

```
In [1]: import pandas as pd
import numpy as np
import seaborn as sns
```

```
In [2]: df=pd.read_excel('health_data_nulls.xlsx')
df.head()
```

Out[2]:

	Person	Age	Income	Alcohol	Exercise	Smoke	Blood Pressure
<b>0</b>	1	61.0	268300.0	41.0	NaN	3.0	62
<b>1</b>	2	55.0	122200.0	51.0	7.0	56.0	53
<b>2</b>	3	53.0	82100.0	37.0	0.0	55.0	42
<b>3</b>	4	30.0	101400.0	41.0	20.0	61.0	48
<b>4</b>	5	64.0	181100.0	NaN	0.0	70.0	81

```
In [3]: df=df.dropna()
df.head()
```

Out[3]:

	Person	Age	Income	Alcohol	Exercise	Smoke	Blood Pressure
<b>1</b>	2	55.0	122200.0	51.0	7.0	56.0	53
<b>2</b>	3	53.0	82100.0	37.0	0.0	55.0	42
<b>3</b>	4	30.0	101400.0	41.0	20.0	61.0	48
<b>8</b>	9	59.0	233500.0	25.0	15.0	33.0	66
<b>9</b>	10	44.0	50400.0	64.0	0.0	85.0	54

```
In [4]: from sklearn import linear_model
from sklearn.model_selection import train_test_split
from matplotlib import pyplot as plt
```

```
In [5]: y = df['Blood Pressure']
df=df[['Age','Income','Alcohol','Exercise','Smoke']]
```

```
In [6]: X_train, X_test, y_train, y_test = train_test_split(df, y, test_size=0.2)
print(X_train.shape, y_train.shape)
print(X_test.shape, y_test.shape)
```

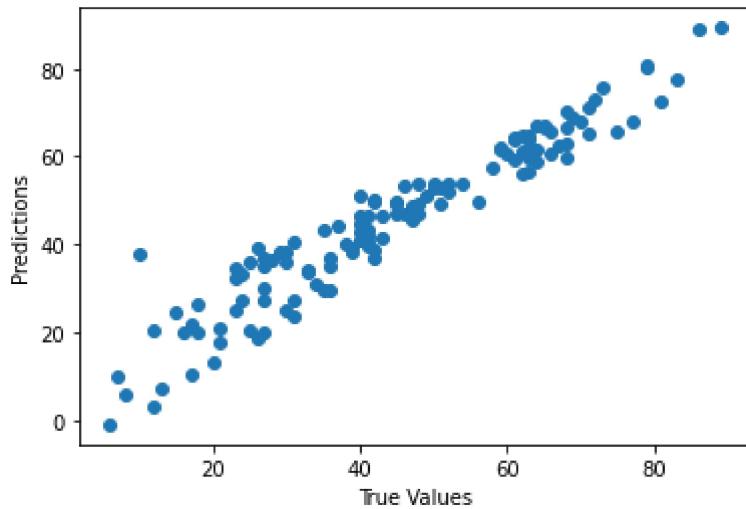
(510, 5) (510,)  
(128, 5) (128,)

```
In [7]: lm=linear_model.LinearRegression()
model=lm.fit(X_train,y_train)
predictions=lm.predict(X_test)
```

```
In [8]: predictions[0:5]
```

```
Out[8]: array([72.635011 , 60.39152806, 33.04744722, -0.98564894, 41.03609251])
```

```
In [9]: plt.scatter(y_test,predictions)
plt.xlabel("True Values")
plt.ylabel("Predictions")
plt.show()
```



```
In [10]: print("Score:", model.score(X_test, y_test))
```

```
Score: 0.9137122610008047
```

```
In [11]: from sklearn.model_selection import KFold
kf = KFold(n_splits=5) #5 splits matches the 80/20 train split
kf.get_n_splits(df) # returns the number of splitting iterations in the cross-validator
print(kf)
KFold(n_splits=5, random_state=None, shuffle=False)
```

```
KFold(n_splits=5, random_state=None, shuffle=False)
```

```
Out[11]: KFold(n_splits=5, random_state=None, shuffle=False)
```

```
In [12]: X=df.to_numpy()
Y=y.to_numpy()
for train_index, test_index in kf.split(X):
    X_train, X_test = X[train_index], X[test_index]
    y_train, t_test = Y[train_index], Y[test_index]
```

```
In [13]: from sklearn.model_selection import cross_val_score, cross_val_predict
from sklearn import metrics
```

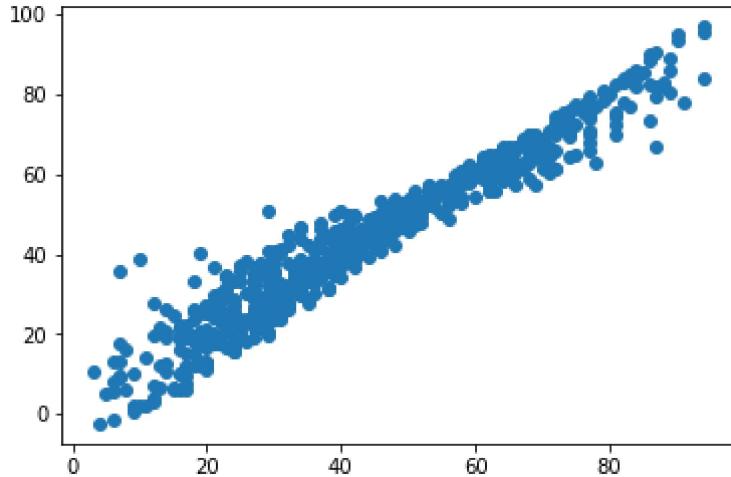
```
In [14]: scores = cross_val_score(model,X, Y, cv=5)
print(scores)
```

```
[0.94136405 0.92254527 0.93177242 0.91735091 0.9503647 ]
```

In [15]:

```
predictions = cross_val_predict(model, X, Y, cv=5)
plt.scatter(Y, predictions)
```

Out[15]: &lt;matplotlib.collections.PathCollection at 0x1e293ce55b0&gt;



In [16]:

```
accuracy = metrics.r2_score(Y,predictions)
accuracy
```

Out[16]: 0.933838939927048

In [17]:

```
#LOOCV Leave One Out Cross Validation, this is equivalent to split equal to observation
#generally only need to use with small data sets
from sklearn.model_selection import LeaveOneOut
loo = LeaveOneOut()
loo.get_n_splits(X)

for train_index, test_index in loo.split(X):
    #print("TRAIN:", train_index, "TEST:", test_index)
    X_train, X_test = X[train_index], X[test_index]
    y_train, y_test = Y[train_index], Y[test_index]
    #print(X_train, X_test, y_train, y_test)
```

In [18]:

#LOOCV is more popular in the classification setting

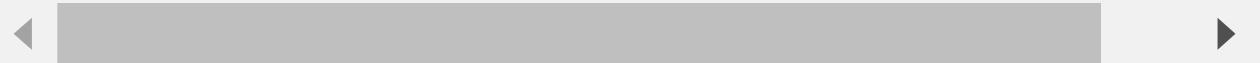
In [20]:

```
mba=pd.ExcelFile('ElecMartSales.xlsx')
df2=mba.parse('Data')
df2.head()
```

Out[20]:

	Date	Day	Time	Region	Card Type	Gender	Buy Category	Items Ordered	Total Cost	High Item	Unnamed: 10
0	2016-03-06	Sun	Morning	West	ElecMart	Female	High	4	136.97	79.97	NaN
1	2016-03-06	Sun	Morning	West	Other	Female	Medium	1	25.55	25.55	NaN

	Date	Day	Time	Region	Card Type	Gender	Buy Category	Items Ordered	Total Cost	High Item	Unnamed: 10
2	2016-03-06	Sun	Afternoon	West	ElecMart	Female	Medium	5	113.95	90.47	NaN
3	2016-03-06	Sun	Afternoon	NorthEast	Other	Female	Low	1	6.82	6.82	NaN
4	2016-03-06	Sun	Afternoon	West	ElecMart	Male	Medium	4	147.32	83.21	NaN



In [21]:

```
two_way=pd.crosstab(df2['Region'], df2['Gender'])
two_way
```

Out[21]:

	Gender	Female	Male
Region			
MidWest		43	28
NorthEast		62	53
South		63	30
West		66	55

In [22]:

```
from scipy.stats import chi2_contingency
stat, p, dof, expected = chi2_contingency(two_way)
```

In [23]:

stat

Out[23]:

5.172525311449733

In [24]:

p

Out[24]:

0.15959121450450045

In [25]:

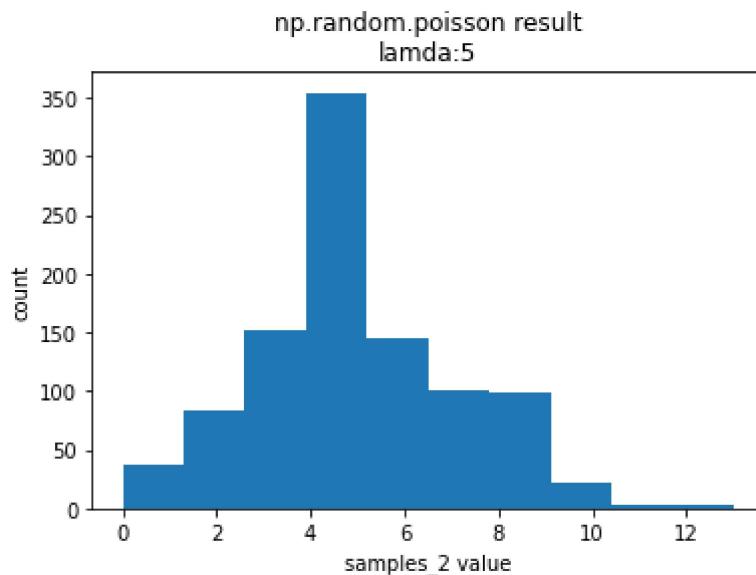
```
from scipy.stats import f_oneway
is_MidWest = df2['Region']=='MidWest'
data1 = df2[is_MidWest]['Total Cost'].to_numpy()
is_NE = df2['Region']=='NorthEast'
data2 = df2[is_NE]['Total Cost'].to_numpy()
is_South = df2['Region']=='South'
data3 = df2[is_South]['Total Cost'].to_numpy()
is_West = df2['Region']=='West'
data4 = df2[is_West]['Total Cost'].to_numpy()
stat, p = f_oneway(data1, data2, data3, data4)
print('stat=% .3f, p=% .3f' % (stat, p))
```

stat=1.964, p=0.119

```
In [27]: np.random.seed(0)
lam, size_1, size_2 = 5, 3, 1000
samples_1 = np.random.poisson(lam, size_1)
samples_2 = np.random.poisson(lam, size_2)
answer_1 = abs(np.mean(samples_1) - lam)
answer_2 = abs(np.mean(samples_2) - lam)
print("|\Lambda - sample mean| with {} samples is {} and with {} samples is {}".format(size_1, answer_1, size_2, answer_2))
```

$|\Lambda - \text{sample mean}|$  with 3 samples is 1.6666666666666667 and with 1000 samples is 0.0579999999999983.

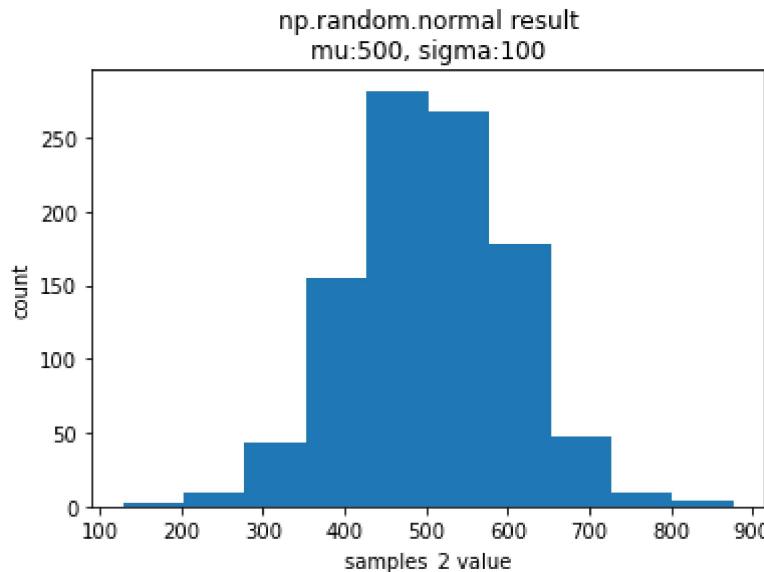
```
In [28]: plt.hist(samples_2)
plt.xlabel('samples_2 value')
plt.ylabel('count')
plt.title('np.random.poisson result\nlamda:5')
plt.show()
```



```
In [29]: np.random.seed(99)
mu, sigma, size_1, size_2 = 500, 100, 3, 1000
samples_1 = np.random.normal(mu, sigma, size_1)
samples_2 = np.random.normal(mu, sigma, size_2)
answer_1 = abs(np.mean(samples_1) - mu)
answer_2 = abs(np.mean(samples_2) - mu)
print("|\mu - sample mean| with {} samples is {} and with {} samples is {}".format(size_1, answer_1, size_2, answer_2))
```

$|\mu - \text{sample mean}|$  with 3 samples is 73.2708278661861 and with 1000 samples is 5.668305231146178.

```
In [30]: plt.hist(samples_2)
plt.xlabel('samples_2 value')
plt.ylabel('count')
plt.title('np.random.normal result\nmu:500, sigma:100')
plt.show()
```



In [31]:

```
#install wordCloud package
from os import path
from PIL import Image
from wordcloud import WordCloud, STOPWORDS, ImageColorGenerator
```

In [33]:

```
df3 = pd.read_csv("winemag-data-130k-v2.csv", index_col=0)
#download from kaggle: https://www.kaggle.com/zynicide/wine-reviews/data
```

In [34]:

```
text=df3.description[0]
```

In [35]:

```
text
```

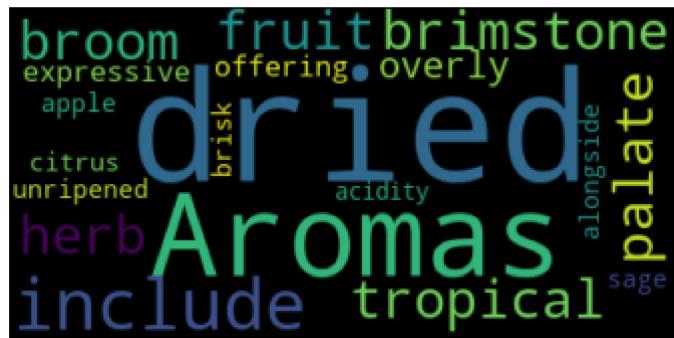
Out[35]: "Aromas include tropical fruit, broom, brimstone and dried herb. The palate isn't overly expressive, offering unripened apple, citrus and dried sage alongside brisk acidity."

In [36]:

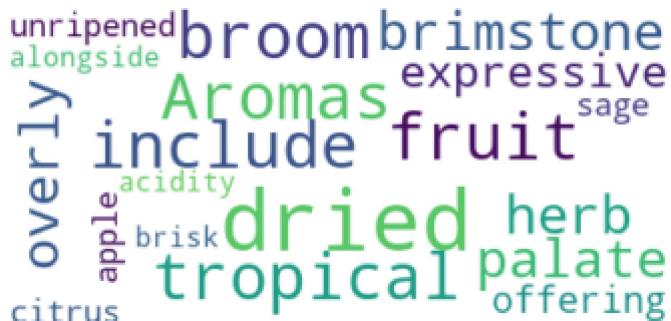
```
wordcloud = WordCloud().generate(text)
```

In [37]:

```
plt.imshow(wordcloud, interpolation='bilinear')
plt.axis('off')
plt.show()
```



```
In [38]: wordcloud = WordCloud(max_font_size=50, max_words=100, background_color="white").generate(text)
plt.figure()
plt.imshow(wordcloud, interpolation="bilinear")
plt.axis("off")
plt.show()
```



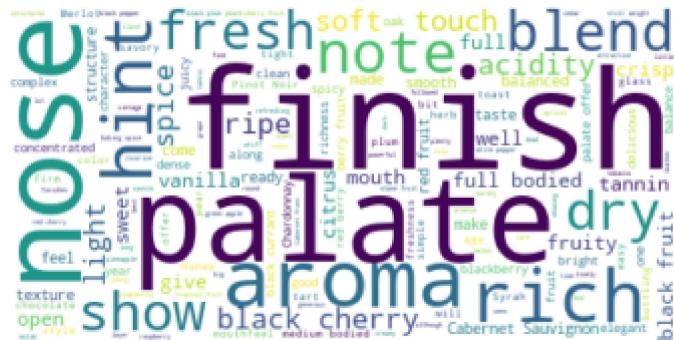
```
In [39]: text = " ".join(review for review in df3.description)
print ("There are {} words in the combination of all review.".format(len(text)))
```

There are 31661073 words in the combination of all review.

```
In [40]: stopwords = set(STOPWORDS)
stopwords.update(["drink", "now", "wine", "flavor", "flavors"])
```

```
In [41]: wordcloud = WordCloud(stopwords=stopwords, background_color="white").generate(text)
```

```
In [42]: plt.imshow(wordcloud, interpolation='bilinear')
plt.axis("off")
plt.show()
```



If you want to learn more about making masks for wordclouds (to take on different shapes than the default), check out this site: <https://www.datacamp.com/community/tutorials/wordcloud-python>