

CSC 711 Neural Networks [10-601] Fall 2019

Homework # 1

Part 2

Team Members:

Loading data and summarize data

```
In [12]: # Listing 6

# Load libraries
from matplotlib import pyplot
from pandas import read_csv
from pandas import set_option
from pandas.plotting import scatter_matrix
import numpy
from numpy import set_printoptions
from sklearn.preprocessing import MinMaxScaler
from sklearn.preprocessing import StandardScaler
from sklearn.preprocessing import Normalizer
from sklearn.preprocessing import Binarizer
```

```
In [14]: # Load dataset
url = 'winequality-white.csv'
names = ['fixed acidity', 'volatile acidity', 'citric acid', 'residual sugar', 'chlorides',
         'free sulfur dioxide', 'total sulfur dioxide', 'density', 'pH', 'sulphates', 'alcohol',
         'quality']
dataset = read_csv(url, sep=';')

# Summarize Data

# Descriptive statistics
# shape
print(dataset.shape)

# List all data types used by the DataFrame to characterize each attribute using dtypes property.
set_option('display.max_rows', 500)
print(dataset.dtypes)
# View first 20 rows
set_option('display.width', 100)
print(dataset.head(20))
# descriptions, change precision to 3 places
set_option('precision', 3)
```

(4898, 12)

fixed acidity	float64
volatile acidity	float64

citric acid	float64
residual sugar	float64
chlorides	float64
free sulfur dioxide	float64
total sulfur dioxide	float64
density	float64
pH	float64
sulphates	float64
alcohol	float64
quality	int64

dtype: object

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide \
0	7.0	0.27	0.36	20.70	0.045	45.0
1	6.3	0.30	0.34	1.60	0.049	14.0
2	8.1	0.28	0.40	6.90	0.050	30.0
3	7.2	0.23	0.32	8.50	0.058	47.0
4	7.2	0.23	0.32	8.50	0.058	47.0
5	8.1	0.28	0.40	6.90	0.050	30.0
6	6.2	0.32	0.16	7.00	0.045	30.0
7	7.0	0.27	0.36	20.70	0.045	45.0
8	6.3	0.30	0.34	1.60	0.049	14.0
9	8.1	0.22	0.43	1.50	0.044	28.0
10	8.1	0.27	0.41	1.45	0.033	11.0
11	8.6	0.23	0.40	4.20	0.035	17.0
12	7.9	0.18	0.37	1.20	0.040	16.0
13	6.6	0.16	0.40	1.50	0.044	48.0
14	8.3	0.42	0.62	19.25	0.040	41.0
15	6.6	0.17	0.38	1.50	0.032	28.0
16	6.3	0.48	0.04	1.10	0.046	30.0
17	6.2	0.66	0.48	1.20	0.029	29.0
18	7.4	0.34	0.42	1.10	0.033	17.0
19	6.5	0.31	0.14	7.50	0.044	34.0

	total sulfur dioxide	density	pH	sulphates	alcohol	quality
0	170.0	1.0010	3.00	0.45	8.8	6
1	132.0	0.9940	3.30	0.49	9.5	6
2	97.0	0.9951	3.26	0.44	10.1	6
3	186.0	0.9956	3.19	0.40	9.9	6
4	186.0	0.9956	3.19	0.40	9.9	6
5	97.0	0.9951	3.26	0.44	10.1	6
6	136.0	0.9949	3.18	0.47	9.6	6
7	170.0	1.0010	3.00	0.45	8.8	6
8	132.0	0.9940	3.30	0.49	9.5	6

9	129.0	0.9938	3.22	0.45	11.0	6
10	63.0	0.9908	2.99	0.56	12.0	5
11	109.0	0.9947	3.14	0.53	9.7	5
12	75.0	0.9920	3.18	0.63	10.8	5
13	143.0	0.9912	3.54	0.52	12.4	7
14	172.0	1.0002	2.98	0.67	9.7	5
15	112.0	0.9914	3.25	0.55	11.4	7
16	99.0	0.9928	3.24	0.36	9.6	6
17	75.0	0.9892	3.33	0.39	12.8	8
18	171.0	0.9917	3.12	0.53	11.3	6
19	133.0	0.9955	3.22	0.50	9.5	5

In [15]: *# The describe() function list 8 statistical properties of each attribute.*

```
print(dataset.describe())
```

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	\
count	4898.000	4898.000	4898.000	4898.000	4898.000	
mean	6.855	0.278	0.334	6.391	0.046	
std	0.844	0.101	0.121	5.072	0.022	
min	3.800	0.080	0.000	0.600	0.009	
25%	6.300	0.210	0.270	1.700	0.036	
50%	6.800	0.260	0.320	5.200	0.043	
75%	7.300	0.320	0.390	9.900	0.050	
max	14.200	1.100	1.660	65.800	0.346	

	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	alcohol	\
count	4898.000	4898.000	4898.000	4898.000	4898.000	4898.000	
mean	35.308	138.361	0.994	3.188	0.490	10.514	
std	17.007	42.498	0.003	0.151	0.114	1.231	
min	2.000	9.000	0.987	2.720	0.220	8.000	
25%	23.000	108.000	0.992	3.090	0.410	9.500	
50%	34.000	134.000	0.994	3.180	0.470	10.400	
75%	46.000	167.000	0.996	3.280	0.550	11.400	
max	289.000	440.000	1.039	3.820	1.080	14.200	

	quality
count	4898.000
mean	5.878
std	0.886
min	3.000
25%	5.000
50%	6.000
75%	6.000
max	9.000

```
In [16]: # Group class distribution
print(dataset.groupby('quality').size())

quality
3      20
4     163
5    1457
6    2198
7     880
8     175
9        5
dtype: int64
```

Listing 6a: Pairwise Pearson correlations

```
In [17]: # Correlations between attributes using Pearson's Correlation Coefficient
# 6a. Pairwise Pearson correlations
print(dataset.corr(method = 'pearson'))
```

Correlations between attributes using Pearson's Correlation Coefficient

Data set after applying Pairwise Pearson Correlations.

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	\
fixed acidity	1.000	-0.023	0.289	0.089	0.023	
volatile acidity	-0.023	1.000	-0.149	0.064	0.071	
citric acid	0.289	-0.149	1.000	0.094	0.114	
residual sugar	0.089	0.064	0.094	1.000	0.089	
chlorides	0.023	0.071	0.114	0.089	1.000	
free sulfur dioxide	-0.049	-0.097	0.094	0.299	0.101	
total sulfur dioxide	0.091	0.089	0.121	0.401	0.199	
density	0.265	0.027	0.150	0.839	0.257	
pH	-0.426	-0.032	-0.164	-0.194	-0.090	
sulphates	-0.017	-0.036	0.062	-0.027	0.017	
alcohol	-0.121	0.068	-0.076	-0.451	-0.360	
quality	-0.114	-0.195	-0.009	-0.098	-0.210	

	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	\
fixed acidity	-4.940e-02	0.091	0.265	-4.259e-01	-0.017	
volatile acidity	-9.701e-02	0.089	0.027	-3.192e-02	-0.036	
citric acid	9.408e-02	0.121	0.150	-1.637e-01	0.062	
residual sugar	2.991e-01	0.401	0.839	-1.941e-01	-0.027	
chlorides	1.014e-01	0.199	0.257	-9.044e-02	0.017	
free sulfur dioxide	1.000e+00	0.616	0.294	-6.178e-04	0.059	
total sulfur dioxide	6.155e-01	1.000	0.530	2.321e-03	0.135	
density	2.942e-01	0.530	1.000	-9.359e-02	0.074	
pH	-6.178e-04	0.002	-0.094	1.000e+00	0.156	
sulphates	5.922e-02	0.135	0.074	1.560e-01	1.000	
alcohol	-2.501e-01	-0.449	-0.780	1.214e-01	-0.017	
quality	8.158e-03	-0.175	-0.307	9.943e-02	0.054	

	alcohol	quality
fixed acidity	-0.121	-0.114
volatile acidity	0.068	-0.195
citric acid	-0.076	-0.009
residual sugar	-0.451	-0.098
chlorides	-0.360	-0.210
free sulfur dioxide	-0.250	0.008
total sulfur dioxide	-0.449	-0.175
density	-0.780	-0.307
pH	0.121	0.099
sulphates	-0.017	0.054
alcohol	1.000	0.436
quality	0.436	1.000

Listing 6b: Skew of Univariate Distributions

```
In [18]: # 6b.Skew of Univariate Distributions
# Knowing an attribute has a skew may allow us to perform data preparation to correct the skew and later
# improve the accuracy of our models.

print(dataset.skew())
```

Knowing an attribute has a skew may allow us to perform data preparation to correct the skew and later improve the accuracy of our models.

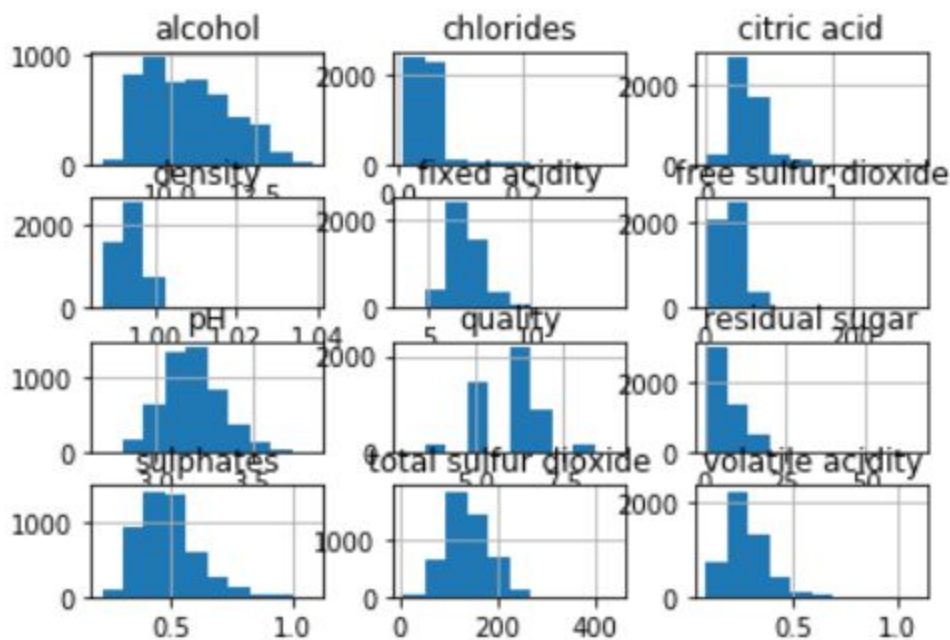
fixed acidity	0.648
volatile acidity	1.577
citric acid	1.282
residual sugar	1.077
chlorides	5.023
free sulfur dioxide	1.407
total sulfur dioxide	0.391
density	0.978
pH	0.458
sulphates	0.977
alcohol	0.487
quality	0.156
dtype:	float64

Listing 6c: Univariate Density Plot

```
In [19]: # 6c. Visualization data with Univariate Plot
# Histograms group data into bin and provide us a count of the number of observations in each bin.
print(dataset.hist())
pyplot(figsize = (8,8))
pyplot.savefig('histograms.png', dpi=300)
pyplot.show()
```

Visualization data with Univariate Plot

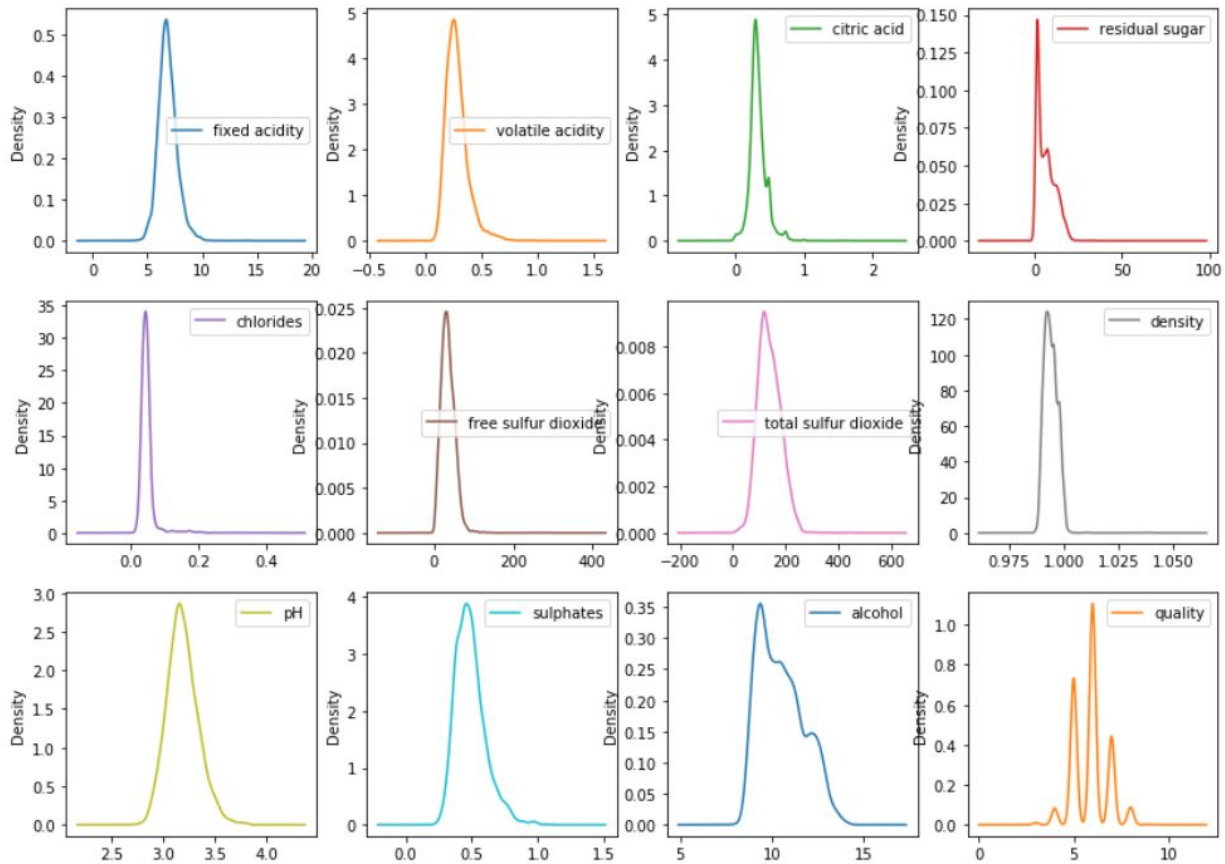
Histograms group data into bin and provide us a count of the number of observations in each bin.



```
In [20]: # Density plots, this help us getting a quick idea of the distribution of each attribute.
# As we can see the distribution for each attribute is clearer than the histograms

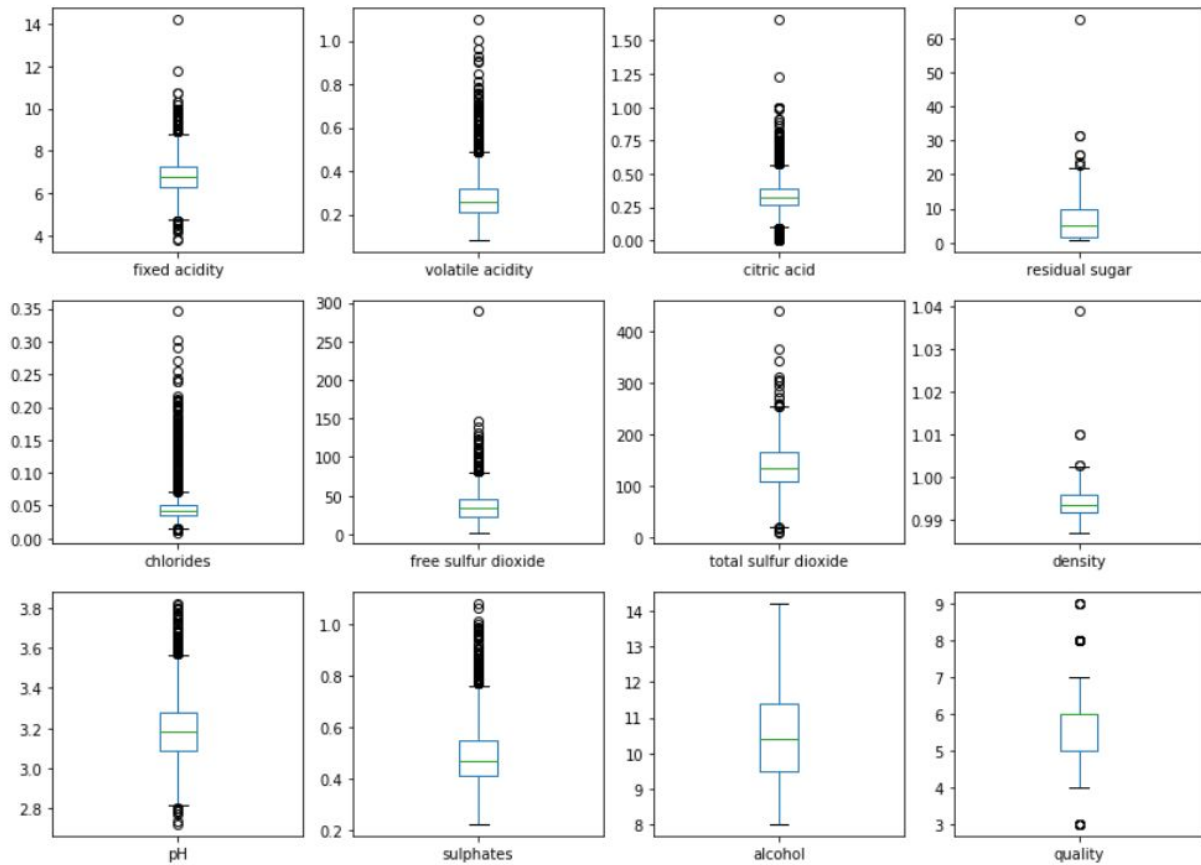
dataset.plot(kind='density', subplots=True, layout=(4,4),sharex=False, figsize = (14,14))
pyplot.show()
```

Density plots, this help us getting a quick idea of the distribution of each attribute.
As we can see the distribution for each attribute is clearer than the histograms




```
In [21]: # Box and Whisker Plots, this give an idea of the spread of data and dot outside of the Whisker show
# candidate outlier values.
dataset.plot(kind='box', subplots=True, layout=(4,4),sharex=False, figsize = (14,14))
pyplot.show()
```

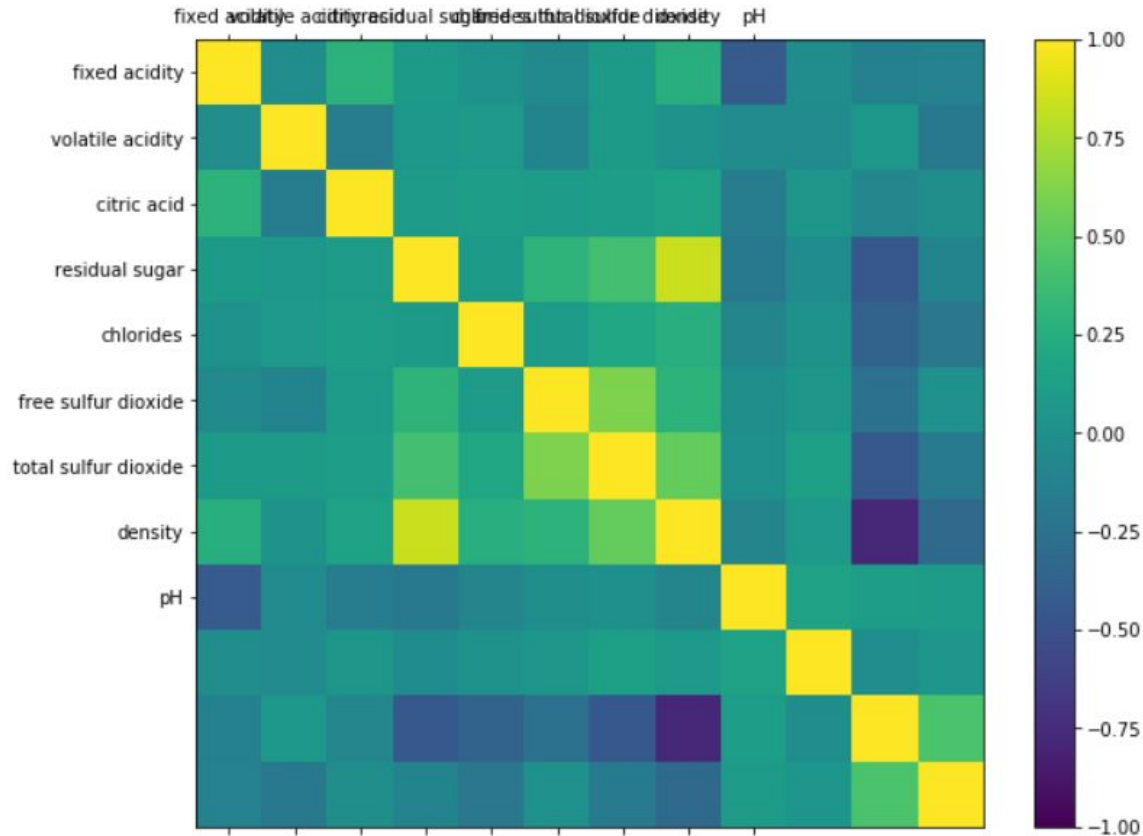
Box and Whisker Plots, this give an idea of the spread of data and dot outside of the Whisker show candidate outlier values.



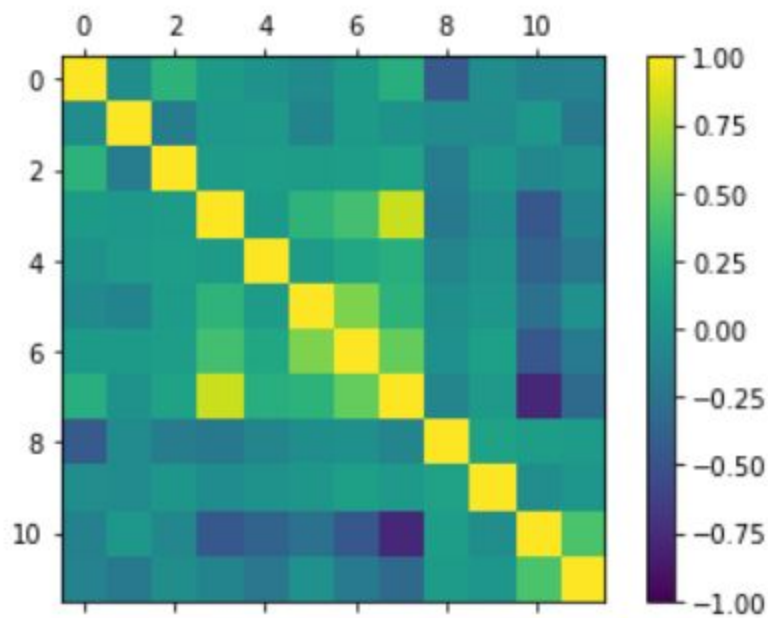
Listing 6d: Correlation Matrix Plot

```
In [30]: # 6d. Correlation Matrix Plot
# This gives an indication of how related the changes are between two variables.
# Plot correlation matrix
fig = pyplot.figure(figsize=(10,8))
ax = fig.add_subplot(111)
cax = ax.matshow(dataset.corr(), vmin=-1, vmax=1)
fig.colorbar(cax)
ticks = numpy.arange(0, 9, 1)
ax.set_xticks(ticks)
ax.set_yticks(ticks)
ax.set_xticklabels(names)
ax.set_yticklabels(names)
pyplot.show()
```

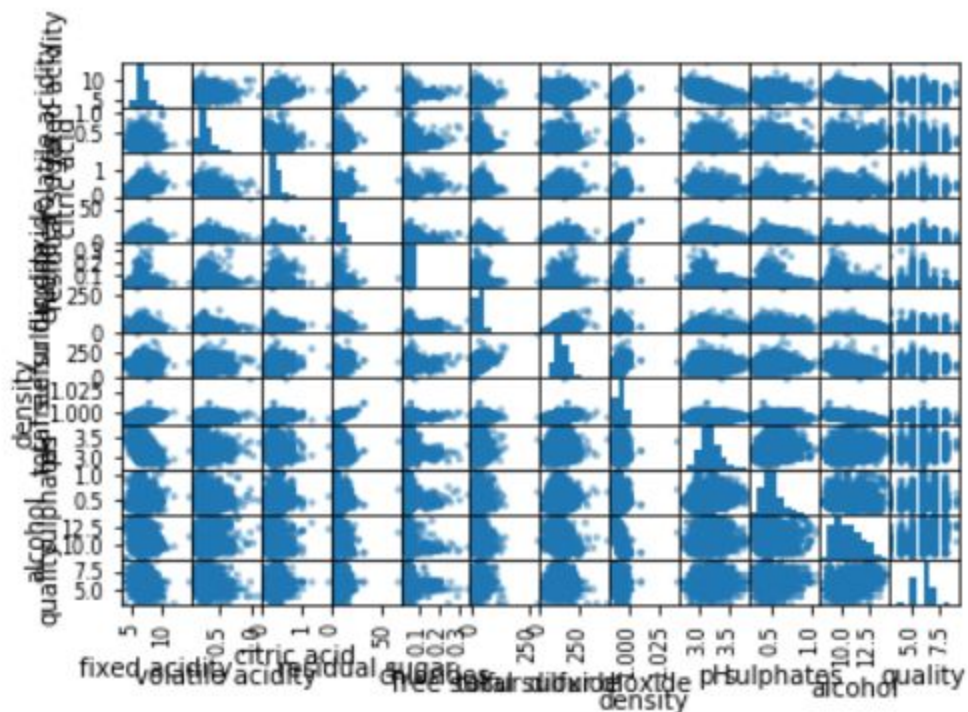
This gives an indication of how related the changes are between two variables.
Plot correlation matrix



```
In [26]: # Correlation Matrix Generic Plot
fig = pyplot.figure()
ax = fig.add_subplot(111)
cax = ax.matshow(dataset.corr(), vmin=-1, vmax=1)
fig.colorbar(cax)
pyplot.show()
```



```
In [35]: # Scatterplot Maxtrix, this shows the relationship between variables
scatter_matrix(dataset)
pyplot.figure(figsize=(20,18))
pyplot.show()
```



Listing 7a: Rescaling Data

```
In [82]: # Listing 7
# 7a. Rescaling data
# After rescaling we can see that all of the values are in the range between 0 and 1.
array = dataset.values
# seperate array into input and output components
X = array[:, 0:11]
Y = array[:, 11]
scaler = MinMaxScaler(feature_range=(0, 1))
rescaledX = scaler.fit_transform(X)
set_printoptions(precision=3)
print(rescaledX[0:5, :])
```

After rescaling we can see that all of the values are in the range between 0 and 1.

```
[[0.308 0.186 0.217 0.308 0.107 0.15 0.374 0.268 0.255 0.267 0.129]
 [0.24 0.216 0.205 0.015 0.119 0.042 0.285 0.133 0.527 0.314 0.242]
 [0.413 0.196 0.241 0.097 0.122 0.098 0.204 0.154 0.491 0.256 0.339]
 [0.327 0.147 0.193 0.121 0.145 0.157 0.411 0.164 0.427 0.209 0.306]
 [0.327 0.147 0.193 0.121 0.145 0.157 0.411 0.164 0.427 0.209 0.306]]
```

Listing 7b: Standardize Data

```
In [84]: # 7b. Standardize Data
X = array[:, 0:11]
Y = array[:, 11]
scaler_standard = StandardScaler().fit(X)
rescaled_standardX = scaler_standard.transform(X)
# summarize transformed data
set_printoptions(precision=3)
print(rescaled_standardX[0:5, :])
```

```
[[ 1.721e-01 -8.177e-02  2.133e-01  2.821e+00 -3.536e-02  5.699e-01
  7.446e-01  2.332e+00 -1.247e+00 -3.492e-01 -1.393e+00]
 [-6.575e-01  2.159e-01  4.800e-02 -9.448e-01  1.477e-01 -1.253e+00
 -1.497e-01 -9.154e-03  7.400e-01  1.342e-03 -8.243e-01]
 [ 1.476e+00  1.745e-02  5.438e-01  1.003e-01  1.935e-01 -3.121e-01
 -9.733e-01  3.587e-01  4.751e-01 -4.368e-01 -3.367e-01]
 [ 4.091e-01 -4.787e-01 -1.173e-01  4.158e-01  5.597e-01  6.875e-01
  1.121e+00  5.259e-01  1.148e-02 -7.873e-01 -4.992e-01]
 [ 4.091e-01 -4.787e-01 -1.173e-01  4.158e-01  5.597e-01  6.875e-01
  1.121e+00  5.259e-01  1.148e-02 -7.873e-01 -4.992e-01]]
```

Listing 7c: Normalize Data

In [86]: # 7c. Normalize Data

```
X = array[:, 0:11]
Y = array[:, 11]
scaler = Normalizer().fit(X)
normalizedX = scaler.transform(X)
# summarize transformed data
set_printoptions(precision=3)
print(normalizedX[0:5, :])
```

```
[[ 3.945e-02  1.522e-03  2.029e-03  1.166e-01  2.536e-04  2.536e-01  9.580e-01
   5.641e-03  1.691e-02  2.536e-03  4.959e-02]
 [ 4.727e-02  2.251e-03  2.551e-03  1.200e-02  3.676e-04  1.050e-01  9.904e-01
   7.458e-03  2.476e-02  3.676e-03  7.128e-02]
 [ 7.891e-02  2.728e-03  3.897e-03  6.722e-02  4.871e-04  2.923e-01  9.450e-01
   9.694e-03  3.176e-02  4.287e-03  9.840e-02]
 [ 3.741e-02  1.195e-03  1.663e-03  4.417e-02  3.014e-04  2.442e-01  9.665e-01
   5.173e-03  1.658e-02  2.078e-03  5.144e-02]
 [ 3.741e-02  1.195e-03  1.663e-03  4.417e-02  3.014e-04  2.442e-01  9.665e-01
   5.173e-03  1.658e-02  2.078e-03  5.144e-02]]
```

Listing 7d: Binarize Data

In [89]: # 7d. Binarize Data

```
binarizer = Binarizer(threshold=0.0).fit(X)
binaryX = binarizer.transform(X)
set_printoptions(precision=3)
print(binaryX[0:5, :])
```

```
[[ 1.  1.  1.  1.  1.  1.  1.  1.  1.  1.  1.]
 [ 1.  1.  1.  1.  1.  1.  1.  1.  1.  1.  1.]
 [ 1.  1.  1.  1.  1.  1.  1.  1.  1.  1.  1.]
 [ 1.  1.  1.  1.  1.  1.  1.  1.  1.  1.  1.]
 [ 1.  1.  1.  1.  1.  1.  1.  1.  1.  1.  1.]]
```

Note that the higher the quality the higher the average alcohol concentration, increased by about 1% at each level. Although lower quality wines have the lowest standard deviation. The chlorides and volatile acidity are less present and presented smaller standard deviation in wines of higher quality. The free sulfur dioxide is higher with higher quality, but their standard deviation decreases with the increase in quality. Higher quality has less fixed acidity, but the standard deviation is slightly higher in mean quality.