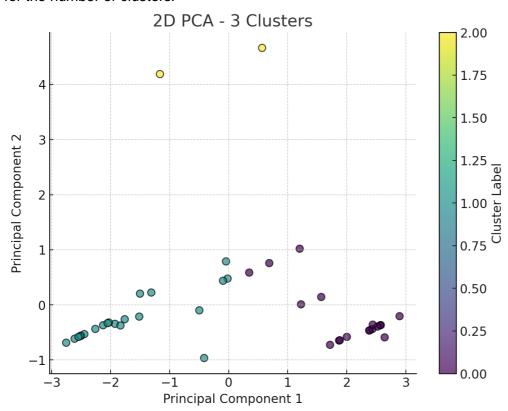
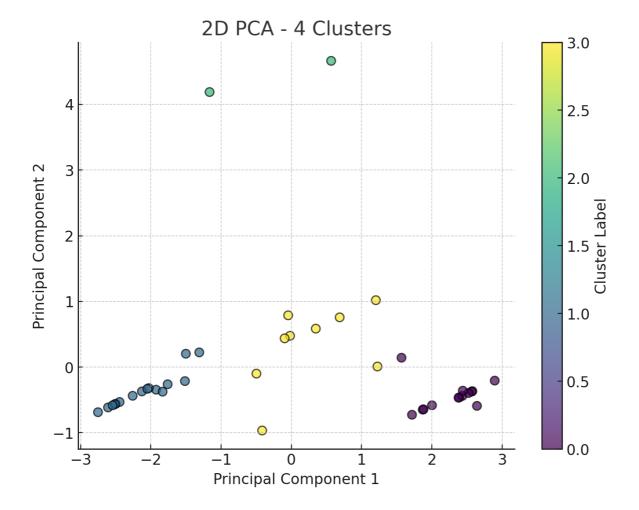


In this plot, it appears that the elbow point is not distinctly sharp, which is common. However, there seems to be a potential elbow point around 3 or 4 clusters, suggesting that either of these could be a reasonable choice for the number of clusters.

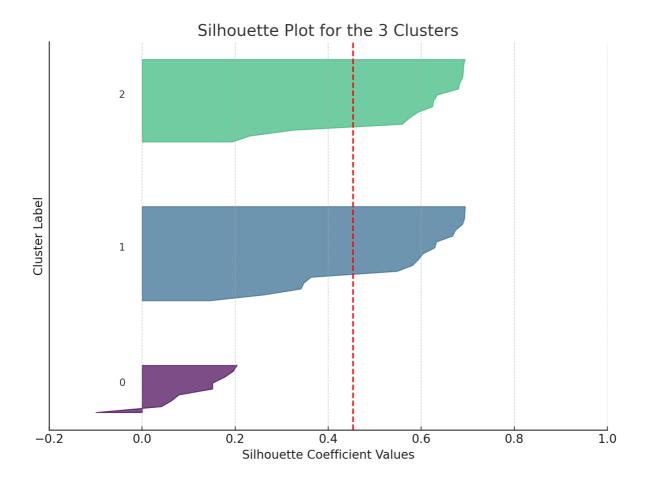




The Random Forest Classifier has ranked the features by their importance in determining the clusters, with the importance score ranging from 0 to 1 (1 being the most important). Here are the top features ranked by their importance:

- 1. **High STIR Signal of the Plantar Fascia (Y/N)**: Importance = 0.2026
- 2. Plantar Fascial Tear (Y/N) linear high signal/fiber interruption: Importance = 0.2003
- 3. Max Fascial Thickness-Coronal (mm): Importance = 0.1604
- 4. Intermediate STIR Signal of the Plantar Fascia (Y/N): Importance = 0.1446
- 5. Edema Deep to fascia (Y/N): Importance = 0.1439
- 6. Plantar Fat Pad Superficial Edema (0/1/2): Importance = 0.1128
- 7. Intermediate signal foci in the Plantar Fascia on T2 (Y/N): Importance = 0.0354
- 8. High signal foci in the Plantar Fascia on T2 (Y/N): Importance = 0.0000

It appears that there are only 7 unique features listed above due to the initial selection of features for this analysis, with the "High signal foci in the Plantar Fascia on T2 (Y/N)" feature showing no importance in this context.



The silhouette plot for the 3 clusters shows the silhouette coefficient values for each sample within the clusters, with the red dashed line indicating the average silhouette score of approximately 0.454. The silhouette coefficient measures how similar an object is to its own cluster (cohesion) compared to other clusters (separation). The range of the silhouette coefficient is between -1 and 1, where a high value indicates that the object is well-matched to its own cluster and poorly matched to neighboring clusters.

In this case, the average silhouette score suggests a reasonable structure has been found, although there's room for improvement. Generally, a score above 0.5 is considered good, indicating tight clusters. The spread of the silhouette scores within each cluster shows some variation, suggesting differences in cohesion and separation across the clusters.

1. Edema Deep to fascia (Y/N)

Chi2: 26.22

• p-value: 2.02e-06

2. Plantar Fat Pad Superficial Edema (0/1/2)

Chi2: 24.86

p-value: 5.36e-05

3. Plantar Fascial Tear (Y/N) linear high signal/fiber interruption

• Chi2: 32.92

• p-value: 7.10e-08

4. Intermediate signal foci in the Plantar Fascia on T2 (Y/N)

• Chi2: 7.48

p-value: 0.0238

5. High signal foci in the Plantar Fascia on T2 (Y/N)

Chi2: 0.00p-value: 1.00

6. Intermediate STIR Signal of the Plantar Fascia (Y/N)

Chi2: 30.86p-value: 1.99e-07

7. High STIR Signal of the Plantar Fascia (Y/N)

Chi2: 36.85p-value: 9.96e-09

The results indicate significant associations between the clusters and most of the categorical features, as evidenced by the low p-values (p < 0.05) for all but one feature. Specifically, features related to STIR signals ("Intermediate STIR Signal of the Plantar Fascia (Y/N)" and "High STIR Signal of the Plantar Fascia (Y/N)" and the "Plantar Fascial Tear (Y/N) linear high signal/fiber interruption" feature show particularly strong associations with the clusters, given their low p-values and high Chi2 values.

The "High signal foci in the Plantar Fascia on T2 (Y/N)" feature shows no association with the clusters, as indicated by a p-value of 1.00.

The Chi-Squared tests of independence between the cluster assignments and each significant pathological feature yield the following results:

1. Significant Myxoid Change

• Chi2: 6.26

• p-value: 0.044

• This result suggests a statistically significant association between cluster assignments and significant myxoid changes, as the p-value is below the typical alpha level of 0.05.

2. Significant Perifascial Changes

• Chi2: 0.59

p-value: 0.744

• There is no statistically significant association between cluster assignments and significant perifascial changes, indicated by a p-value well above 0.05.

3. Significant FCM Cells

• Chi2: 1.38

p-value: 0.501

• Similarly, there is no statistically significant association between cluster assignments and the presence of significant FCM cells.

4. Significant Loss of Collagen

Chi2: 1.57

p-value: 0.456

• There is no statistically significant association between cluster assignments and significant loss of collagen.

These results indicate that among the pathological features considered, only the presence of significant myxoid changes shows a significant association with the cluster assignments within the dataset. This suggests that the clustering may reflect differences in the underlying pathology, particularly regarding myxoid changes.

Feature	Significant_Myxoid_Change	Significant_Perifascial_Changes	Significant_FCM_Cells	Significant_Loss_of_Collagen
Significant_Myxoid_Change	1	0.095	0.11	0.156
Significant_Perifascial_Changes	0.095	1	0.139	0.02
Significant_FCM_Cells	0.11	0.139	1	-0.326
Significant_Loss_of_Collagen	0.156	0.02	-0.326	1

Key observations from the correlation matrix:

- Significant Myxoid Change has a slight positive correlation with Significant Loss of Collagen (0.156), suggesting a mild association where cases with significant myxoid changes might also present significant loss of collagen more frequently than by chance.
- Significant FCM Cells and Significant Loss of Collagen show a notable negative correlation (-0.326), indicating that cases with significant FCM cells are less likely to present significant loss of collagen, and vice versa. This suggests a distinct pathological process or stage of disease progression.
- Correlations between other pairs of significant pathological features are relatively low, indicating that these conditions may occur independently of each other within the dataset.
- Optimal groupings in the data suggest 3 or 4 distinct patterns based on imaging features related to
 plantar fasciitis, with specific features like STIR signals and fascial tears playing a key role in
 differentiating these groups.
- Significant myxoid changes were statistically linked to cluster assignments, highlighting a potential relationship between specific imaging features and pathological changes.
- A correlation analysis revealed a notable negative correlation between significant FCM cells and significant loss of collagen, suggesting distinct disease processes or stages might be captured by our clustering.