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RED WINE QUALITY DATA ANALYSIS USING NUMPY PART-2

```
In [ ]:
#Importing Necessary Modules
In [41]:
import numpy as np
In [42]:
wines = np.genfromtxt("winequality-red.csv", delimiter=";", skip_header=1)
In [ ]:
# Sum Of All Residual Sugar Values
In [3]:
y=wines[:,3]
sum(y)
Out[3]:
4059.5500000000003
In [ ]:
# Sum Of Every Feature Value
In [4]:
sum(wines)
Out[4]:
                  , 843.985
array([13303.1
                                   433.29
                                                4059.55
                                                              139.859
                 , 74302.
                                  1593.79794,
                                                5294.47
       25384.
                                                             1052.38
                  , 9012.
                                ])
       16666.35
In [ ]:
# Sum Of Every Row
```

```
In [5]:
z=wines[:,:].sum(axis=1)
Out[5]:
array([ 74.5438 , 123.0548 , 99.699 , ..., 100.48174, 105.21547,
        92.49249])
In [ ]:
# Size
In [6]:
wines.shape[0]
Out[6]:
1599
In [ ]:
# Maximum Residual Sugar value in red wines data
In [7]:
p=wines[:,3]
p=y.astype('int32')
р
Out[7]:
array([1, 2, 2, ..., 2, 2, 3])
In [ ]:
# Maximum Residual Sugar value
In [8]:
max(p)
Out[8]:
15
In [ ]:
# Minimum Residual Sugar value
In [9]:
min(p)
Out[9]:
0
```

```
In [ ]:
# Average Residual Sugar value in red wines data
In [10]:
np.mean(y)
Out[10]:
2.53880550343965
In [ ]:
# 25 percentile residual sugar value
In [11]:
np.percentile(y,25)
Out[11]:
1.9
In [ ]:
# 75 percentile residual sugar value
In [12]:
np.percentile(y,75)
Out[12]:
2.6
In [ ]:
# Average of each feature value
In [13]:
t=wines[:,:]
t.mean(axis=0)
Out[13]:
array([ 8.31963727, 0.52782051, 0.27097561, 2.5388055, 0.08746654,
       15.87492183, 46.46779237, 0.99674668, 3.3111132, 0.65814884,
       10.42298311, 5.63602251])
Numpy Array Comparisons
In [ ]:
# Wines with quality >5
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In [14]:
wines[:,11]>5
Out[14]:
array([False, False, False, ..., True, False, True])
In [ ]:
# Wines with quality >7
In [15]:
k=wines[:,11]>7
k
Out[15]:
array([False, False, False, False, False, False])
In [ ]:
# True for the condition quality >7
In [16]:
True in k
Out[16]:
True
In [ ]:
# First 3 rows whetre wine quality >7,call it high_quality
In [17]:
high_quality=wines[:,11]>7
high_quality
Out[17]:
array([False, False, False, False, False, False])
In [ ]:
# Only top 3 rows and columns of high_quality wines data
```

```
In [18]:
wines[high_quality][0:3]
Out[18]:
array([[7.900e+00, 3.500e-01, 4.600e-01, 3.600e+00, 7.800e-02, 1.500e+01,
        3.700e+01, 9.973e-01, 3.350e+00, 8.600e-01, 1.280e+01, 8.000e+00],
       [1.030e+01, 3.200e-01, 4.500e-01, 6.400e+00, 7.300e-02, 5.000e+00,
        1.300e+01, 9.976e-01, 3.230e+00, 8.200e-01, 1.260e+01, 8.000e+00],
       [5.600e+00, 8.500e-01, 5.000e-02, 1.400e+00, 4.500e-02, 1.200e+01,
        8.800e+01, 9.924e-01, 3.560e+00, 8.200e-01, 1.290e+01, 8.000e+00]])
In [ ]:
# Wines with alcohol >10 and high wine quality >7
In [19]:
alcohol=wines[:,10]>10
alcohol
Out[19]:
array([False, False, False, ..., True, True, True])
In [20]:
h=alcohol & k
h
Out[20]:
array([False, False, False, False, False, False])
In [ ]:
# Alcohol and wine quality columns
In [21]:
wines[h,10:]
Out[21]:
               8.],
array([[12.8,
       [12.6,
               8.],
               8.],
       [12.9,
               8.],
       [13.4,
       [11.7,
               8.],
               8.],
       [11.,
       \lceil 11. , \rceil
               8. ],
       [14.,
               8.],
               8.],
       [12.7,
               8.],
       [12.5,
       [11.8,
               8.],
               8.],
       [13.1,
              8.],
       [11.7,
       [14.,
               8.],
       [11.3,
               8.],
       [11.4,
               8. ]])
```

Combining NumPy Arrays

```
In [ ]:
# Combine red wine and white wine data
In [ ]:
# Open White wine dataset
In [22]:
white_wines=np.genfromtxt("winequality-white.csv",delimiter=";",skip_header=1)
In [ ]:
# Size of White_wines
In [23]:
white_wines.shape
Out[23]:
(4898, 12)
In [ ]:
# Combine wines and white_wines data frames using vstack and call it all_wines
In [24]:
all_wines=np.vstack((wines,white_wines))
In [25]:
all_wines.shape
Out[25]:
(6497, 12)
In [ ]:
# Combine wines and white_wines data frames using concatenate method
In [26]:
all_wines1=np.concatenate((wines,white_wines),axis=0)
In [27]:
all_wines1.shape
Out[27]:
```

(6497, 12)

```
In [ ]:
# Transpose of wines and its size
In [28]:
tran=wines.T
tran.shape
Out[28]:
(12, 1599)
In [ ]:
# Convert wines data into 1D array
In [29]:
wines.ravel()
Out[29]:
array([ 7.4 , 0.7 , 0. , ..., 0.66, 11. , 6. ])
In [ ]:
# Size
In [30]:
wines.ravel().shape
Out[30]:
(19188,)
In [ ]:
# Reshape second row of wines into a 2-dim array with 2rows and 6 columns
In [31]:
wines[1].reshape((2,6))
Out[31]:
                               , 2.6 , 0.098 , 25.
array([[ 7.8
              , 0.88 , 0.
              , 0.9968, 3.2
                                                           ]])
                                , 0.68 , 9.8 , 5.
In [ ]:
# Sort alcohol column ascending order
In [34]:
sorted_alcohol=np.sort(wines[:,-2])
```

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In [35]:
sorted_alcohol
Out[35]:
array([ 8.4, 8.4, 8.5, ..., 14., 14., 14.9])
In [ ]:
# Make sorting to take place in-place
In [36]:
wines[:,-2].sort()
In [37]:
wines[:,-2]
Out[37]:
array([ 8.4, 8.4, 8.5, ..., 14., 14., 14.9])
In [ ]:
# Sort alcohol column Descending order
In [38]:
sorted_alcohol_desc=np.sort(wines[:,10])[::-1]
In [39]:
sorted_alcohol_desc
Out[39]:
array([14.9, 14., 14., ..., 8.5, 8.4, 8.4])
In [ ]:
# will original data be modified?
In [40]:
wines[:,-2]
Out[40]:
array([ 8.4, 8.4, 8.5, ..., 14., 14., 14.9])
In [ ]:
```