

```
In [1]: def mean(a):
#5100 : row
#36 : colu
#a.shape 5100 36
# a.shape[1] 36
# a.shape[0] 5100
result=[]

a= np.array(a)
for i in range(a.shape[1]):
    sum = 0
    counter = 0
    for j in range(a.shape[0]):
        sum = sum + a[j][i]
        counter = counter + 1
    mean = sum/counter
    result.append(mean)
return np.array(result)
```

```
In [2]: def std(a):
a = np.array(a)
result=[]
mean1 = mean(a)
var=[]
for i in range(a.shape[1]):
    sum = 0
    counter = 0
    for j in range(a.shape[0]):
        sum = sum + (a[j][i] - mean1[i]) ** 2
        counter = counter + 1
    answer=sum / counter
    var.append(answer)
for i in range(len(var)):
    result.append(var[i] ** (1/2))
return result
```

```
In [3]: def flip(a):
answer=a[::-1]
return np.array(answer)
```

```
In [4]: def outer(a,b):
result=[]
for i in range(len(a)):
    result.append([])
for i in range(len(a)):
    answer = 0
    for j in range(len(a)):
        answer = a[i]*b[j]
        result[i].append(answer)
return np.array(result)
```

```
In [5]: def signfunction(a):
if a < 0:
    return -1
elif a == 0:
    return 0
else:
    return 1
```

```
In [6]: def normalise(a):
answer = 0
for i in range(len(a)):
```

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    answer = answer + a[i]**2
sqrt = answer ** (1/2)
return sqrt
```

```
In [7]: def sumfunction(a):
        sum = 0
        for i in range(len(a)):
            sum = sum + a[i]
        return sum
```

```
In [8]: def diag(v):
        res=[]
        for i in range(len(v)):
            for j in range(len(v[i])):
                if i == j:
                    res.append(v[i][j])
        return np.array(res)
```

```
In [9]: def zeros(n):
        answer = []
        for i in range(n):
            answer.append(0)
        return np.array(answer)
```

```
In [10]: def eyes(n):
        answer=[]
        for i in range(n):
            answer.append([])
        for i in range(n):
            for j in range(n):
                if i==j:
                    answer[i].append(1.)
                else:
                    answer[i].append(0.)
        return np.array(answer)
```

```
In [11]: def householder_reflection(a, e):
        '''
        Given a vector a and a unit vector e,
        (where a is non-zero and not collinear with e)
        returns an orthogonal matrix which maps a
        into the line of e.
        '''

        assert a.ndim == 1
        assert np.allclose(1, sumfunction(e**2))

        u = a - signfunction(a[0]) * normalise(a) * e
        v = u / normalise(u)
        H = eyes(len(a)) - 2 * outer(v, v)

        return H
```

```
In [12]: def qr_decomposition(A):
        '''
        Given an n x m invertable matrix A, returns the pair:
            Q an orthogonal n x m matrix
            R an upper triangular m x m matrix
        such that QR = A.
        '''
        n, m = A.shape
        assert n >= m
```

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Q = eyes(n)
R = A.copy()

for i in range(m - int(n==m)):
    r = R.iloc[i:, i]

    if np.allclose(r[1:], 0):
        continue

    # e is the i-th basis vector of the minor matrix.
    e = zeros(n-i)
    e[0] = 1

    H = eyes(n)
    H[i:, i:] = householder_reflection(r, e)
    Q = H.T
    R = H @ R
    return Q, R

```

```

In [13]: def eigen_decomposition(A, max_iter=100):
    A_k = A
    Q_k = eyes(A.shape[1])

    for k in range(max_iter):
        Q, R = qr_decomposition(A_k)
        Q_k = Q_k @ Q
        A_k = R @ Q

    eigenvalues = diag(A_k)
    eigenvectors = Q_k
    return eigenvalues, eigenvectors

```

```

In [14]: class PCA:
    def __init__(self, n_components=None, whiten=False):
        self.n_components = n_components
        self.whiten = bool(whiten)

    def fit(self, X):
        n, m = X.shape

        # subtract off the mean to center the data.
        self.mu = mean(X)
        X = X - self.mu

        # whiten if necessary
        if self.whiten:
            self.std = std(X)
            X = X / self.std

        # Eigen Decomposition of the covariance matrix
        C = X.T @ X / (n-1)
        self.eigenvalues, self.eigenvectors = eigen_decomposition(C)

        # truncate the number of components if doing dimensionality reduction
        if self.n_components is not None:
            self.eigenvalues = self.eigenvalues[0:self.n_components]
            self.eigenvectors = self.eigenvectors[:, 0:self.n_components]

        # the QR algorithm tends to puts eigenvalues in descending order
        # but is not guarenteed to. To make sure, we use argsort.
        descending_order = flip(np.argsort(self.eigenvalues))
        self.eigenvalues = self.eigenvalues[descending_order]
        self.eigenvectors = self.eigenvectors[:, descending_order]

    return self

```

```

def transform(self, X):
    X = X - self.mu

    if self.whiten:
        X = X / self.std

    return X @ self.eigenvectors

@property
def proportion_variance_explained(self):
    return self.eigenvalues / sumfunction(self.eigenvalues)

return Q, R

```

```

In [15]: import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
from mpl_toolkits.mplot3d import Axes3D

# Importing Data file

X = pd.read_csv('DataSet_2.csv')
display(X.head())

```

	V1	V2	V3	V4	V5	V6	V7	V8	V9	V10	...	V27	V28	V29	V30	V31	V32	V33	V34	V35	V36
0	46	40	119	139	42	30	135	157	42	30	...	115	113	50	46	111	116	44	31	131	142
1	47	37	119	133	44	34	124	143	44	34	...	100	85	50	39	118	132	43	29	133	143
2	80	95	100	74	64	64	104	96	46	36	...	100	81	82	91	92	78	78	83	96	74
3	56	51	72	60	59	54	72	60	59	51	...	71	50	57	55	74	61	57	55	78	65
4	44	34	129	140	44	34	124	136	44	34	...	133	139	43	31	128	135	43	29	128	132

5 rows × 36 columns

```

In [16]: # Calculating Co-variance Matrix

X_white = (X - mean(X))/std(X)
C = X_white.T @ X_white / (X_white.shape[0] - 1)

# Visualizing Co-variance Matrix

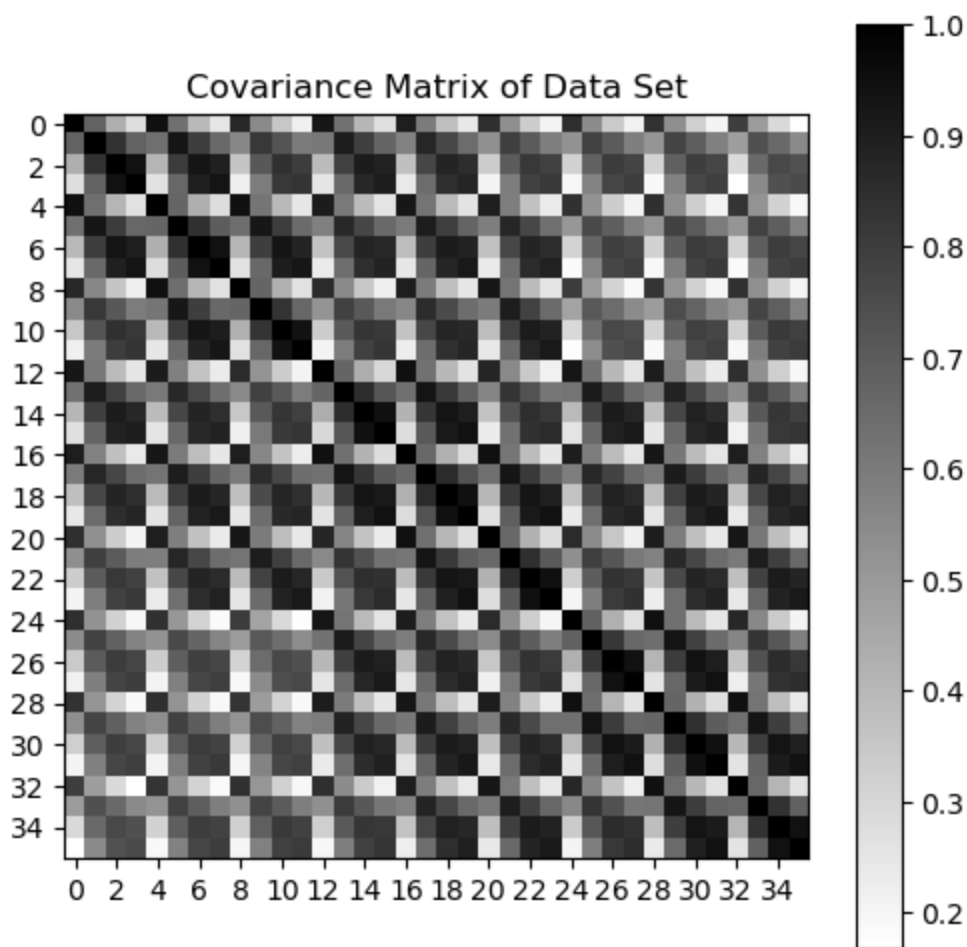
plt.figure(figsize=(6,6))
plt.imshow(C, cmap='binary')
plt.title("Covariance Matrix of Data Set")
plt.xticks(np.arange(0, 36, 2))
plt.yticks(np.arange(0, 36, 2))
plt.colorbar()

```

```

Out[16]: <matplotlib.colorbar.Colorbar at 0x2a7b80234d0>

```

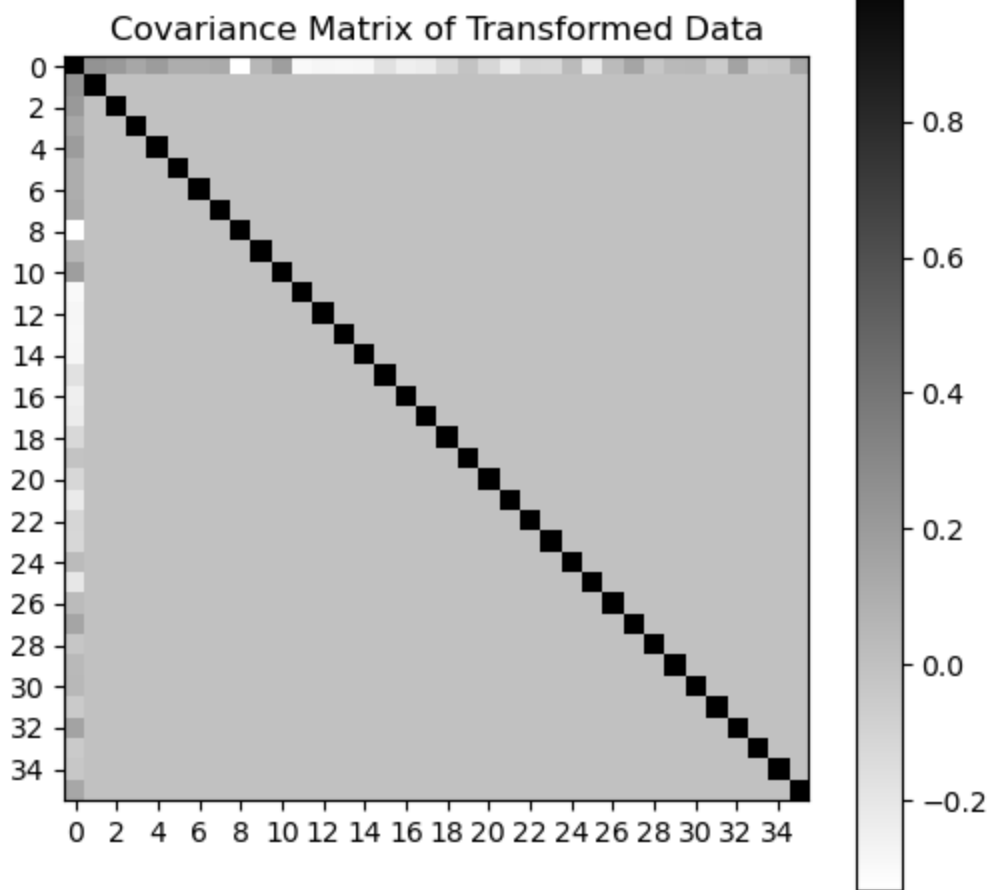


In [17]: *# Calculating*

```
pca = PCA(whiten=True)
pca.fit(X)
X_prime = pca.transform(X)
pca.eigenvalues
C_prime = pca.eigenvectors
X_white = (C_prime - mean(C_prime))/std(C_prime)
C_prime = X_white.T @ X_white / (X_white.shape[0] - 1)

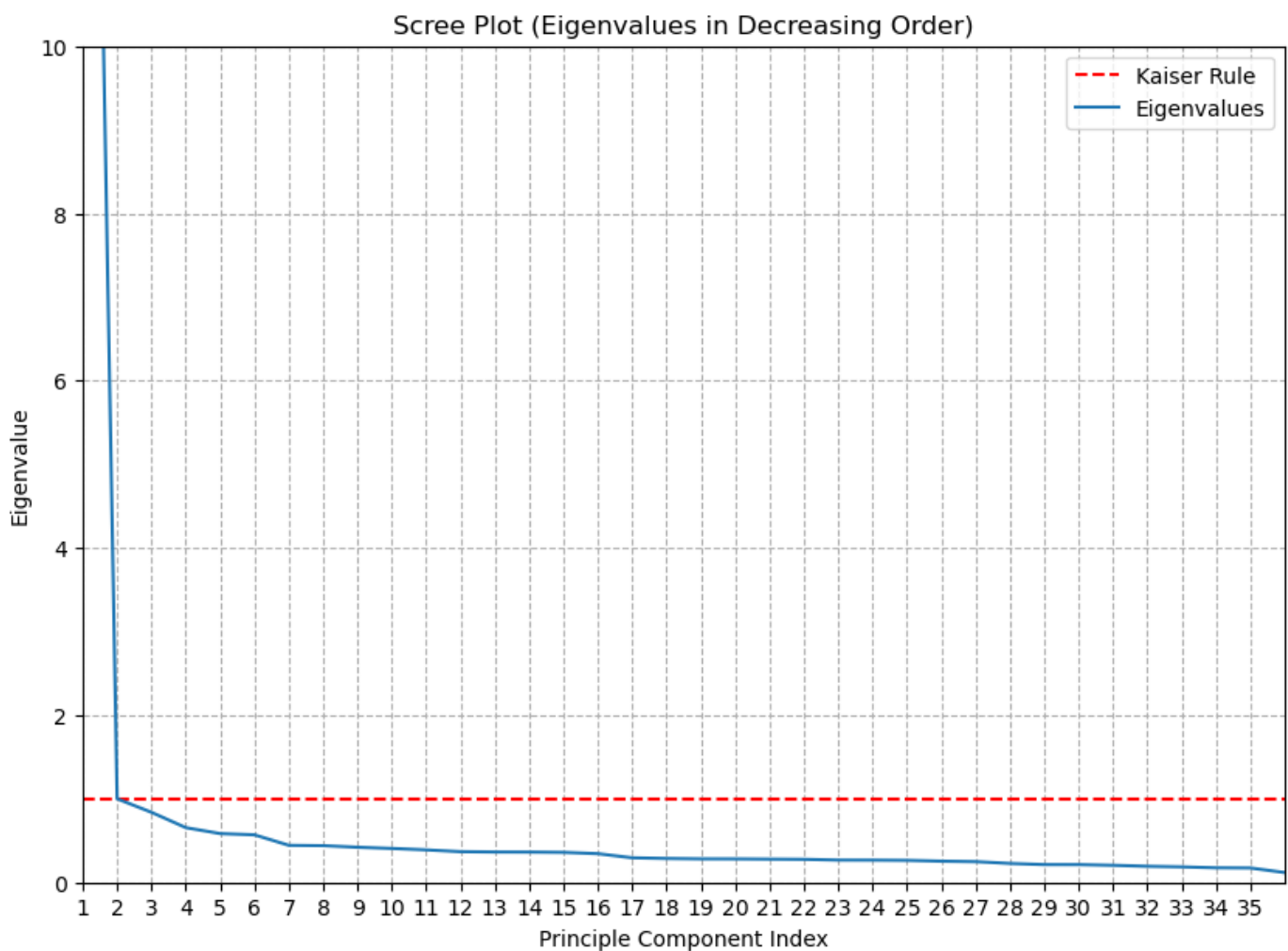
# Visualizing Co-Variance Matrix of the Transformed Data

plt.figure(figsize=(6,6))
plt.imshow(C_prime, cmap='binary')
plt.title("Covariance Matrix of Transformed Data")
plt.xticks(np.arange(0, 36, 2))
plt.yticks(np.arange(0, 36, 2))
plt.colorbar()
plt.show()
```



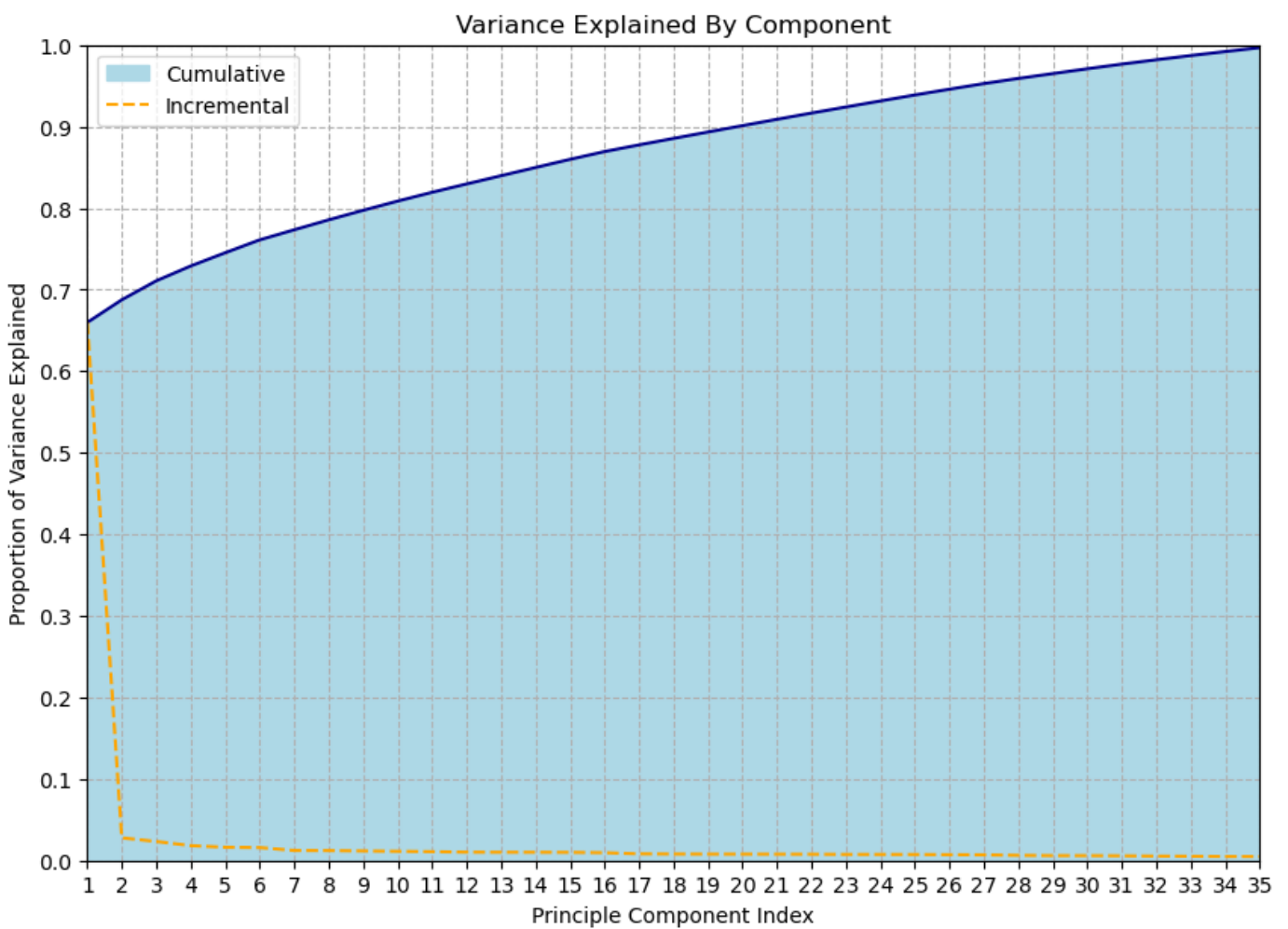
In [18]: *# Plotting Eigen Vectors*

```
fig = plt.figure(figsize=(10, 7))
plt.title("Scree Plot (Eigenvalues in Decreasing Order)")
plt.plot([1, 36], [1, 1], color='red', linestyle='--', label="Kaiser Rule")
plt.xticks(np.arange(1, 36, 1))
plt.xlim(1, 36)
plt.ylim(0, 10)
plt.ylabel("Eigenvalue")
plt.xlabel("Principle Component Index")
plt.grid(linestyle='--')
plt.plot(range(1, 37), pca.eigenvalues, label="Eigenvalues")
plt.legend()
plt.show()
```



```
In [19]: # Variance covered by the Eigen Vectors

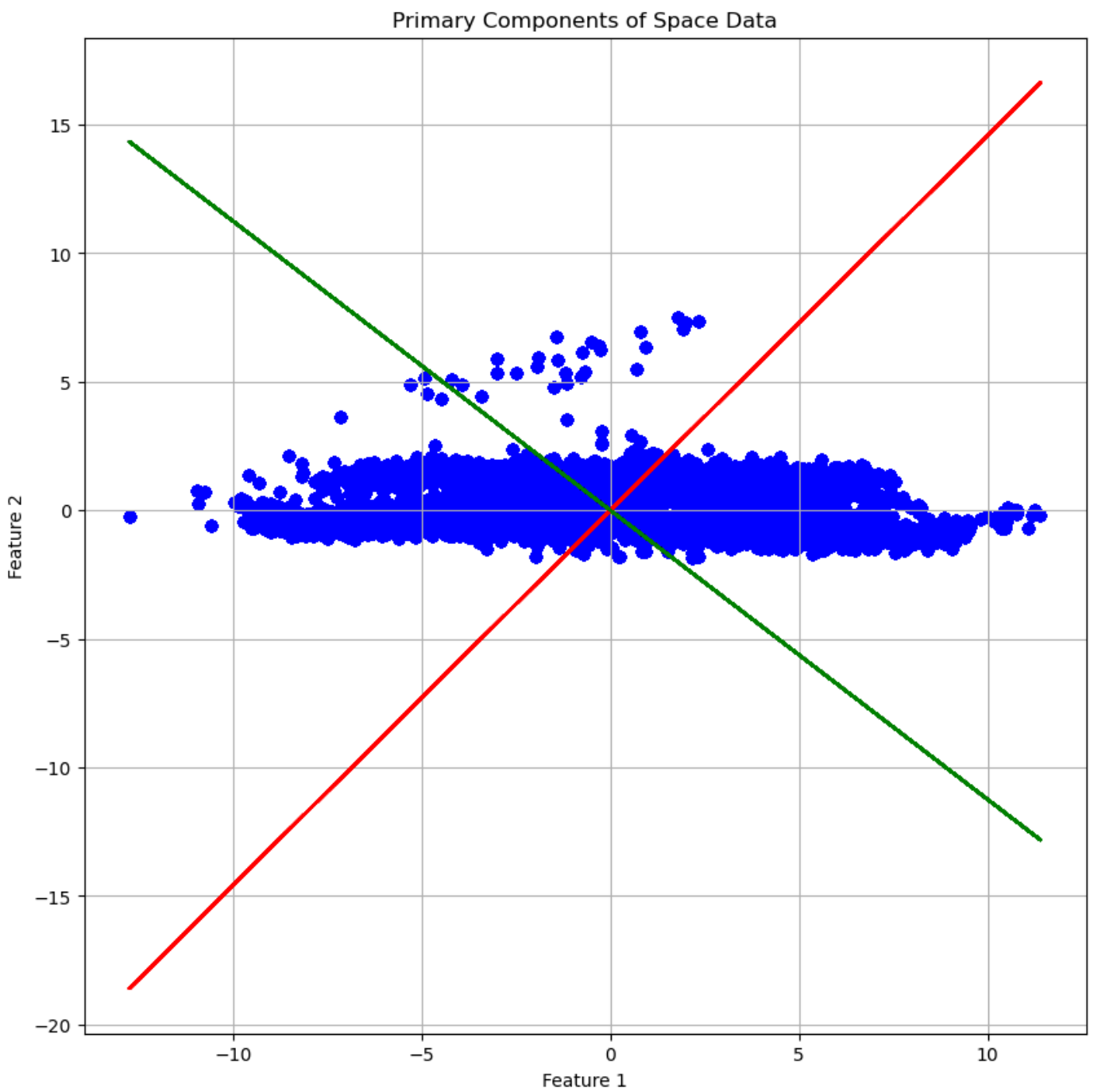
fig = plt.figure(figsize=(10, 7))
plt.title("Variance Explained By Component")
plt.xticks(np.arange(1, 36, 1))
plt.yticks(np.arange(0, 1.0001, 0.1))
plt.xlim(1, 35)
plt.ylim(0, 1)
plt.ylabel("Proportion of Variance Explained")
plt.xlabel("Principle Component Index")
plt.grid(linestyle='--')
plt.fill_between(
    range(1, 37),
    np.cumsum(pca.proportion_variance_explained),
    0,
    color="lightblue",
    label="Cumulative")
plt.plot(
    range(1, 37),
    np.cumsum(pca.proportion_variance_explained),
    0,
    color="darkblue")
plt.plot(
    range(1, 37),
    pca.proportion_variance_explained,
    label="Incremental",
    color="orange",
    linestyle="--")
plt.legend(loc='upper left')
plt.show()
```



```
In [20]: # Visualizing Eigen Vectors and Tranformed Data

plt.figure(figsize=(10, 4))
x_data = X.iloc[:,0]
y_data= (pca.eigenvectors[1,0]/pca.eigenvectors[0,0])*X_prime.iloc[:,0]
y_data1= (pca.eigenvectors[1,1]/pca.eigenvectors[0,1])*X_prime.iloc[:,0]
fig = plt.figure(figsize=(10, 10))
for c in np.unique(X):
    X_class = X_prime
    plt.plot(X_prime.iloc[:,0],y_data,color='red')
    plt.plot(X_prime.iloc[:,0],y_data1,color='green')
    plt.scatter(X_class.iloc[:,0],X_class.iloc[:,1],alpha=0.1,color='blue')
plt.title("Primary Components of Space Data")
plt.xlabel("Feature 1")
plt.ylabel("Feature 2")
plt.grid()
plt.show()
```

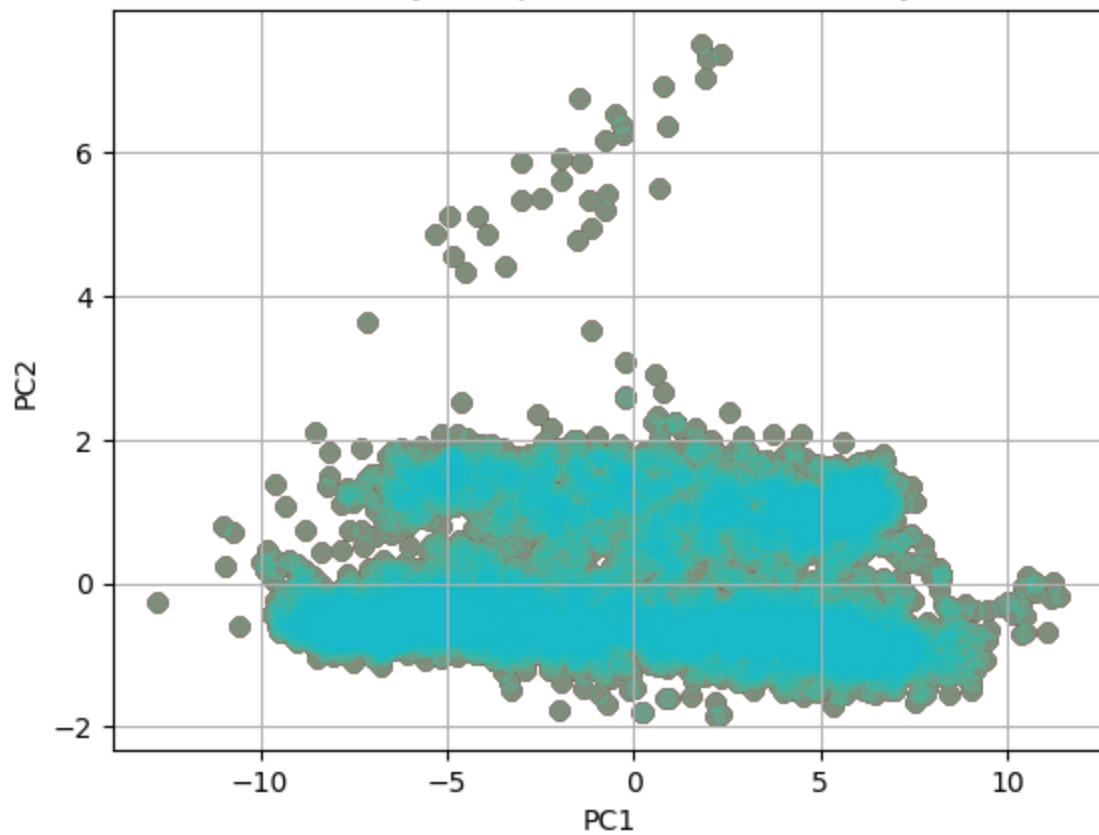
<Figure size 1000x400 with 0 Axes>



In [21]: *# 2D Scatter plot of Tranformed Data*

```
for c in np.unique(X):
    X_class = X_prime
    plt.scatter(X_class.iloc[:, 0], X_class.iloc[:, 1], alpha=0.2)
plt.title("Primary Components of Wine Quality")
plt.xlabel("PC1")
plt.ylabel("PC2")
plt.grid()
plt.show()
```

Primary Components of Wine Quality



```
In [22]: # 3D Scatter plot of Tranformed Data

fig = plt.figure(figsize=(10, 8))
ax = fig.add_subplot(111, projection='3d')
ax.view_init(15, -45)
for c in np.unique(X):
    X_class = X_prime
    ax.scatter(X_class.iloc[:, 0], X_class.iloc[:, 2], X_class.iloc[:, 1], alpha=0.2)
plt.title("First 3 Primary Components of Wine Quality")
ax.set_xlabel('PC1')
ax.set_ylabel('PC2')
ax.set_zlabel('PC3')
plt.show()
```

First 3 Primary Components of Wine Quality

