Objective: To understand the importance of scaling on PCA

```
In [8]: from sklearn.decomposition import PCA
    from sklearn import preprocessing
    from sklearn import metrics
    import numpy as np
    import matplotlib.pyplot as plt
    from sklearn.datasets import load_wine
```

Task 0: Write the function to compute the pca using Eigenvector approach

```
In [9]: from numpy.linalg import svd
def pca(X):
    U, S, P_trans = svd(X, full_matrices = False)
    Sigma = np.diag(S)
    T = np.dot(U,Sigma)
    P = P_trans.T
    return T, Sigma, P #Score, Variace, Loadings

In [10]: features, target = load_wine(return_X_y=True)
    X=features
    y=target
```

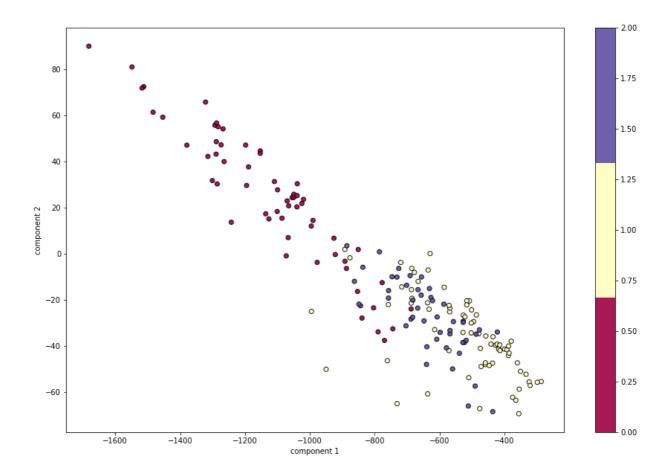
Three different ways of scaling

- Scaling by removing the mean and divising by the standard deviation #standard_scaling=preprocessing.StandardScaler() #X_standard=standard_scaling.fit_transform(X)
- Scaling to min and maximum values of each feature #minmax_scaling=preprocessing.MinMaxScaler() #X_minmax=minmax_scaling.fit_transform(X)

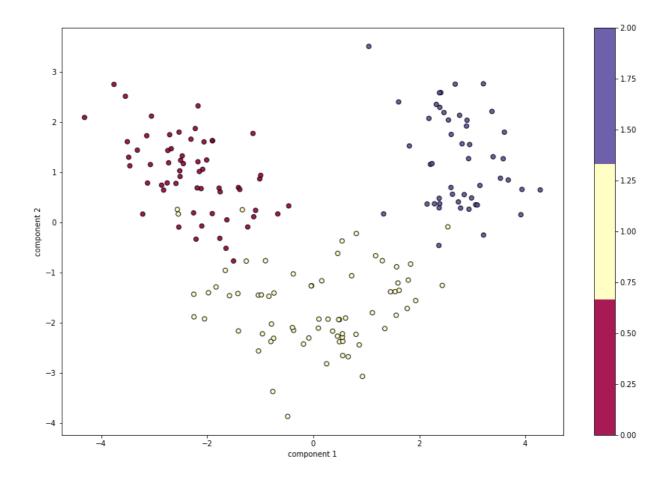
(178, 13)

 Scaling by diving by the maximum absolute values of each features #max_abs_scaler=preprocessing.MaxAbsScaler() #X maxabs=max abs scaler.fit transform(X)

Task 1: Create the scores plot without any scaling

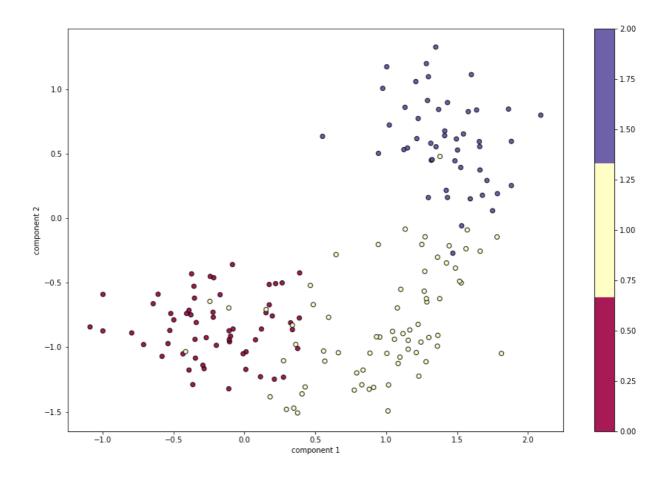


Task 2: Create the scores plot with standard scaling

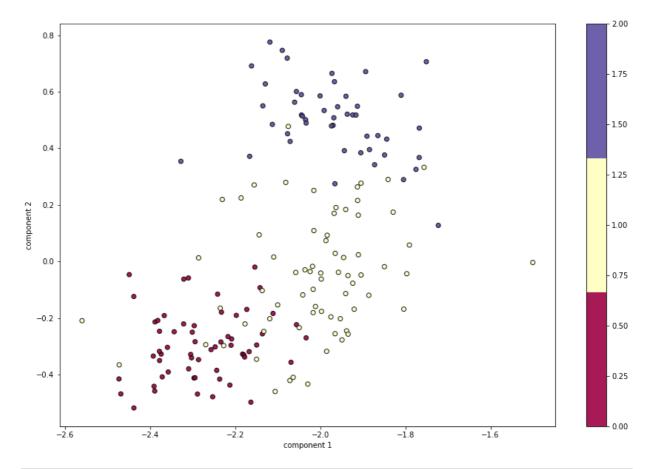


Task 3: Create the scores plot with min max scaling

1680.0



Task 4: Create the scores plot with max abs scaling



In []: