Wine_visualisation

January 16, 2025

1 Wine Data Visualisation in Jupyter Notebook - using atoti

Link to the tutorial: https://www.youtube.com/watch?v=Y49662c3EL4&ab_channel=LightsOnData - **Youtube: How to Create a Data Visualization in Jupyter Notebook Using atoti**

[2]:
```
import atoti as tt
import numpy as np
import pandas as pd
import seaborn as sns
```

2 Read the data for the red and white wines

2.1 Red Wine

[6]:
```
wine_red = pd.read_csv(
    "https://data.atoti.io/notebooks/wine-analytics/winequality-red.csv", sep=";
 ↪"
)
wine_red.head()
```

[6]:

	fixed acidity	volatile acidity	citric acid	residual sugar	chlorides	free sulfur dioxide	total sulfur dioxide	density	pH	sulphates	alcohol	quality
0	7.4	0.70	0.00	1.9	0.076	11.0	34.0	0.9978	3.51	0.56	9.4	5
1	7.8	0.88	0.00	2.6	0.098	25.0	67.0	0.9968	3.20	0.68	9.8	5
2	7.8	0.76	0.04	2.3	0.092	15.0	54.0	0.9970	3.26	0.65	9.8	5
3	11.2	0.28	0.56	1.9	0.075	17.0	60.0	0.9980	3.16	0.58		
4	7.4	0.70	0.00	1.9	0.076	11.0	34.0	0.9978	3.51	0.56		

```
3      9.8        6
4      9.4        5
```

2.2 White Wine

[7]:
```
wine_white = pd.read_csv(
    "https://data.atoti.io/notebooks/wine-analytics/winequality-white.csv",␣
 ↪sep=";"
)
wine_white.head()
```

[7]:
```
   fixed acidity  volatile acidity  citric acid  residual sugar  chlorides  \
0            7.0              0.27         0.36            20.7      0.045
1            6.3              0.30         0.34             1.6      0.049
2            8.1              0.28         0.40             6.9      0.050
3            7.2              0.23         0.32             8.5      0.058
4            7.2              0.23         0.32             8.5      0.058

   free sulfur dioxide  total sulfur dioxide  density    pH  sulphates  \
0                 45.0                 170.0   1.0010  3.00       0.45
1                 14.0                 132.0   0.9940  3.30       0.49
2                 30.0                  97.0   0.9951  3.26       0.44
3                 47.0                 186.0   0.9956  3.19       0.40
4                 47.0                 186.0   0.9956  3.19       0.40

   alcohol  quality
0      8.8        6
1      9.5        6
2     10.1        6
3      9.9        6
4      9.9        6
```

2.3 Add a new column “category” to the datasets

[9]:
```
wine_red["category"] = "Red"
wine_red.head()
```

[9]:
```
   fixed acidity  volatile acidity  citric acid  residual sugar  chlorides  \
0            7.4              0.70         0.00             1.9      0.076
1            7.8              0.88         0.00             2.6      0.098
2            7.8              0.76         0.04             2.3      0.092
3           11.2              0.28         0.56             1.9      0.075
4            7.4              0.70         0.00             1.9      0.076

   free sulfur dioxide  total sulfur dioxide  density    pH  sulphates  \
0                 11.0                  34.0   0.9978  3.51       0.56
1                 25.0                  67.0   0.9968  3.20       0.68
```

```
2                 15.0                  54.0   0.9970  3.26       0.65
3                 17.0                  60.0   0.9980  3.16       0.58
4                 11.0                  34.0   0.9978  3.51       0.56

   alcohol  quality category
0      9.4        5      Red
1      9.8        5      Red
2      9.8        5      Red
3      9.8        6      Red
4      9.4        5      Red
```

[11]:
```
wine_white["category"] = "White"
wine_white.head()
```

[11]:
```
   fixed acidity  volatile acidity  citric acid  residual sugar  chlorides  \
0            7.0              0.27         0.36            20.7      0.045
1            6.3              0.30         0.34             1.6      0.049
2            8.1              0.28         0.40             6.9      0.050
3            7.2              0.23         0.32             8.5      0.058
4            7.2              0.23         0.32             8.5      0.058

   free sulfur dioxide  total sulfur dioxide  density    pH  sulphates  \
0                 45.0                 170.0   1.0010  3.00       0.45
1                 14.0                 132.0   0.9940  3.30       0.49
2                 30.0                  97.0   0.9951  3.26       0.44
3                 47.0                 186.0   0.9956  3.19       0.40
4                 47.0                 186.0   0.9956  3.19       0.40

   alcohol  quality category
0      8.8        6    White
1      9.5        6    White
2     10.1        6    White
3      9.9        6    White
4      9.9        6    White
```

[12]:
```
wines = pd.concat([wine_red, wine_white], axis=0, ignore_index=True)
wines.index.set_names("wine index", inplace=True)
wines.head()
```

[12]:
```
            fixed acidity  volatile acidity  citric acid  residual sugar  \
wine index
0                     7.4              0.70         0.00             1.9
1                     7.8              0.88         0.00             2.6
2                     7.8              0.76         0.04             2.3
3                    11.2              0.28         0.56             1.9
4                     7.4              0.70         0.00             1.9
```

```
             chlorides  free sulfur dioxide  total sulfur dioxide  density  \
wine index
0                0.076                 11.0                  34.0   0.9978
1                0.098                 25.0                  67.0   0.9968
2                0.092                 15.0                  54.0   0.9970
3                0.075                 17.0                  60.0   0.9980
4                0.076                 11.0                  34.0   0.9978

               pH  sulphates  alcohol  quality category
wine index
0            3.51       0.56      9.4        5      Red
1            3.20       0.68      9.8        5      Red
2            3.26       0.65      9.8        5      Red
3            3.16       0.58      9.8        6      Red
4            3.51       0.56      9.4        5      Red
```

[13]:
```
wines["alcohol range"] = wines["alcohol"].apply(np.floor)
```

[14]:
```
wines["Rating"] = "Good"
wines.loc[wines["quality"] < 7, "Rating"] = "Average"
wines.loc[wines["quality"] < 5, "Rating"] = "Poor"
wines
```

[14]:
```
             fixed acidity  volatile acidity  citric acid  residual sugar  \
wine index
0                      7.4              0.70         0.00             1.9
1                      7.8              0.88         0.00             2.6
2                      7.8              0.76         0.04             2.3
3                     11.2              0.28         0.56             1.9
4                      7.4              0.70         0.00             1.9
…                      …                 …            …                …
6492                   6.2              0.21         0.29             1.6
6493                   6.6              0.32         0.36             8.0
6494                   6.5              0.24         0.19             1.2
6495                   5.5              0.29         0.30             1.1
6496                   6.0              0.21         0.38             0.8

             chlorides  free sulfur dioxide  total sulfur dioxide  density  \
wine index
0                0.076                 11.0                  34.0  0.99780
1                0.098                 25.0                  67.0  0.99680
2                0.092                 15.0                  54.0  0.99700
3                0.075                 17.0                  60.0  0.99800
4                0.076                 11.0                  34.0  0.99780
…                …                      …                     …      …
6492             0.039                 24.0                  92.0  0.99114
6493             0.047                 57.0                 168.0  0.99490
```

```
6494            0.041                  30.0                  111.0  0.99254
6495            0.022                  20.0                  110.0  0.98869
6496            0.020                  22.0                   98.0  0.98941

              pH  sulphates  alcohol  quality category  alcohol range   Rating
wine index
0           3.51       0.56      9.4        5      Red            9.0  Average
1           3.20       0.68      9.8        5      Red            9.0  Average
2           3.26       0.65      9.8        5      Red            9.0  Average
3           3.16       0.58      9.8        6      Red            9.0  Average
4           3.51       0.56      9.4        5      Red            9.0  Average
…            …          …         …        …       …              …      …
6492        3.27       0.50     11.2        6    White           11.0  Average
6493        3.15       0.46      9.6        5    White            9.0  Average
6494        2.99       0.46      9.4        6    White            9.0  Average
6495        3.34       0.38     12.8        7    White           12.0     Good
6496        3.26       0.32     11.8        6    White           11.0  Average

[6497 rows x 15 columns]
```

[25]:

[25]:
```
      fixed acidity  volatile acidity  citric acid  residual sugar  chlorides  \
0               7.4             0.700         0.00             1.9      0.076
1               7.8             0.880         0.00             2.6      0.098
2               7.8             0.760         0.04             2.3      0.092
3              11.2             0.280         0.56             1.9      0.075
4               7.4             0.700         0.00             1.9      0.076
…               …                 …            …                …        …
1594            6.2             0.600         0.08             2.0      0.090
1595            5.9             0.550         0.10             2.2      0.062
1596            6.3             0.510         0.13             2.3      0.076
1597            5.9             0.645         0.12             2.0      0.075
1598            6.0             0.310         0.47             3.6      0.067

      free sulfur dioxide  total sulfur dioxide  density    pH  sulphates  \
0                    11.0                  34.0  0.99780  3.51       0.56
1                    25.0                  67.0  0.99680  3.20       0.68
2                    15.0                  54.0  0.99700  3.26       0.65
3                    17.0                  60.0  0.99800  3.16       0.58
4                    11.0                  34.0  0.99780  3.51       0.56
…                     …                     …        …     …          …
1594                 32.0                  44.0  0.99490  3.45       0.58
1595                 39.0                  51.0  0.99512  3.52       0.76
1596                 29.0                  40.0  0.99574  3.42       0.75
1597                 32.0                  44.0  0.99547  3.57       0.71
1598                 18.0                  42.0  0.99549  3.39       0.66
```

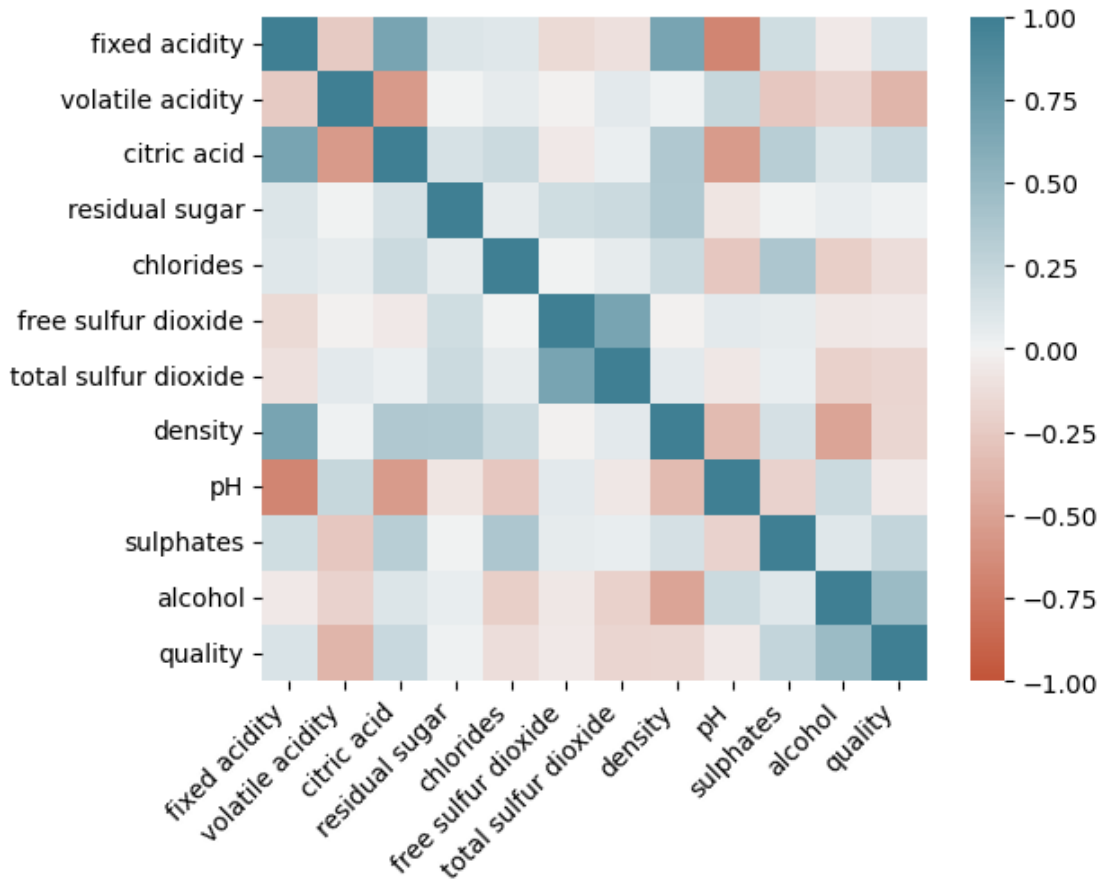
```
      alcohol  quality
0         9.4        5
1         9.8        5
2         9.8        5
3         9.8        6
4         9.4        5
…         …        …
1594     10.5        5
1595     11.2        6
1596     11.0        6
1597     10.2        5
1598     11.0        6

[1599 rows x 12 columns]
```

2.3.1 Prepare the data for corelation

[27]:
```
onlyRedWines = wine_red[wine_red['category'] == 'Red']
redWinesReadyForCorelation = onlyRedWines.drop(columns=['category'])
```

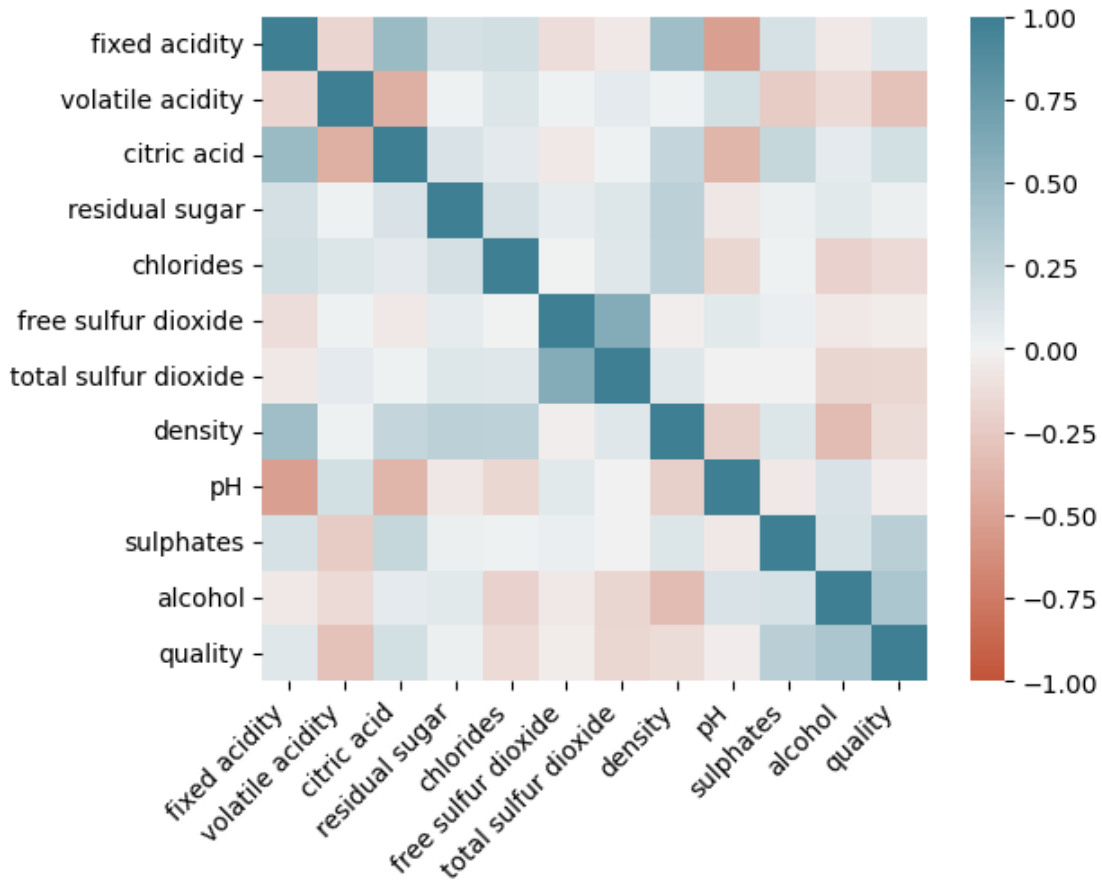
2.4 Default corelation => using Pearson method

[28]:
```
corr_red_pearson = redWinesReadyForCorelation.corr()
ax_red = sns.heatmap(
    corr_red_pearson,
    vmin=-1,
    vmax=1,
    center=0,
    cmap=sns.diverging_palette(20, 220, n=200),
    square=True,
)
ax_red.set_xticklabels(
    ax_red.get_xticklabels(), rotation=45, horizontalalignment="right"
);
```



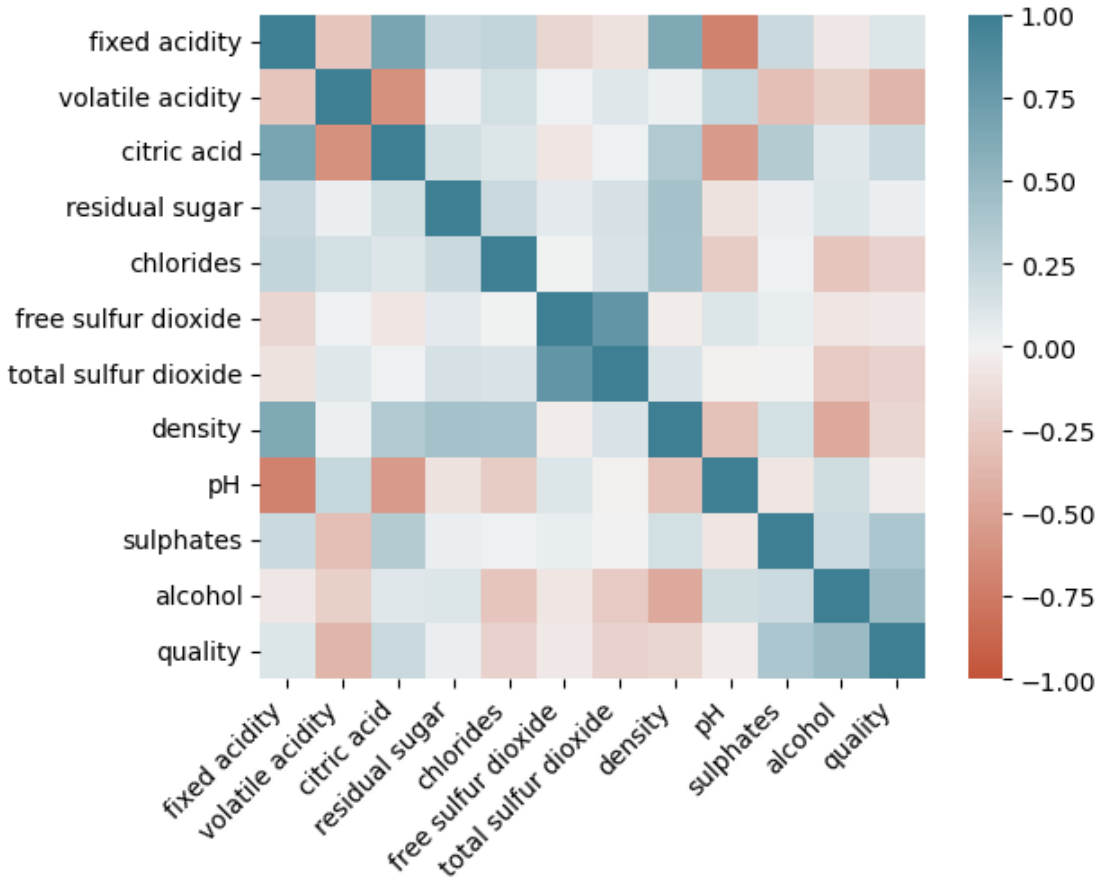
2.5 Corelation using kendall method

[29]:
```
corr_red_kendall = redWinesReadyForCorelation.corr(method='kendall')
ax_red = sns.heatmap(
    corr_red_kendall,
    vmin=-1,
    vmax=1,
    center=0,
    cmap=sns.diverging_palette(20, 220, n=200),
    square=True,
)
ax_red.set_xticklabels(
    ax_red.get_xticklabels(), rotation=45, horizontalalignment="right"
);
```



2.6 Corelation using spearman method

[30]:
```
corr_red_spearman = redWinesReadyForCorelation.corr(method='spearman')
ax_red = sns.heatmap(
    corr_red_spearman,
    vmin=-1,
    vmax=1,
    center=0,
    cmap=sns.diverging_palette(20, 220, n=200),
    square=True,
)
ax_red.set_xticklabels(
    ax_red.get_xticklabels(), rotation=45, horizontalalignment="right"
);
```



[40]:
```
print("Pearson corelation values")
corr_red_pearson[(corr_red_pearson['quality'] > 0.25) |␣
 ↪(corr_red_pearson['quality'] < -0.25)]["quality"]
```

```
Pearson corelation values
```

[40]:
```
volatile acidity   -0.390558
sulphates           0.251397
alcohol             0.476166
quality             1.000000
Name: quality, dtype: float64
```

[42]:
```
print("Kendall corelation values")
corr_red_kendall[(corr_red_kendall['quality'] > 0.25) |␣
 ↪(corr_red_kendall['quality'] < -0.25)]["quality"]
```

```
Kendall corelation values
```

[42]:
```
volatile acidity   -0.300779
sulphates           0.299270
```

```
alcohol              0.380367
quality              1.000000
Name: quality, dtype: float64
```

[43]:
```
print("Spearman corelation values")
corr_red_spearman[(corr_red_spearman['quality'] > 0.25) |␣
 ↪(corr_red_spearman['quality'] < -0.25)]["quality"]
```

```
Spearman corelation values
```

[43]:
```
volatile acidity   -0.380647
sulphates           0.377060
alcohol             0.478532
quality             1.000000
Name: quality, dtype: float64
```

Pearson - Linear relashiopnship - continuous variables - normally distributed data - homoscedasticity - the variability is the same between dependent and non dependent variables

Spearman - Monotonic relashionship - continous or ordinal - non-parametric => does not assume data's distribution - variables are ranked => a ranking has to be done before the analasys is done

BOTH: - look at linear relations (they only decrease or increase…) -

[ ]:
```

```

[ ]:
```
b
```