n []: [import pandas as pd import numpy as np import matplotlib.pyplot as plt import seaborn as sns
]:	SK-Learn Library Import ''' SK-Learn Library Import''' from sklearn.preprocessing import LabelEncoder, OneHotEncoder from sklearn.linear_model import LogisticRegression from sklearn.linear_model import RandomizedLasso, LassoLarsCV from sklearn.exceptions import ConvergenceWarning
[]:	
[]:	<pre>import warnings warnings.filterwarnings('ignore') sns.set() %matplotlib inline</pre>
[[11]: [[11]:	Display The head> To Check if Data is Properly Imported df = pd.read_csv(r'https://raw.githubusercontent.com/dsrscientist/dataset1/master/abalone.csv') df.head() Sex Length Diameter Height Whole weight Shucked weight Viscera weight Shell weight Rings
	Sex Length Diameter Height Whole Weight Shucked Weight Viscera Weight Shell Weight Rings 0 M 0.455 0.365 0.095 0.5140 0.2245 0.1010 0.150 15 1 M 0.350 0.265 0.090 0.2255 0.0995 0.0485 0.070 7 2 F 0.530 0.420 0.135 0.6770 0.2565 0.1415 0.210 9 3 M 0.440 0.365 0.125 0.5160 0.2155 0.1140 0.155 10 4 I 0.330 0.255 0.080 0.2050 0.0895 0.0395 0.0395 7 Feature Information of the DataSet
,	<pre>calass 'pandas.core.frame.DataFrame'> RangeIndex: 4177 entries, 0 to 4176 Data columns (total 9 columns): # Column Non-Null Count Dtype </pre>
[13]: -	Length Diameter Height Whole weight Shucked weight Viscera weight Shell weight Rings count 4177.000000 4177.000000 4177.000000 4177.000000 4177.000000 4177.000000 4177.000000 4177.000000 mean 0.523992 0.407881 0.139516 0.828742 0.359367 0.180594 0.238831 9.933684 std 0.120093 0.099240 0.041827 0.490389 0.221963 0.109614 0.139203 3.224169 min 0.075000 0.055000 0.000000 0.001000 0.000500 0.001500 1.000000 25% 0.450000 0.350000 0.141500 0.186000 0.093500 0.130000 8.000000 50% 0.545000 0.480000 0.165000 0.150000 0.523000 0.234000 9.000000 max 0.815000 0.650000 1.130000 2.825500 1.488000 0.760000 1.005000 29.000000
[]:	According to Described Information: 1)No Feature has Minimum Value = 0, except Height 2)All Features are not Normally Distributed, (Theortically if feature is normally distributed, Mean = Median = Mode). 3)But Features are close to Normality 4)All numerical, Except Sex 5)Each Feature has Different Scale
[15]:	<pre>nf = df.select_dtypes(include=[np.number]).columns cf = df.select_dtypes(include=[np.object]).columns df.hist(figsize=(20,20), grid = True, layout = (2,4), bins = 30)</pre>
	<pre>-Axessubplot:title('center':Malaght')>, -Axessubplot:title('center':Malaght')>, -</pre>
	Shucked weight Viscera weight Shell weight Rings 350 300 300 250 250 250 250 300 300 300 300 300 300 300 300 300 3
	\$ 588 1 487 12 287 6 789 131 203 132 203 133 203 134 203 134 203 135 203 136 203 137 203 138 2
[28]:	<pre>import seaborn as sns sns.countplot(x='Sex', data = df) <axessubplot:xlabel='sex', ylabel="count"> 1600 1200 1200 1200 1200 1200 1200 120</axessubplot:xlabel='sex',></pre>
[29]:	Sex Ratio in Abalone print("\nSex Count in Percentage") print(df. Sex. value_counts(normalize = True)) print("\nSex Count in Numbers") print(df. Sex. value_counts()) Sex Count in Percentage M
[30]: [30]:	Small Feature Engineering, Deriving Age from Rings Column, Age = Rings + 1.5" df['Age'] = df['Rings'] + 1.5 df['Age'].head(5) 0 16.5 1 8.5 2 10.5 3 11.5 4 8.5
	4 8.5 Name: Age, dtype: float64 Sex and Age Visulization''' import seaborn as sns plt.figure(figsize = (20,7)) sns.swarmplot(x = 'Sex', y = 'Age', data = df, hue = 'Sex')
	<pre>sns.violinplot(x = 'Sex', y = 'Age', data = df) D:\ANACONDA\lib\site-packages\seaborn\categorical.py:1296: UserWarning: 56.2% of the points cannot be placed; you may want to decrease the size of the material or use stripplot. warnings.warn(msg, UserWarning) D:\ANACONDA\lib\site-packages\seaborn\categorical.py:1296: UserWarning: 52.2% of the points cannot be placed; you may want to decrease the size of the material or use stripplot. warnings.warn(msg, UserWarning)</pre>
[32]:	D:\ANACONDA\lib\site-packages\seaborn\categorical.py:1296: UserWarning: 58.5% of the points cannot be placed; you may want to decrease the size of the mornings.warn(msg, UserWarning) <axessubplot:xlabel='sex', ylabel="Age"> Sex M F 15 15</axessubplot:xlabel='sex',>
	df.groupby('Sex')[['Length', 'Diameter', 'Height', 'Whole weight', 'Age']].mean().sort_values(by = 'Age', ascending = False) Length Diameter Height Whole weight Shucked weight Viscera weight', 'Shell weight Age
[]: [<pre>Preprocessing Data for the Model df['Sex'] = LabelEncoder().fit_transform(df['Sex'].tolist()) transformed_sex_feature = OneHotEncoder().fit_transform(df['Sex'].values.reshape(-1,1)).toarray() df_sex_encoded = pd.DataFrame(transformed_sex_feature, columns = ["Sex_"+str(int(i)) for i in range(transformed_sex_feature.shape[1])]) df = pd.concat([df, df_sex_encoded], axis=1)</pre>
	df · he ad () Sex Length Diameter Height Whole weight Shucked weight Viscera weight Shell weight Rings Age 0 M 0.455 0.365 0.095 0.5140 0.2245 0.1010 0.1500 15 16.5 1 M 0.350 0.265 0.090 0.2255 0.0995 0.0485 0.070 7 8.5
	2 F 0.530 0.420 0.135 0.6770 0.2565 0.1415 0.210 9 10.5 3 M 0.440 0.365 0.125 0.5160 0.2155 0.1140 0.155 10 11.5 4 I 0.330 0.255 0.080 0.2050 0.0895 0.0395 0.0395 0.055 7 8.5
	THANK YOU