from matplotlib import pyplot as plt from sklearn.preprocessing import StandardScaler from sklearn import preprocessing import seaborn as sns from sklearn.neighbors import KNeighborsClassifier from sklearn.metrics import confusion\_matrix from sklearn.model\_selection import train\_test\_split from sklearn.metrics import accuracy\_score from sklearn.model\_selection import cross\_val\_score sns.set() df = pd.read\_csv('Zoo.csv') In [11]: In [14]: df1 = df.copy()df1.head() In [15]: Out[15]: animal name hair feathers eggs milk airborne aquatic predator toothed backbone breathes venomous fins legs tail domestic catsize type aardvark 1 1 1 1 1 antelope 0 1 0 2 0 0 1 0 1 1 1 0 1 0 1 0 bass 3 1 bear 1 1 1 1 0 0 1 0 0 1 1 boar df1.describe() In [17]: breathes domestic Out[17]: feathers eggs airborne aquatic predator toothed backbone venomous fins legs tail catsize type 101.000000 101.000000 101.000000 101.000000 101.000000 101.000000 101.000000 101.000000 **count** 101.000000 101.000000 101.000000 101.000000 101.000000 101.000000 101.000000 101.000000 101.000000 0.425743 0.198020 0.584158 0.405941 0.237624 0.356436 0.554455 0.603960 0.821782 0.792079 0.079208 0.168317 2.841584 0.742574 0.128713 0.435644 2.831683 mean 0.496921 0.400495 0.495325 0.493522 0.427750 0.481335 0.499505 0.491512 0.384605 0.407844 0.271410 0.376013 2.033385 0.439397 0.336552 0.498314 2.102709 std 0.000000 0.000000 0.000000 0.000000 0.000000 0.000000 0.000000 0.000000 0.000000 0.000000 0.000000 0.000000 0.000000 0.000000 0.000000 0.000000 1.000000 min 0.000000 0.000000 0.000000 0.000000 0.000000 0.000000 0.000000 0.000000 1.000000 1.000000 0.000000 0.000000 2.000000 0.000000 0.000000 0.000000 1.000000 25% 0.000000 0.000000 1.000000 0.000000 0.000000 0.000000 1.000000 1.000000 1.000000 1.000000 0.000000 0.000000 4.000000 1.000000 0.000000 0.000000 2.000000 0.000000 1.000000 1.000000 1.000000 1.000000 0.000000 1.000000 1.000000 1.000000 1.000000 1.000000 0.000000 0.000000 4.000000 1.000000 0.000000 4.000000 75% 1.000000 1.000000 1.000000 1.000000 1.000000 1.000000 1.000000 1.000000 1.000000 1.000000 1.000000 1.000000 8.000000 1.000000 1.000000 1.000000 7.000000 sns.factorplot('type', data=df1, kind="count", size = 5, aspect = 2) In [18]: C:\Users\HP\anaconda3\lib\site-packages\seaborn\categorical.py:3704: UserWarning: The `factorplot` function has been renamed to `catplot`. The original name will be removed in a fu ture release. Please update your code. Note that the default `kind` in `factorplot` (`'point'`) has changed `'strip'` in `catplot`. warnings.warn(msg) C:\Users\HP\anaconda3\lib\site-packages\seaborn\categorical.py:3710: UserWarning: The `size` parameter has been renamed to `height`; please update your code. warnings.warn(msg, UserWarning) C:\Users\HP\anaconda3\lib\site-packages\seaborn\\_decorators.py:36: FutureWarning: Pass the following variable as a keyword arg: x. From version 0.12, the only valid positional argu ment will be `data`, and passing other arguments without an explicit keyword will result in an error or misinterpretation. warnings.warn( Out[18]: <seaborn.axisgrid.FacetGrid at 0x1741ec5ed00> 40 35 30 25 8 20 15 10 5 0 2 3 4 6 7 type df1.plot(kind='density', subplots=True, layout=(4,5), figsize=(13,20), sharex=False, sharey=False) plt.show() 1.2 hair 2.00 1.2 milk 1.75 1.75 1.0 1.0 1.50 1.0 1.50 1.25 0.8 0.8 1.25 Density 9.0 Density 9.0 <u>₽</u> 1.00 1.00 0.6 ص 0.75 0.75 0.4 0.4 0.4 0.50 0.50 0.2 0.25 0.00 0.0 0 0 1.4 1.2 breathes 2.0 1.75 1.2 1.0 1.0 1.50 1.0 0.8 1.5 1.25 0.8 o.8 Density 9.0 Density 0.1 1.00 0.75 ± 0.6 backbone 를 0.6 0.75 0.4 0.4 0.4 0.50 0.2 0.2 0.2 0.25 0.00 0.0 0.0 0 0 3.5 1.75 fins 0.200 domestic 2.5 2.0 0.175 3.0 1.50 0.150 2.0 2.5 1.25 1.5 Zige 2.0 0.125 ≥ 1.00 1.5 호 0.100 venomous<sup>2</sup> 를 1.5 0.075 1.0 1.0 0.50 0.050 0.5 0.5 0.25 0.025 0.000 0 0 0 0 10 0 catsize 0.25 1.0 0.20 0.8 0.15 Density 9.0 回 0.10 0.2 0.0 0 0 df1.plot(kind='box', subplots=True, layout=(4,5), figsize=(13,20), sharex=False, sharey=False) plt.show() 1.0 1.0 1.0 1.0 1.0 0 0.8 0.8 0.8 0.8 0.8 0.6 0.6 0.6 0.6 0.6 0.4 0.2 0.2 0.2 0.2 0.2 0.0 0.0 0.0 0.0 0.0 feathers hair milk airborne 1.0 1.0 1.0 1.0 0.8 0.8 0.8 0.8 0.8 0.6 0.6 0.6 0.6 0.6 0.4 0.4 0.4 0.4 0.4 0.2 0.2 0.2 0.2 0.2 0.0 0.0 0 0.0 predator toothed backbone breathes aquatic 1.0 1.0 0 1.0 1.0 0 0.8 0.8 0.8 0.8 0.6 0.6 0.6 0.4 0.4 0.4 0.4 0.2 0.2 0.2 0.2 0.0 0.0 0.0 0.0 fins domestic venomous legs 1.0 0.8 0.6 0.2 0.0 catsize Finding correlation between the variables in the data cor = df1.corr(method='pearson') cor.style.background\_gradient(cmap='coolwarm') aquatic feathers airborne predator toothed backbone breathes fins legs domestic catsize Out[22]: hair eggs milk venomous type 0.394009 1.000000 -0.427851 -0.817382 0.878503 -0.198431 -0.473554 -0.154769 0.441149 -0.104245 -0.280313 0.048973 0.455020 -0.562384 hair 0.492531 0.191681 0.207208 0.419248 -0.410761 -0.058552 -0.104430 -0.135934 -0.197520 feathers -0.427851 1.000000 0.656553 -0.613631 0.231403 0.254588 -0.145739 -0.223541 -0.206686 0.292569 0.031586 0.817382 0.419248 1.000000 -0.938848 -0.155610 -0.514650 0.376646 0.376244 0.011605 -0.642150 -0.340420 -0.382777 0.098689 0.164796 -0.224918 -0.221090 0.661825 eggs -0.410761 -0.938848 0.163928 milk 0.878503 1.000000 -0.366765 -0.362613 -0.029721 0.628168 0.384958 0.423527 -0.242449 -0.156328 0.214196 0.210026 0.574906 -0.723683 -0.198431 0.376646 -0.366765 1.000000 -0.172638 -0.295181 -0.594311 -0.104718 0.286039 0.043712 0.009482 0.063274 -0.349768 0.022677 airborne 0.656553 0.008528 -0.251157 -0.172638 -0.034642 -0.224308 -0.111866 -0.473554 -0.058552 0.376244 -0.362613 1.000000 0.375978 0.053150 0.022463 -0.637506 0.087915 0.604492 -0.360638 0.326639 aquatic predator -0.154769 -0.104430 0.011605 -0.029721 -0.295181 0.375978 1.000000 0.129452 0.051022 -0.262931 0.115391 0.190302 -0.099723 0.018947 -0.309794 0.144790 0.061179 0.492531 -0.613631 -0.642150 0.628168 -0.594311 0.053150 0.129452 1.000000 0.575085 -0.065690 -0.062344 0.364292 -0.193476 0.310368 0.069430 0.344010 -0.471527 toothed 0.384958 -0.104718 0.022463 1.000000 -0.246611 0.101733 -0.828845 backbone **0.191681 0.231403** -0.340420 0.051022 0.575085 0.207666 0.209499 -0.432856 0.731762 0.356976 -0.120752 -0.637506 -0.262931 1.000000 -0.617219 0.124068 0.204125 -0.519308 0.441149 0.254588 -0.382777 0.423527 0.286039 -0.065690 0.207666 0.369868 0.088952 breathes -0.120752 0.098689 0.008528 0.115391 -0.062344 -0.246611 1.000000 -0.033956 -0.162724 -0.003252 -0.183748 -0.104245 -0.145739 -0.242449 0.087915 0.022964 0.321476 -0.033956 1.000000 -0.605652 -0.093887 -0.280313 -0.223541 **0.164796** -0.156328 -0.251157 0.604492 0.190302 0.364292 0.209499 -0.617219 0.204349 0.031705 0.099430 fins -0.206686 -0.224918 0.214196 0.043712 -0.360638 -0.099723 -0.193476 -0.432856 0.369868 0.022964 -0.605652 1.000000 -0.348295 0.073931 0.068791 0.131693 legs 0.394009 0.023434 0.243277 -0.631830 -0.221090 0.210026 0.009482 -0.034642 0.018947 0.310368 0.731762 0.088952 -0.162724 0.204349 -0.348295 1.000000 tail 0.063274 -0.224308 -0.309794 -0.003252 -0.093887 0.073931 0.023434 1.000000 0.020073 -0.181043 domestic 0.207208 0.031586 -0.155610 0.163928 0.069430 0.101733 0.124068 -0.349768 -0.111866 0.144790 -0.135934 0.020073 -0.521030 0.455020 -0.514650 0.574906 0.344010 0.356976 0.204125 -0.183748 0.031705 0.068791 0.243277 1.000000 catsize -0.562384 -0.197520 <mark>0.661825</mark> -0.723683 type -0.519308 0.321476 0.099430 0.131693 -0.631830 -0.181043 -0.521030 1.000000 As seen in the above graph, there is a high correlation exists between some of the variables. We can use PCA to reduce the hight correlated variables KNN Finding optimal number of K In [23]: X = df1.iloc[:,1:17]y = df1.iloc[:,17]X\_train, X\_test, y\_train, y\_test = train\_test\_split(X, y, test\_size=0.2, random\_state=1, stratify=y)  $k_{values} = np.arange(1, 25)$ In [26]: train\_accuracy = [] test\_accuracy = [] for i, k in enumerate(k\_values): knn = KNeighborsClassifier(n\_neighbors=k) knn.fit(X\_train,y\_train) train\_accuracy.append(knn.score(X\_train, y\_train)) test\_accuracy.append(knn.score(X\_test, y\_test)) plt.figure(figsize=[13,8]) plt.plot(k\_values, test\_accuracy, label = 'Testing Accuracy') plt.plot(k\_values, train\_accuracy, label = 'Training Accuracy') plt.legend() plt.title('-value VS Accuracy') plt.xlabel('Number of Neighbors') plt.ylabel('Accuracy') plt.xticks(k\_values) plt.show() -value VS Accuracy Testing Accuracy 1.00 Training Accuracy 0.95 0.90 Accuracy 58.0 0.80 0.75 0.70 0.65 15 16 17 18 19 20 21 22 12 14 Number of Neighbors As shown in the graph, with K=5 we can achive accurary of 90%. Applying the algorithm knn = KNeighborsClassifier(n\_neighbors=5) In [29]: knn.fit(X\_train, y\_train) In [30]: y\_pred\_KNeighborsClassifier = knn.predict(X\_test) In [31]: scores = [] cv\_scores = [] score = accuracy\_score(y\_pred\_KNeighborsClassifier,y\_test) In [32]: scores.append(score) In [33]: score\_knn=cross\_val\_score(knn, X,y, cv=10) C:\Users\HP\anaconda3\lib\site-packages\sklearn\model\_selection\\_split.py:670: UserWarning: The least populated class in y has only 4 members, which is less than n\_splits=10. warnings.warn(("The least populated class in y has only %d" score\_knn.mean() 0.890909090909091 score\_knn.std()\*2 In [38]: cv\_score = score\_knn.mean() In [41]: cv\_scores.append(cv\_score) cv\_scores Out[42]: [0.890909090909091] Conclusion Support Vector Machine Accuracy: 0.90 (+/- 0.14)

import numpy as np
import pandas as pd