# pca-cancer-dataset

December 19, 2023

[106]:
```
import matplotlib.pyplot as plt
import pandas as pd
import numpy as np
import seaborn as sns

from sklearn.preprocessing import StandardScaler
from sklearn.decomposition import PCA
```

[107]:
```
df=pd.read_csv("/kaggle/input/breast-cancer-wisconsin-data/data.csv")
df.head()
```

[107]:

| | id | diagnosis | radius_mean | texture_mean | perimeter_mean | area_mean | \ |
|---|---|---|---|---|---|---|---|
| 0 | 842302 | M | 17.99 | 10.38 | 122.80 | 1001.0 | |
| 1 | 842517 | M | 20.57 | 17.77 | 132.90 | 1326.0 | |
| 2 | 84300903 | M | 19.69 | 21.25 | 130.00 | 1203.0 | |
| 3 | 84348301 | M | 11.42 | 20.38 | 77.58 | 386.1 | |
| 4 | 84358402 | M | 20.29 | 14.34 | 135.10 | 1297.0 | |

| | smoothness_mean | compactness_mean | concavity_mean | concave points_mean | \ |
|---|---|---|---|---|---|
| 0 | 0.11840 | 0.27760 | 0.3001 | 0.14710 | |
| 1 | 0.08474 | 0.07864 | 0.0869 | 0.07017 | |
| 2 | 0.10960 | 0.15990 | 0.1974 | 0.12790 | |
| 3 | 0.14250 | 0.28390 | 0.2414 | 0.10520 | |
| 4 | 0.10030 | 0.13280 | 0.1980 | 0.10430 | |

| | … | texture_worst | perimeter_worst | area_worst | smoothness_worst | \ |
|---|---|---|---|---|---|---|
| 0 | … | 17.33 | 184.60 | 2019.0 | 0.1622 | |
| 1 | … | 23.41 | 158.80 | 1956.0 | 0.1238 | |
| 2 | … | 25.53 | 152.50 | 1709.0 | 0.1444 | |
| 3 | … | 26.50 | 98.87 | 567.7 | 0.2098 | |
| 4 | … | 16.67 | 152.20 | 1575.0 | 0.1374 | |

| | compactness_worst | concavity_worst | concave points_worst | symmetry_worst | \ |
|---|---|---|---|---|---|
| 0 | 0.6656 | 0.7119 | 0.2654 | 0.4601 | |
| 1 | 0.1866 | 0.2416 | 0.1860 | 0.2750 | |
| 2 | 0.4245 | 0.4504 | 0.2430 | 0.3613 | |
| 3 | 0.8663 | 0.6869 | 0.2575 | 0.6638 | |

```
4             0.2050           0.4000                0.1625          0.2364

   fractal_dimension_worst  Unnamed: 32
0                  0.11890          NaN
1                  0.08902          NaN
2                  0.08758          NaN
3                  0.17300          NaN
4                  0.07678          NaN

[5 rows x 33 columns]
```

[108]:
```
df.shape
```

[108]:
```
(569, 33)
```

[109]:
```
df.info()
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 569 entries, 0 to 568
Data columns (total 33 columns):
 #   Column                   Non-Null Count  Dtype
---  ------                   --------------  -----
 0   id                       569 non-null    int64
 1   diagnosis                569 non-null    object
 2   radius_mean              569 non-null    float64
 3   texture_mean             569 non-null    float64
 4   perimeter_mean           569 non-null    float64
 5   area_mean                569 non-null    float64
 6   smoothness_mean          569 non-null    float64
 7   compactness_mean         569 non-null    float64
 8   concavity_mean           569 non-null    float64
 9   concave points_mean      569 non-null    float64
 10  symmetry_mean            569 non-null    float64
 11  fractal_dimension_mean   569 non-null    float64
 12  radius_se                569 non-null    float64
 13  texture_se               569 non-null    float64
 14  perimeter_se             569 non-null    float64
 15  area_se                  569 non-null    float64
 16  smoothness_se            569 non-null    float64
 17  compactness_se           569 non-null    float64
 18  concavity_se             569 non-null    float64
 19  concave points_se        569 non-null    float64
 20  symmetry_se              569 non-null    float64
 21  fractal_dimension_se     569 non-null    float64
 22  radius_worst             569 non-null    float64
 23  texture_worst            569 non-null    float64
 24  perimeter_worst          569 non-null    float64
```

```
 25  area_worst               569 non-null    float64
 26  smoothness_worst         569 non-null    float64
 27  compactness_worst        569 non-null    float64
 28  concavity_worst          569 non-null    float64
 29  concave points_worst     569 non-null    float64
 30  symmetry_worst           569 non-null    float64
 31  fractal_dimension_worst  569 non-null    float64
 32  Unnamed: 32              0 non-null      float64
dtypes: float64(31), int64(1), object(1)
memory usage: 146.8+ KB
```

[110]:
```
# Drop unnecessary columns
df.drop(["Unnamed: 32","id"],axis=1,inplace=True)
df.head()
```

[110]:

| | diagnosis | radius_mean | texture_mean | perimeter_mean | area_mean | \ |
|---|---|---|---|---|---|---|
| 0 | M | 17.99 | 10.38 | 122.80 | 1001.0 | |
| 1 | M | 20.57 | 17.77 | 132.90 | 1326.0 | |
| 2 | M | 19.69 | 21.25 | 130.00 | 1203.0 | |
| 3 | M | 11.42 | 20.38 | 77.58 | 386.1 | |
| 4 | M | 20.29 | 14.34 | 135.10 | 1297.0 | |

| | smoothness_mean | compactness_mean | concavity_mean | concave points_mean | \ |
|---|---|---|---|---|---|
| 0 | 0.11840 | 0.27760 | 0.3001 | 0.14710 | |
| 1 | 0.08474 | 0.07864 | 0.0869 | 0.07017 | |
| 2 | 0.10960 | 0.15990 | 0.1974 | 0.12790 | |
| 3 | 0.14250 | 0.28390 | 0.2414 | 0.10520 | |
| 4 | 0.10030 | 0.13280 | 0.1980 | 0.10430 | |

| | symmetry_mean | … | radius_worst | texture_worst | perimeter_worst | \ |
|---|---|---|---|---|---|---|
| 0 | 0.2419 | … | 25.38 | 17.33 | 184.60 | |
| 1 | 0.1812 | … | 24.99 | 23.41 | 158.80 | |
| 2 | 0.2069 | … | 23.57 | 25.53 | 152.50 | |
| 3 | 0.2597 | … | 14.91 | 26.50 | 98.87 | |
| 4 | 0.1809 | … | 22.54 | 16.67 | 152.20 | |

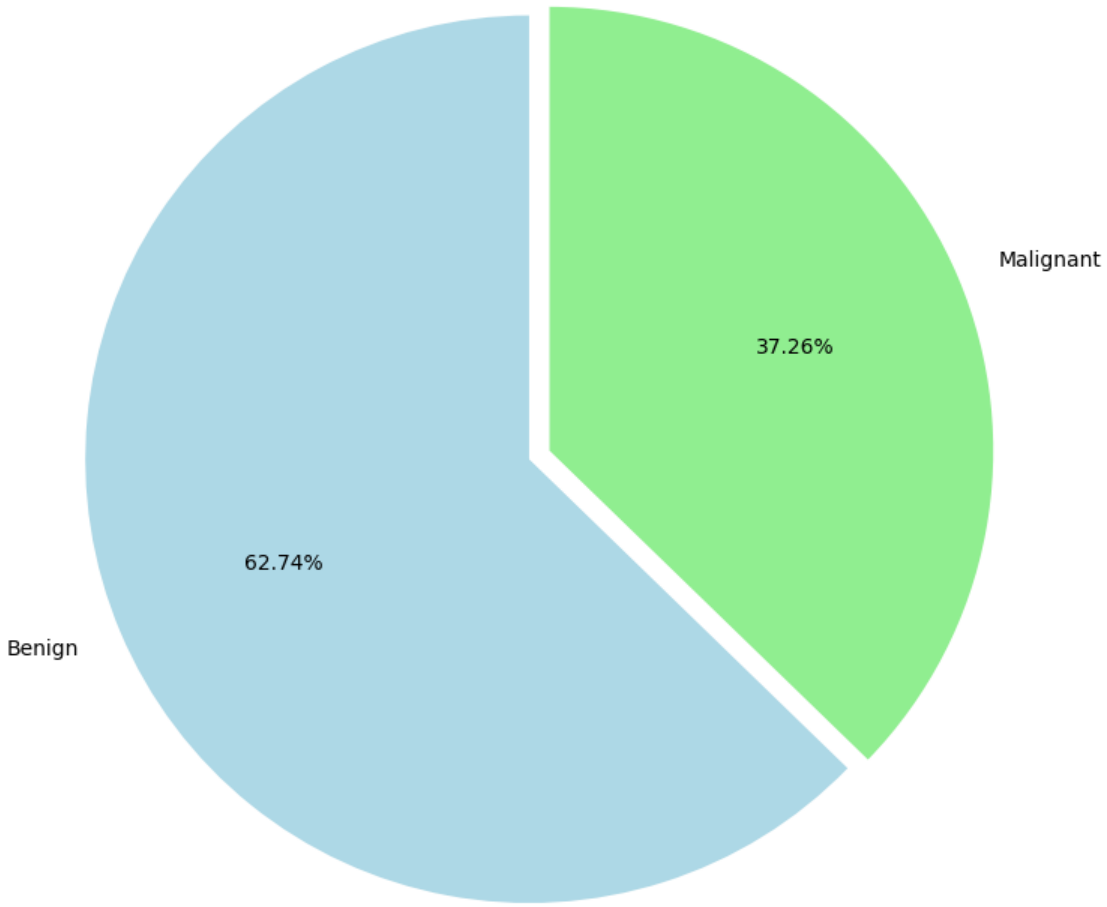
| | area_worst | smoothness_worst | compactness_worst | concavity_worst | \ |
|---|---|---|---|---|---|
| 0 | 2019.0 | 0.1622 | 0.6656 | 0.7119 | |
| 1 | 1956.0 | 0.1238 | 0.1866 | 0.2416 | |
| 2 | 1709.0 | 0.1444 | 0.4245 | 0.4504 | |
| 3 | 567.7 | 0.2098 | 0.8663 | 0.6869 | |
| 4 | 1575.0 | 0.1374 | 0.2050 | 0.4000 | |

| | concave points_worst | symmetry_worst | fractal_dimension_worst |
|---|---|---|---|
| 0 | 0.2654 | 0.4601 | 0.11890 |
| 1 | 0.1860 | 0.2750 | 0.08902 |
| 2 | 0.2430 | 0.3613 | 0.08758 |

```
3                0.2575          0.6638                0.17300
4                0.1625          0.2364                0.07678

[5 rows x 31 columns]
```

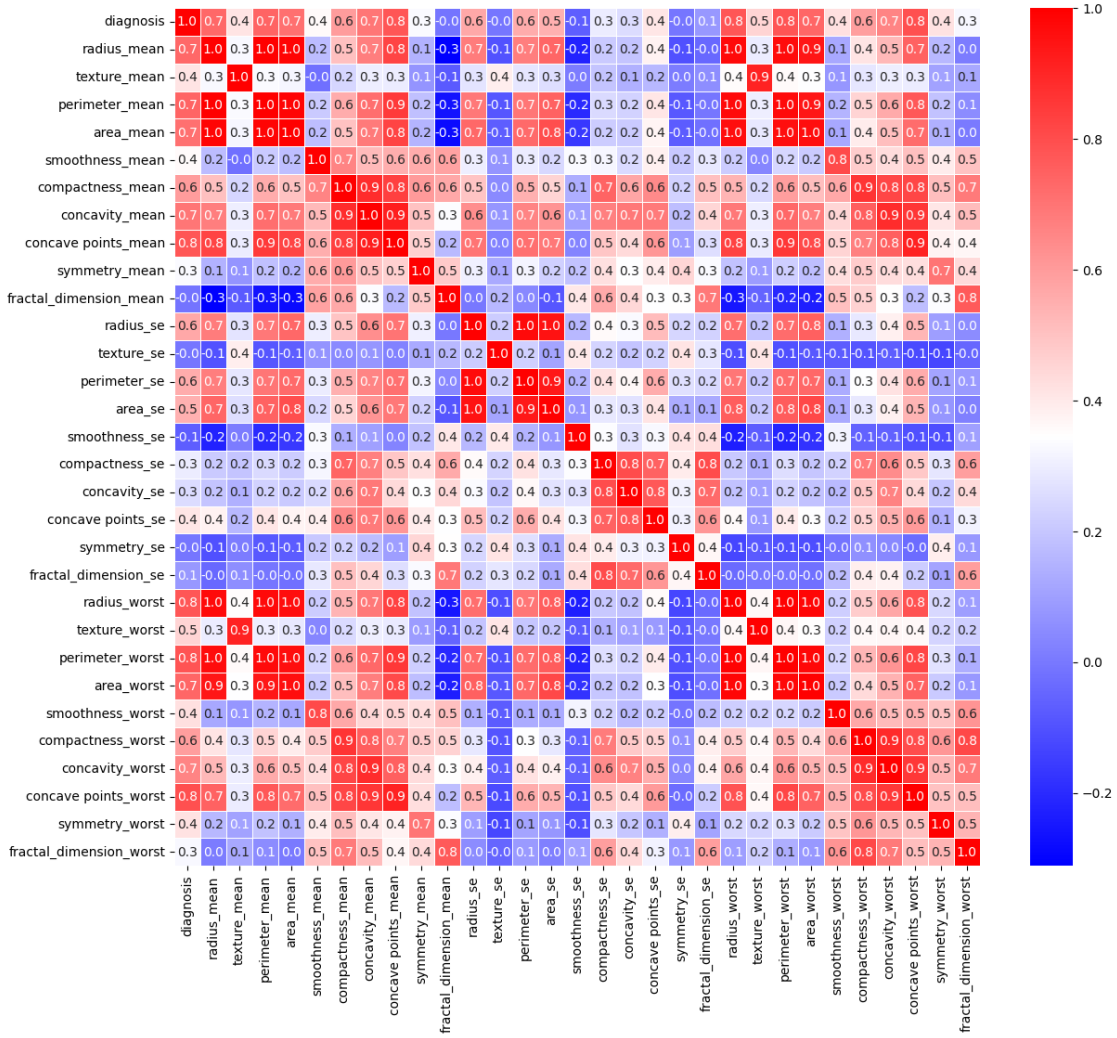
[111]:
```
# Categorical data convert to numeric data
df["diagnosis"] = [
    1 if item == "M"
    else 0  for item in df["diagnosis"]]
```

[112]:
```
# Pie chart
plt.pie(df.diagnosis.value_counts(), startangle=90, explode=[0.05, 0.05],␣
↪autopct='%0.2f%%',
        labels=['Benign', 'Malignant'], colors=['#add8e6', '#90ee90'], radius=2)
plt.show()
```




### Correlation Matrix

[113]:
```
import seaborn as sns
f,ax = plt.subplots(figsize=(14,12))
sns.heatmap(df.corr(), cmap="bwr", annot=True, linewidths=0.5, fmt= '.1f',ax=ax)
plt.show()
```




## PCA (Principal Component Analysis)

[114]:
```
from sklearn.preprocessing import StandardScaler
Y = df["diagnosis"]
X = df.drop('diagnosis', axis=1)
```

[115]:
```
sc = StandardScaler()
X = sc.fit_transform(X)
X.shape
```

[115]: (569, 30)

[116]:
```
#PCA
from sklearn.decomposition import PCA
n_components = 3
pca = PCA(n_components=n_components)
pca.fit(X)
components = pca.transform(X)
X.shape
```

[116]: (569, 30)

[117]:
```
components.shape
```

[117]: (569, 3)

[118]:
```
pca.explained_variance_ratio_
np.cumsum(pca.explained_variance_ratio_)
```

[118]: array([0.44272026, 0.63243208, 0.72636371])

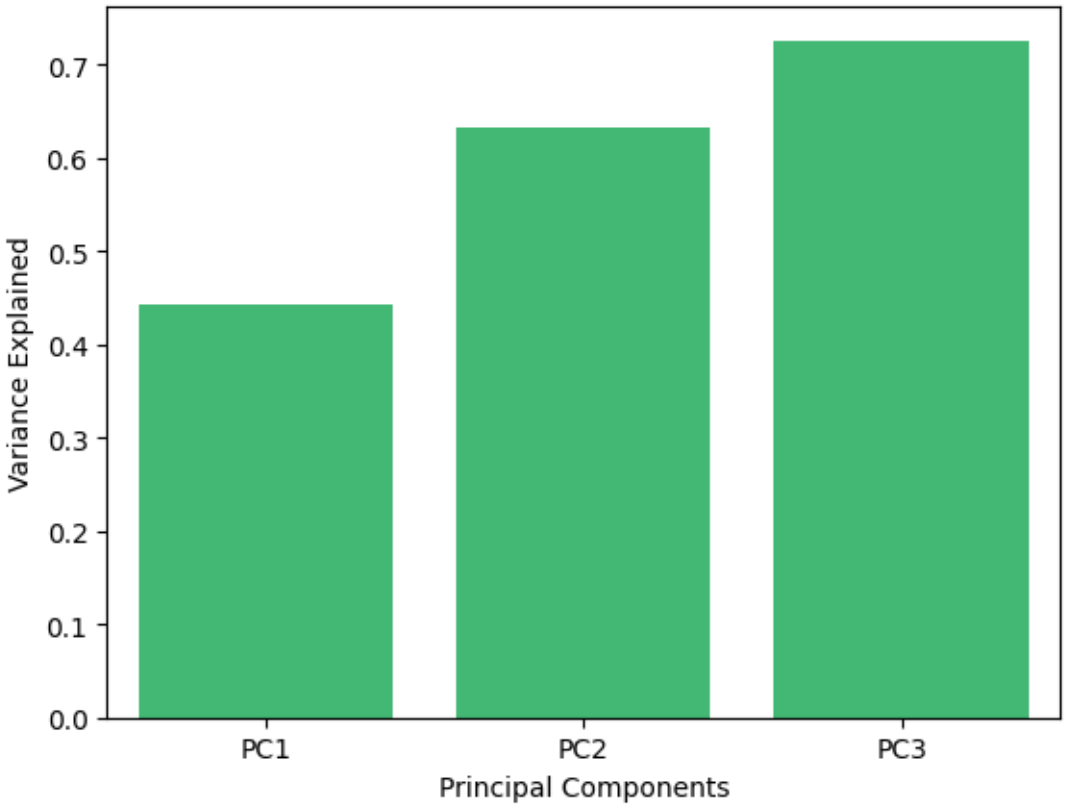
[119]:
```
df_pca=pd.DataFrame({"PC":["PC1","PC2","PC3"],
                     "var": np.cumsum(pca.explained_variance_ratio_)})

df_pca
```

[119]:

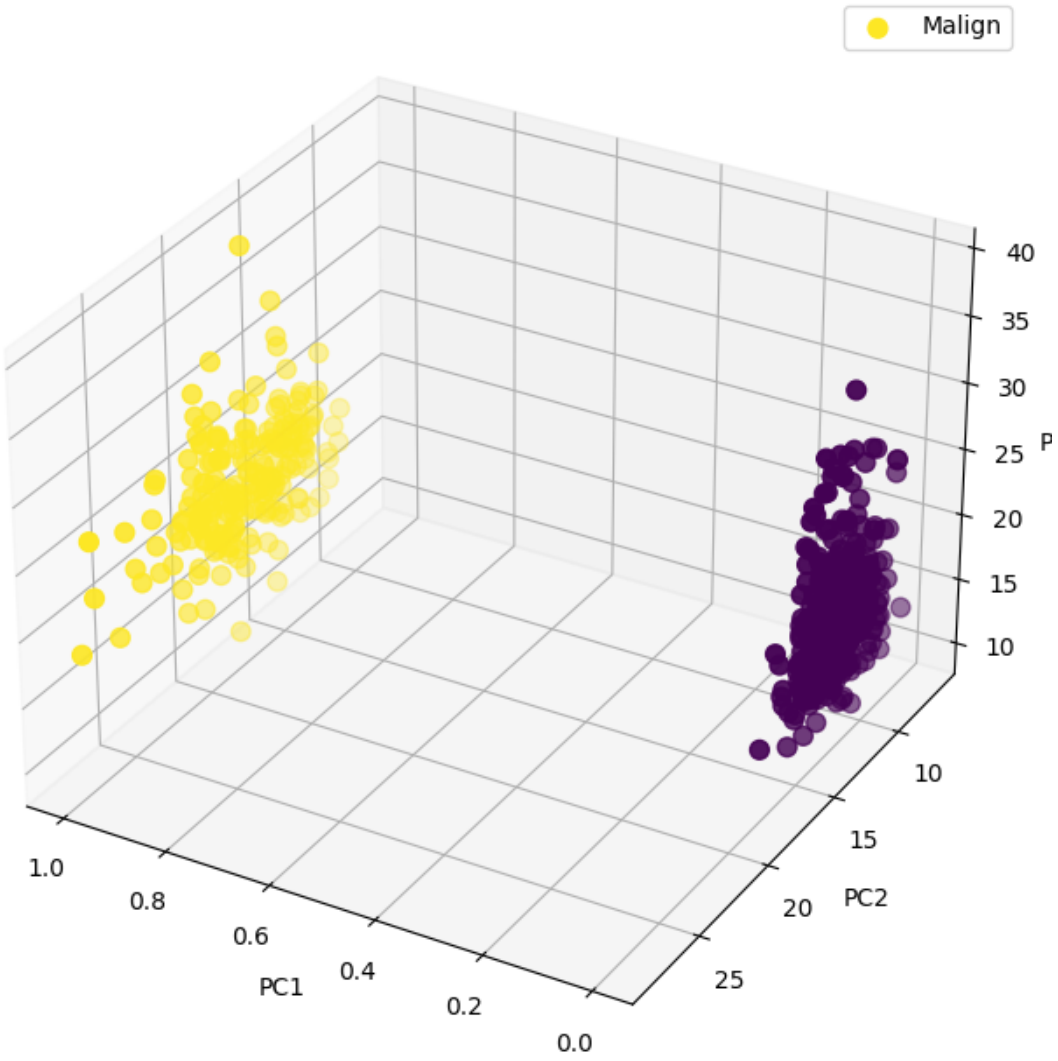
| | PC | var |
|---|---|---|
| 0 | PC1 | 0.442720 |
| 1 | PC2 | 0.632432 |
| 2 | PC3 | 0.726364 |

### PCA Visualization

[120]:
```
sns.barplot(x="PC", y="var", data=df_pca, color="#2ecc71")
plt.ylabel("Variance Explained")
plt.xlabel("Principal Components")
plt.show()
```

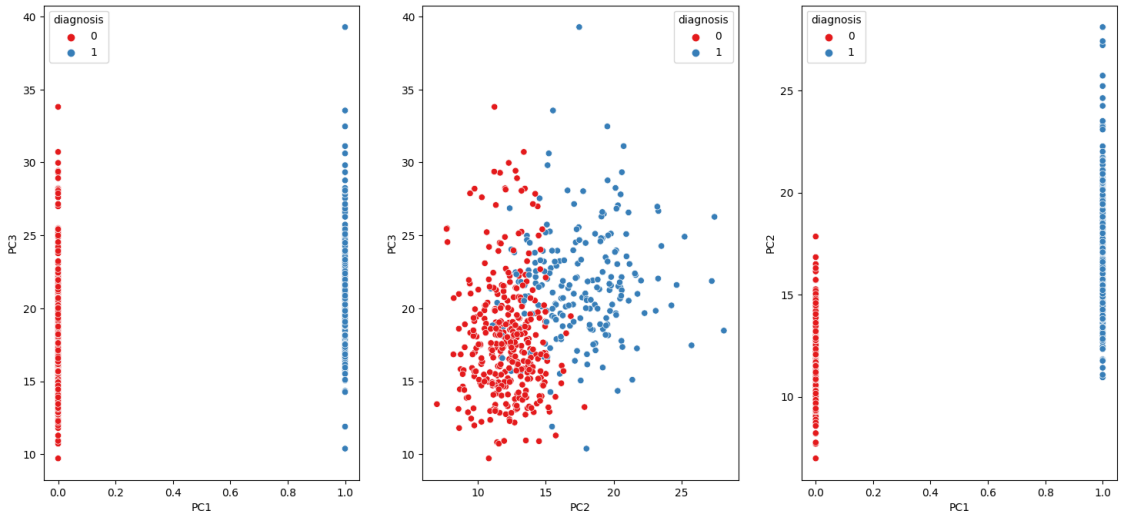



[121]:
```
from mpl_toolkits.mplot3d import Axes3D
fig = plt.figure(figsize=(15, 8))
ax = fig.add_subplot(111, projection='3d')
ax.scatter(df.iloc[:, 0], df.iloc[:, 1], df.iloc[:, 2], c=df['diagnosis'], s=60)
ax.legend(['Malign'])
ax.set_xlabel('PC1')
ax.set_ylabel('PC2')
ax.set_zlabel('PC3')
ax.view_init(30, 120)
```




[122]:
```
# First subplot
plt.figure(figsize=(18, 8))
plt.subplot(1, 3, 1)
sns.scatterplot(x=df.iloc[:, 0], y=df.iloc[:, 2], hue=df['diagnosis'],␣
 ↪palette='Set1')
plt.xlabel('PC1')
plt.ylabel('PC3')

# Second subplot
plt.subplot(1, 3, 2)
sns.scatterplot(x=df.iloc[:, 1], y=df.iloc[:, 2], hue=df['diagnosis'],␣
 ↪palette='Set1')
plt.xlabel('PC2')
```

```
plt.ylabel('PC3')


# 3rd subplot
plt.subplot(1, 3, 3)
sns.scatterplot(x=df.iloc[:, 0], y=df.iloc[:, 1], hue=df['diagnosis'],␣
 ↪palette='Set1')
plt.xlabel('PC1')
plt.ylabel('PC2')

plt.show()
```




[123]:
```
pca = PCA(n_components=3)  # Set the appropriate number of components
pca.fit(df)

# Create a DataFrame for principal components
df_pc = pd.DataFrame(pca.components_, columns=df.columns)

# Display the DataFrame
df_pc
```

[123]:

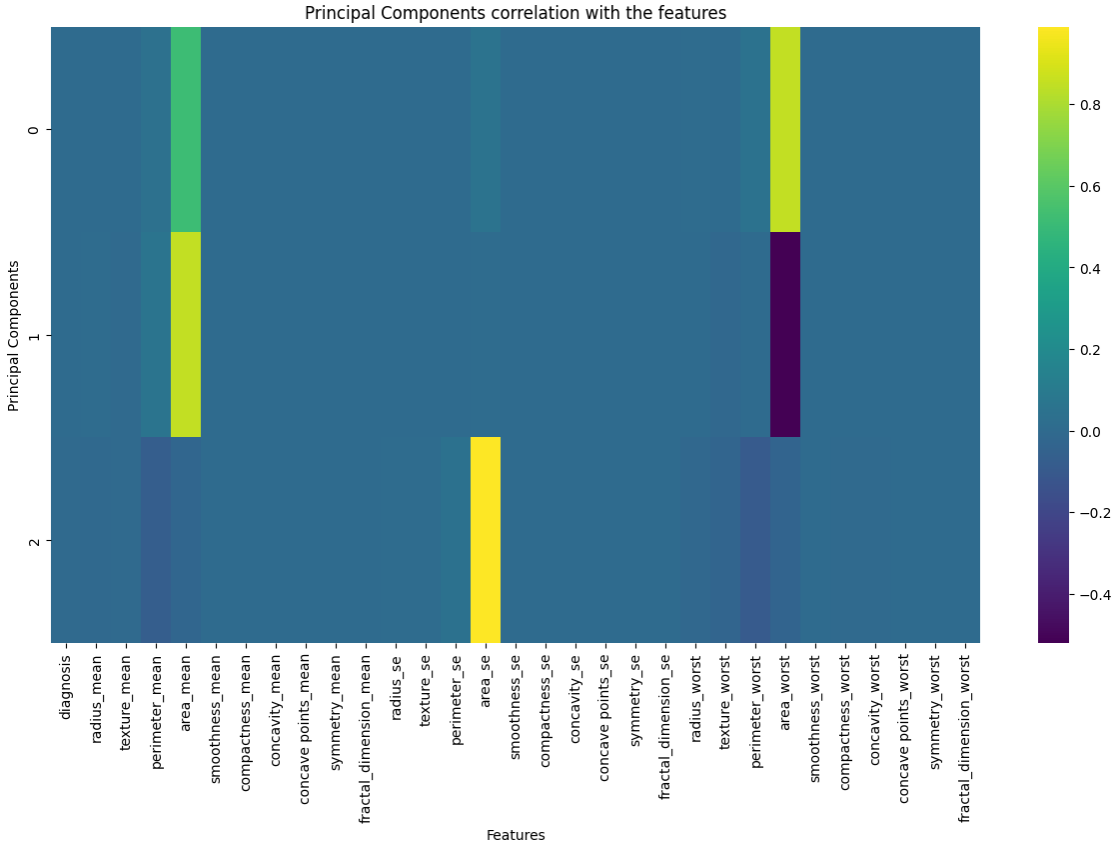
| | diagnosis | radius_mean | texture_mean | perimeter_mean | area_mean | \ |
|---|---|---|---|---|---|---|
| 0 | 0.000532 | 0.005086 | 0.002197 | 0.035076 | 0.516826 | |
| 1 | -0.000220 | 0.009287 | -0.002882 | 0.062748 | 0.851824 | |
| 2 | -0.001755 | -0.012343 | -0.006356 | -0.071671 | -0.027894 | |

| | smoothness_mean | compactness_mean | concavity_mean | concave points_mean | \ |
|---|---|---|---|---|---|
| 0 | 0.000004 | 0.000041 | 0.000082 | 0.000048 | |
| 1 | -0.000015 | -0.000003 | 0.000075 | 0.000046 | |

```
2         0.000073          0.000102        0.000266              0.000036

   symmetry_mean  …  radius_worst  texture_worst  perimeter_worst  \
0       0.000007  …      0.007155       0.003067         0.049458
1      -0.000025  …     -0.000569      -0.013215        -0.000186
2       0.000141  …     -0.015566      -0.031546        -0.092316

   area_worst  smoothness_worst  compactness_worst  concavity_worst  \
0    0.852063          0.000006           0.000101         0.000169
1   -0.519742         -0.000077          -0.000256        -0.000175
2   -0.039317         -0.000042          -0.000765        -0.000847

   concave points_worst  symmetry_worst  fractal_dimension_worst
0              0.000074        0.000018                 0.000002
1             -0.000031       -0.000157                -0.000055
2             -0.000334       -0.000350                -0.000041

[3 rows x 31 columns]
```

[124]:
```
plt.figure(figsize=(15, 8))
sns.heatmap(df_pc, cmap='viridis')
plt.title('Principal Components correlation with the features')
plt.xlabel('Features')
plt.ylabel('Principal Components')
```

[124]:
```
Text(158.22222222222223, 0.5, 'Principal Components')
```


Principal Components correlation with the features
Principal Components
0
1
2
Features
diagnosis
radius_mean
texture_mean
perimeter_mean
area_mean
smoothness_mean
compactness_mean
concavity_mean
concave points_mean
symmetry_mean
fractal_dimension_mean
radius_se
texture_se
perimeter_se
area_se
smoothness_se
compactness_se
concavity_se
concave points_se
symmetry_se
fractal_dimension_se
radius_worst
texture_worst
perimeter_worst
area_worst
smoothness_worst
compactness_worst
concavity_worst
concave points_worst
symmetry_worst
fractal_dimension_worst
0.8
0.6
0.4
0.2
0.0
−0.2
−0.4