```
import numpy as np
from sklearn.preprocessing import StandardScaler as StandardScaler
from sklearn.decomposition import PCA as PCA
import pandas as pd

X_array = [0, 1, 2, 2, 3, 3, 4]
Y_array = [1, 1, 1, 3, 2, 3, 5]
raw_data = np.column_stack((X_array, Y_array))
```

(a) PCA after standardizing data

```
scaler = StandardScaler()
_ = scaler.fit(raw_data)
standardized_data = scaler.transform(raw_data)
standardized_pca = PCA(n_components=2)
_ = standardized_pca.fit(standardized_data)
print("First Two Components of Standardized PCA")
print(standardized_pca.components_)
First Two Components of Standardized PCA
[[ 0.70710678  0.70710678]
[0.70710678 - 0.70710678]]
standardized_pca_transformed_data = standardized_pca.fit_transform(standardized_data)
standardized_pca_dataframe = pd.DataFrame(data = standardized_pca_transformed_data
            , columns = ['PC 1', 'PC 2'])
standardized_pca_dataframe["Standardized X"] = standardized_data[:, 0]
standardized_pca_dataframe["Standardized Y"] = standardized_data[:, 1]
print(standardized_pca_dataframe)
      PC 1 PC 2 Standardized X Standardized Y
                                    -0.928279
0 -1.873053 -0.560268 -1.720618
                        -0.917663
1 -1.305278 0.007507
                                        -0.928279
                        -0.114708
2 -0.737503 0.575282
                                       -0.928279
3 0.283552 -0.445773
                        -0.114708
                                         0.515711
4 0.340799 0.632529
                         0.688247
                                        -0.206284
5 0.851327 0.122002
                         0.688247
                                         0.515711
6 2.440157 -0.331278
                         1.491202
                                         1.959700
```

```
print(np.matmul(standardized_data, standardized_pca.components_).round(2))

[[-1.87 -0.56]
  [-1.31 0.01]
  [-0.74 0.58]
```

```
[ 0.28 -0.45]
[ 0.34 0.63]
[ 0.85 0.12]
[ 2.44 -0.33]]
```

(b) PCA without standardizing data

```
raw_data_pca = PCA(n_components=2)
_ = raw_data_pca.fit(raw_data)
print("First Two Components of Raw PCA")
print(raw_data_pca.components_)
First Two Components of Raw PCA
[[ 0.65908697  0.75206673]
 [ 0.75206673 -0.65908697]]
raw_pca_transformed_data = raw_data_pca.fit_transform(raw_data)
raw_pca_dataframe = pd.DataFrame(data = raw_pca_transformed_data
            , columns = ['PC 1', 'PC 2'])
raw_pca_dataframe["Raw X"] = raw_data[:, 0]
raw_pca_dataframe["Raw Y"] = raw_data[:, 1]
print(raw_pca_dataframe)
      PC 1 PC 2 Raw X Raw Y
1 -1.720185 -0.012107 1
2 -1.061098 0.739959 2
                                1
                               1
3 0.443035 -0.578215
                              3
                        3
4 0.350055 0.832939
                              2
5 1.102122 0.173852 3
6 3.265343 -0.392255 4
                              3
                                5
```

```
print(np.matmul(raw_data, raw_data_pca.components_).round(2))

[[ 0.75 -0.66]
  [ 1.41   0.09]
  [ 2.07   0.85]
  [ 3.57 -0.47]
  [ 3.48   0.94]
  [ 4.23   0.28]
  [ 6.4   -0.29]]
```

Since manually multiplying the Principal Components with the Raw Dataset does not yield the same result as the sklearn PCA model, we know that PCA is not scale invariant.

Therefore, it is best to standardize the data before the procedure.

```
import numpy as np
from sklearn.preprocessing import PolynomialFeatures
from sklearn.linear_model import LinearRegression
from sklearn.model_selection import KFold
from sklearn.model_selection import cross_val_score
from matplotlib import pyplot as plt
raw_poly_data = np.loadtxt("poly_data.csv", delimiter=" ")
print(str("Shape of Data: {}").format(raw_poly_data.shape))
Shape of Data: (200, 2)
number_of_splits = 5
polynomial_degree_array = np.arange(1, 41, 1)
mse_error_array = []
for poly_degree in polynomial_degree_array:
    polynomial_features = PolynomialFeatures(degree=poly_degree)
    transformed_poly_data = polynomial_features.fit_transform(raw_poly_data[:, 0].reshape(-1, 1))
    linear_regression_obj = LinearRegression()
    _ = linear_regression_obj.fit(transformed_poly_data, raw_poly_data[:, 1])
    if (poly_degree % 5 == 0):
        print(str("Polynomial Degree: {:2d}; Training Accuracy: {:3.1f}%").format(poly_degree,
                                                                                  linear_regression_c
                                                                                      transformed_pol
                                                                                      raw_poly_data[:
    cross_validation_obj = KFold(n_splits=number_of_splits)
    computed_cross_val_scores = -1 * cross_val_score(linear_regression_obj, transformed_poly_data, ra
                                                     scoring='neg_mean_squared_error', cv=cross_valid
    mse_error_array.append(computed_cross_val_scores)
Polynomial Degree: 5; Training Accuracy: 94.7%
Polynomial Degree: 10; Training Accuracy: 94.8%
Polynomial Degree: 15; Training Accuracy: 94.9%
Polynomial Degree: 20; Training Accuracy: 95.0%
Polynomial Degree: 25; Training Accuracy: 95.1%
Polynomial Degree: 30; Training Accuracy: 76.3%
Polynomial Degree: 35; Training Accuracy: 60.2%
Polynomial Degree: 40; Training Accuracy: 54.9%
```

```
plt.rcParams['figure.figsize'] = (12, 5)
fig, axs = plt.subplots(1, 2, sharex="all")
mse_error_array = np.array(mse_error_array)
for poly_degree in range(polynomial_degree_array[-1]):
    for split in range(number_of_splits):
        axs[0].scatter(poly_degree, mse_error_array[poly_degree][split], c="Purple", s=2)
        axs[0].set_yscale('linear')
for poly_degree in range(polynomial_degree_array[-1]):
    for split in range(number_of_splits):
        axs[1].scatter(poly_degree, mse_error_array[poly_degree][split], c="Blue", s=2)
        axs[1].set_yscale('log')
axs[0].grid()
axs[1].grid()
fig.text(0.5, 0.04, 'Polynomial Degree', ha='center')
fig.text(0.04, 0.5, 'MSE Loss', va='center', rotation='vertical')
plt.show()
       400
       300
MSE Loss
                                                        10^{2}
       200
       100
                     10
                               20
                 5
                                    25
                                        30
                                             35
                                                  40
                                                                                               35
                                                Polynomial Degree
```

Which Polynomial Degree fits the data the best?

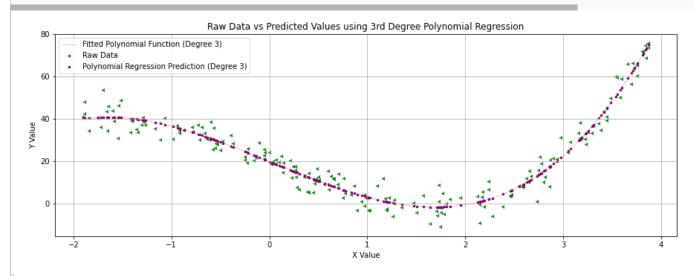
Based on median MSE error computed above for each polynomial degree.

```
median_mse_error_array = np.median(mse_error_array, axis=1)
min_mse = min(median_mse_error_array)
lowest_error_polynomial_degree = median_mse_error_array.tolist().index(min_mse) + 1
print(str("Best Fitting Polynomial Degree is: {}").format(lowest_error_polynomial_degree))

Best Fitting Polynomial Degree is: 3
```

Polynomial Degree: 3; Training Accuracy: 94.7%

Fitted Polynomial Function: $19.8 + (X * -19.0) + (X^2 * -0.1) + (X^3 * 2.2)$



```
import numpy as np
from matplotlib import pyplot as plt
import scipy.stats as stats
```

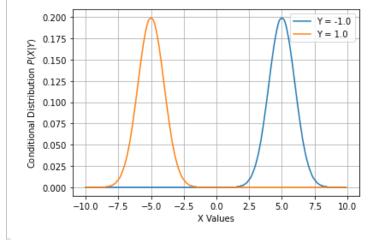
```
def evaluate_conditional_distribution(x, y):
    return (1.0 / (2 * np.sqrt(2 * np.pi)) * np.exp(-(x + 5 * y) ** 2 / 2.0))

vectorized_conditional = np.vectorize(evaluate_conditional_distribution)
```

```
X_array = np.arange(-10.0, 10.0, 0.1)
Y_array = [-1.0, 1.0]
```

```
for y in Y_array:
    evaluated_distribution = vectorized_conditional(X_array, y).round(3)
    plt.plot(X_array, evaluated_distribution, label="Y = " + str(y))

plt.grid()
plt.legend()
plt.xlabel("X Values")
plt.ylabel(r"Conditional Distribution $P(X | Y)$")
plt.show()
```



```
print("Classification Error of Bayes Optimal Classifier described in (c): {:.3e}".format(
    1 - stats.norm.cdf(5, loc=0, scale=1)))
```

Classification Error of Bayes Optimal Classifier described in (c): 2.867e-07

Given:
$$P(X=n, Y=y) = P(Y=y)P(X=x|Y=y)$$

Distailation support $x \in \mathbb{R}, y \in \{-1, +1\}$

$$P(Y=+1) = P(Y=-1) = \frac{1}{2}$$

$$P(X=x|Y=+1) = \frac{1}{\sqrt{2\pi}} e^{-(x-5)^2}$$

$$P(x=x|Y=-1) = \frac{1}{\sqrt{2\pi}} e^{-\frac{(x+5)^2}{2}}$$

Perove
$$P(X = n / Y = y) = 1 e^{-(n+s)^2}$$
 \Rightarrow This is easy just multiply $P(Y = y)$ for both $+1 - 8 - 1$ cares

$$P(X=n \mid Y=+1)$$

$$P(X=n \mid Y=-1) \quad \text{in a single figure}$$

This phreshold can be 0 because each
$$1=18-1$$
 is unimodal & symmetric peaks with $1=18-1$ is $1=1$ whimodal & symmetric peaks $1=18-1$ is $1=18-1$ is $1=18-1$ is

phinal classifies
$$n = \begin{cases} < 0 & y = 1 \\ > 0 & y = -1 \end{cases}$$

3d) compute error of Bayes optimal classifier

$$Pr(h^*(x) \neq y) = E[1]_{h^*(x)} \neq y]$$

$$(x,y) \sim P$$

$$(x,y) \sim P$$

result should be of the parm $1-\overline{\Phi}(c)$ where $\overline{\Phi}$ is Gaussian WF

$$3e$$
 $p(Y=+1|X=n) = \frac{1}{1+e^{-\beta_0-\beta_1x}}$

show that the given distribution eatisping their assumption

$$P(Y=+1|X=X) = P(Y=+1) P(Y=+1) P(Y=+1) P(X=n)$$

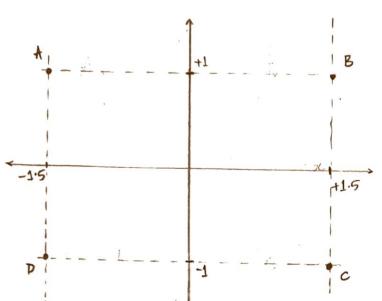
$$P(X=n) = P(X=n) P(X=n)$$

$$P(x=n) = P(x=n|Y=H) \cdot P(Y=H) + P(x=n|Y=-1) \cdot P(Y=H)$$

$$= \frac{1}{2} \left(P(x=n|Y=H) + P(X=n|Y=H) \right)$$

$$P(Y=+1 \mid X=x) = \frac{1}{2} \cdot \frac{1}{\sqrt{2\pi}} e^{-\frac{(x-5)^2}{2}} e^{-\frac{2}{2}} e^{-\frac{2}{2}$$

Given:
$$\min_{C_1,...,C_k} \sum_{i=1}^n \|x_i - c(x_i)\|_2^2$$



To Prove: K-means algorithm does not always find the optimal Solution to the above objective

Step 0: centeroid arrangements

1 A & D 2 B & C ...

Step 1: compute alyective

$$= 2 \cdot \|1 - 0\|_{L}^{2} = 2$$

Step 2: Optimal abjective = 2+2=4 = OPT

New consider a bad initialization:

Step 0: antoroid arrangements

Step 1: compute objective

 $= 2 \cdot || 1 \cdot 5||_{2}^{2} = 4 \cdot 5$

In this case, the next step of the algorithm does not change any point assignments because A&B are closer to (0,1) than (0,-1)

Then, when we recalculate the cluster contended, that doesn't change either because (0,1) is the center of A&B, same por C&D. Hence, the algorithm converges

New, we wish to generalize this example to P dimensions

n data points

k clusters

4 types of points:

(t+0.5,t)

$$(-t-0.5,-t)$$

Points =
$$\begin{cases} (-t+0.5, t) : t \in [1, T] \\ (-t-0.5, -t) \\ (t+0.5, -t) \end{cases}$$

This could work for any $t \geqslant 0$; 4T points; 2T duters
The optimal objective punction cost =

All the optimal clusters are the midpoints of the pain of points that have the same y-value & are symmetrically reflected across the y-anis; i.e.

Thus, optimal value of abjective would be:

$$\sum_{t=1}^{T} 2 \cdot t^2 = 2 \sum_{t=1}^{T} t^2$$

In the non-optimal case, centraids would be located:

The value of the objective function in this care

$$\frac{T}{\sum_{t=1}^{T}} 2 \cdot (t+0.5)^{2} = 2 \sum_{t=1}^{T} t^{2} + 2 \sum_{t=1}^{T} (0.25+t)$$

$$= 2 \sum_{t=1}^{T} t^{2} + \sum_{t=1}^{T} (t+0.5)$$

$$> 2 \sum_{t=1}^{T} t^{2}$$

$$> 2 \sum_{t=1}^{T} t^{2}$$

$$> 2 \sum_{t=1}^{T} t^{2}$$

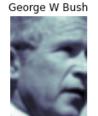
$$> 2 \sum_{t=1}^{T} t^{2}$$

Thus, since we have proved suboptimal objective is greater in this sotup, we have proved k-means is not ALWAYS optimal.

```
from sklearn.datasets import fetch_lfw_people
from sklearn.decomposition import PCA
import numpy as np
from matplotlib import pyplot as plt
```

```
faces_dataset = fetch_lfw_people(min_faces_per_person = 60)
print("Shape of Dataset: {}".format(faces_dataset.data.shape))
index = np.random.randint(0, faces_dataset.data.shape[0], size=16)
plt.rcParams['figure.figsize']=(10, 10)
i = 0
fig, axs = plt.subplots(4, 4)
for r in range(4):
   for c in range(4):
        axs[r, c].imshow(faces_dataset.images[index[i]], cmap=plt.cm.bone)
        axs[r, c].set_title(str(faces_dataset.target_names[faces_dataset.target[index[i]]]))
        axs[r, c].axis('off')
        i+=1
```

Shape of Dataset: (1348, 2914)



George W Bush



Ariel Sharon



George W Bush



George W Bush



Tony Blair



Gerhard Schroeder



George W Bush



George W Bush



Colin Powell



Colin Powell



Donald Rumsfeld



George W Bush



George W Bush



George W Bush



Tony Blair



PCA on the dataset to find the first 150 components, using randomized PCA from sklearn

```
faces_pca = PCA(n_components=150, whiten=True, svd_solver='randomized')
_ = faces_pca.fit(faces_dataset.data)
```

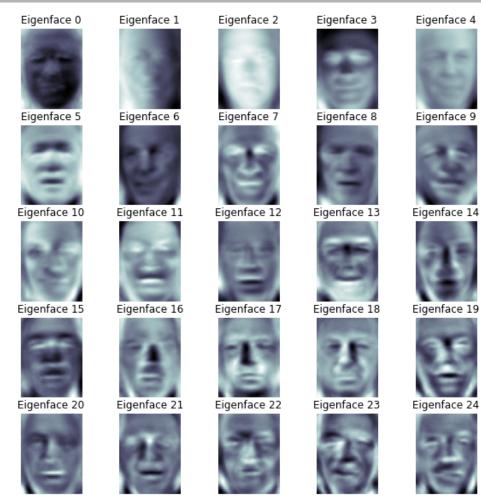
Eigenfaces associated with first 25 principal components

```
pca_eigenfaces = faces_pca.components_
plt.rcParams['figure.figsize']=(10, 10)

i = 0

fig, axs = plt.subplots(5, 5)

for r in range(5):
    axs[r, c].imshow(pca_eigenfaces[i].reshape(faces_dataset.images.shape[1:]), cmap=plt.cm.bone)
    axs[r, c].set_title(str("Eigenface {}").format(i))
    axs[r, c].axis('off')
    i+=1
```



```
plt.rcParams['figure.figsize']=(5.5, 10)
index = np.random.randint(0, faces_dataset.data.shape[0], size=5)

i = 0
fig, axs = plt.subplots(5, 2)
for r in range(5):
    c = 0
    axs[r, c].imshow(faces_dataset.images[index[i]], cmap=plt.cm.bone)
    axs[r, c].set_title(str(faces_dataset.target_names[faces_dataset.target[index[i]]]))
    axs[r, c].axis('off')

pca_transformed_face = faces_pca.transform(faces_dataset.images[index[i]].reshape(1, -1))
    axs[r, c+1].imshow(faces_pca.inverse_transform(pca_transformed_face).reshape(faces_dataset.images
    axs[r, c+1].set_title(str("Recreated: " + faces_dataset.target_names[faces_dataset.target[index[i]]])
    axs[r, c+1].axis('off')
    i+=1
```

Junichiro Koizumi

Recreated: Junichiro Koizumi



Gerhard Schroeder



Recreated: Gerhard Schroeder



George W Bush



Recreated: George W Bush



George W Bush



Recreated: George W Bush



Colin Powell



Recreated: Colin Powell



