category_counts = species.groupby(['category', 'is_protected'])\
      .scientific_name.nunique().reset_index()
```

```
[18]: category_counts.head()
```

```
[18]:
```

	category	is_protected	scientific_name
0	Amphibian	False	72
1	Amphibian	True	7
2	Bird	False	413
3	Bird	True	75
4	Fish	False	115

```
[19]: category_pivot = category_counts.pivot(columns='is_protected',
      index='category',
      values='scientific_name')\
      .reset_index()
```

```
[20]: category_pivot
```

```
[20]:
```

is_protected	category	False	True
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

```
[21]: category_pivot.columns = ['category', 'not_protected', 'protected']
```

```
[22]: category_pivot['percent_protected'] = category_pivot.protected / \
      (category_pivot.protected + \
      ↪category_pivot.not_protected)
```

```
[23]: category_pivot
```

```
[23]:
```

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

```
[24]: contingency = [[30, 146],  
                    [75, 413]]
```

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

```
[26]: chi2_contingency(contingency)
```

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

```
[27]: contingency = [[30, 146],  
                    [5, 73]]  
chi2_contingency(contingency)
```

```
[27]: Chi2ContingencyResult(statistic=4.289183096203645, pvalue=0.03835559022969898,  
dof=1, expected_freq=array([[ 24.2519685, 151.7480315],  
[ 10.7480315, 67.2519685]]))
```

```
[28]: observations = pd.read_csv('observations.csv')  
observations.head()
```

```
[28]:
```

	scientific_name	park_name	observations
0	Vicia benghalensis	Great Smoky Mountains National Park	68
1	Neovison vison	Great Smoky Mountains National Park	77
2	Prunus subcordata	Yosemite National Park	138
3	Abutilon theophrasti	Bryce National Park	84
4	Githopsis specularioides	Great Smoky Mountains National Park	85

```
[29]: # Does "Sheep" occur in this string?  
str1 = 'This string contains Sheep'  
'Sheep' in str1
```

```
[29]: True
```

```
[30]: # Does "Sheep" occur in this string?  
str2 = 'This string contains Cows'  
'Sheep' in str2
```

```
[30]: False
```

```
[31]: species['is_sheep'] = species.common_names.apply(lambda x: 'Sheep' in x)  
species.head()
```

```
[31]:
```

	category	scientific_name	\
0	Mammal	Clethrionomys gapperi	gapperi
1	Mammal	Bos bison	

```

2  Mammal          Bos taurus
3  Mammal          Ovis aries
4  Mammal          Cervus elaphus

```

```

                                common_names conservation_status \
0          Gapper's Red-Backed Vole      No Intervention
1          American Bison, Bison        No Intervention
2  Aurochs, Aurochs, Domestic Cattle (Feral), Dom... No Intervention
3  Domestic Sheep, Mouflon, Red Sheep, Sheep (Feral) No Intervention
4          Wapiti Or Elk                No Intervention

```

```

    is_protected  is_sheep
0          False    False
1          False    False
2          False    False
3          False     True
4          False    False

```

```
[32]: species[species.is_sheep]
```

```

[32]:          category          scientific_name \
3          Mammal          Ovis aries
1139  Vascular Plant      Rumex acetosella
2233  Vascular Plant      Festuca filiformis
3014          Mammal      Ovis canadensis
3758  Vascular Plant      Rumex acetosella
3761  Vascular Plant      Rumex paucifolius
4091  Vascular Plant      Carex illota
4383  Vascular Plant  Potentilla ovina var. ovina
4446          Mammal      Ovis canadensis sierrae

```

```

                                common_names conservation_status \
3  Domestic Sheep, Mouflon, Red Sheep, Sheep (Feral)      No Intervention
1139          Sheep Sorrel, Sheep Sorrell      No Intervention
2233          Fineleaf Sheep Fescue      No Intervention
3014          Bighorn Sheep, Bighorn Sheep  Species of Concern
3758  Common Sheep Sorrel, Field Sorrel, Red Sorrel,...      No Intervention
3761  Alpine Sheep Sorrel, Fewleaved Dock, Meadow Dock      No Intervention
4091          Sheep Sedge, Smallhead Sedge      No Intervention
4383          Sheep Cinquefoil      No Intervention
4446          Sierra Nevada Bighorn Sheep      Endangered

```

```

    is_protected  is_sheep
3          False     True
1139          False     True
2233          False     True
3014          True      True

```

3758	False	True
3761	False	True
4091	False	True
4383	False	True
4446	True	True

```
[33]: sheep_species = species[(species.is_sheep) & (species.category == 'Mammal')]
      sheep_species
```

```
[33]:      category      scientific_name \
3      Mammal      Ovis aries
3014    Mammal      Ovis canadensis
4446    Mammal  Ovis canadensis sierrae

      common_names conservation_status \
3      Domestic Sheep, Mouflon, Red Sheep, Sheep (Feral)      No Intervention
3014      Bighorn Sheep, Bighorn Sheep      Species of Concern
4446      Sierra Nevada Bighorn Sheep      Endangered

      is_protected  is_sheep
3      False      True
3014      True      True
4446      True      True
```

```
[34]: sheep_observations = observations.merge(sheep_species)
      sheep_observations
```

```
[34]:      scientific_name      park_name \
0      Ovis canadensis      Yellowstone National Park
1      Ovis canadensis      Bryce National Park
2      Ovis canadensis      Yosemite National Park
3      Ovis canadensis  Great Smoky Mountains National Park
4  Ovis canadensis sierrae      Yellowstone National Park
5  Ovis canadensis sierrae      Yosemite National Park
6  Ovis canadensis sierrae      Bryce National Park
7  Ovis canadensis sierrae  Great Smoky Mountains National Park
8      Ovis aries      Yosemite National Park
9      Ovis aries  Great Smoky Mountains National Park
10     Ovis aries      Bryce National Park
11     Ovis aries      Yellowstone National Park

      observations category      common_names \
0      219      Mammal      Bighorn Sheep, Bighorn Sheep
1      109      Mammal      Bighorn Sheep, Bighorn Sheep
2      117      Mammal      Bighorn Sheep, Bighorn Sheep
3      48      Mammal      Bighorn Sheep, Bighorn Sheep
4      67      Mammal      Sierra Nevada Bighorn Sheep
```

5	39	Mammal		Sierra Nevada Bighorn Sheep
6	22	Mammal		Sierra Nevada Bighorn Sheep
7	25	Mammal		Sierra Nevada Bighorn Sheep
8	126	Mammal	Domestic Sheep, Mouflon, Red Sheep, Sheep (Feral)	
9	76	Mammal	Domestic Sheep, Mouflon, Red Sheep, Sheep (Feral)	
10	119	Mammal	Domestic Sheep, Mouflon, Red Sheep, Sheep (Feral)	
11	221	Mammal	Domestic Sheep, Mouflon, Red Sheep, Sheep (Feral)	

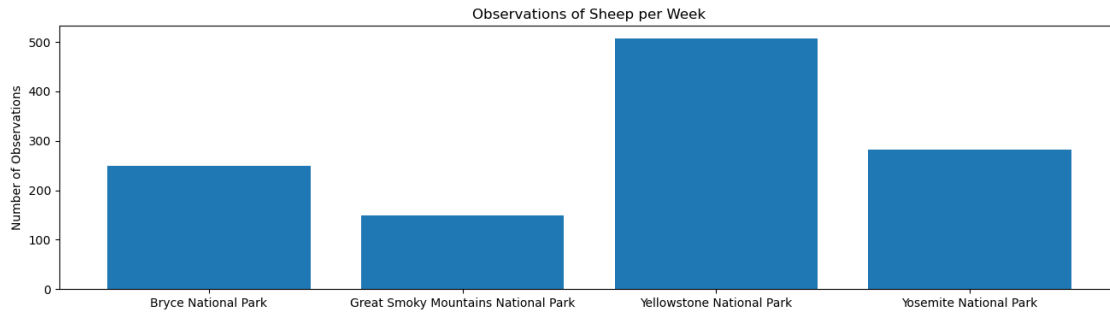
	conservation_status	is_protected	is_sheep
0	Species of Concern	True	True
1	Species of Concern	True	True
2	Species of Concern	True	True
3	Species of Concern	True	True
4	Endangered	True	True
5	Endangered	True	True
6	Endangered	True	True
7	Endangered	True	True
8	No Intervention	False	True
9	No Intervention	False	True
10	No Intervention	False	True
11	No Intervention	False	True

```
[35]: obs_by_park = sheep_observations.groupby('park_name').observations.sum().
      ↪reset_index()
obs_by_park
```

```
[35]:
```

	park_name	observations
0	Bryce National Park	250
1	Great Smoky Mountains National Park	149
2	Yellowstone National Park	507
3	Yosemite National Park	282

```
[36]: plt.figure(figsize=(16, 4))
ax = plt.subplot()
plt.bar(range(len(obs_by_park)),
        obs_by_park.observations.values)
ax.set_xticks(range(len(obs_by_park)))
ax.set_xticklabels(obs_by_park.park_name.values)
plt.ylabel('Number of Observations')
plt.title('Observations of Sheep per Week')
plt.show()
```



```
[37]: minimum_detectable_effect = 100 * 0.05 / 0.15
      minimum_detectable_effect
```

```
[37]: 33.333333333333336
```

```
[38]: baseline = 15
```

```
[39]: sample_size_per_variant = 870
      # Note: This could be 890 if you used 33% for the "Minimum Detectable Effect"
      ↪ instead of 33.33%. That's fine.
```

```
[40]: sample_size_per_variant = 870
      # Note: This could be 890 if you used 33% for the "Minimum Detectable Effect"
      ↪ instead of 33.33%. That's fine.
```

```
[41]: bryce = 870 / 250.
      yellowstone = 810 / 507.

      # Approximately 3.5 weeks at Bryce and 1.5 weeks at Yellowstone.
```

```
[ ]:
```