



# UNSUPERVISED MACHINE LEARNING FOR STRATIFYING PATIENT SUBGROUPS IN AUTOIMMUNE LIVER DISEASE

Abstract No.  
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Authors

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## Project Aims & Objectives

Autoimmune liver disease is a complex and heterogeneous condition that challenges effective diagnosis and management. Manual grouping of patients based on disease grade and clinical parameters is a tedious task and makes it challenging to derive meaningful insights from each group. This study aimed to use machine learning based clustering techniques to identify distinct and clinically relevant patient subgroups from the patient data, enhancing the understanding of disease heterogeneity and supporting more personalized care strategies

## Materials And Methods

The study analyzed a comprehensive dataset containing clinical, biochemical, immunological, and demographic variables from patients diagnosed with autoimmune liver disease. Eight unsupervised clustering algorithms were applied to the data. The elbow method was used to identify the optimal number of clusters, and clustering performance was evaluated using Silhouette Score, Davies-Bouldin Index, and Calinski-Harabasz Index. A novel Combined Performance Score was used to compare model performance. The features were assessed to determine the most significant ones driving cluster formation.

## Charts and Figures

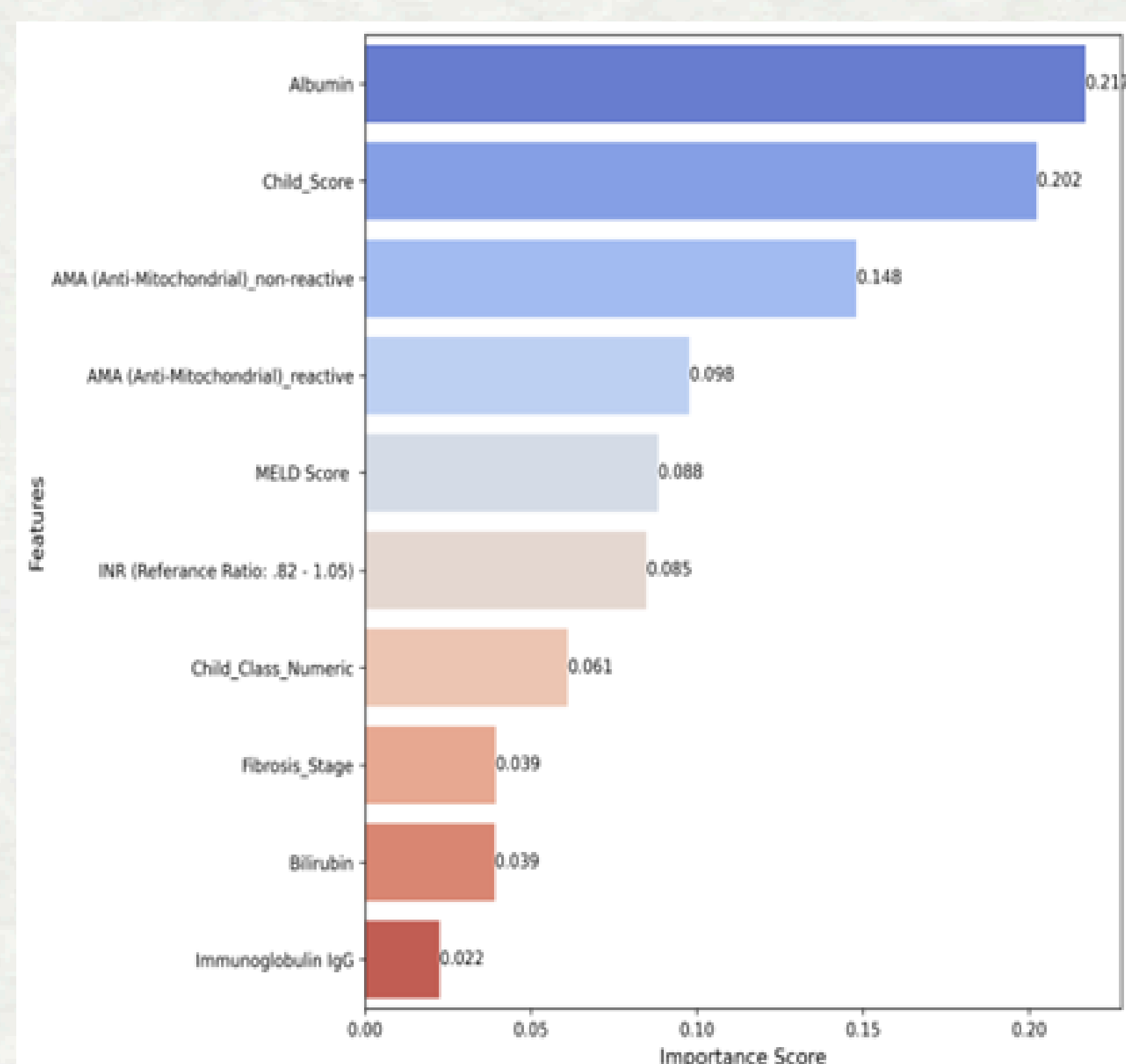


Fig. 1. Feature Importance Plot

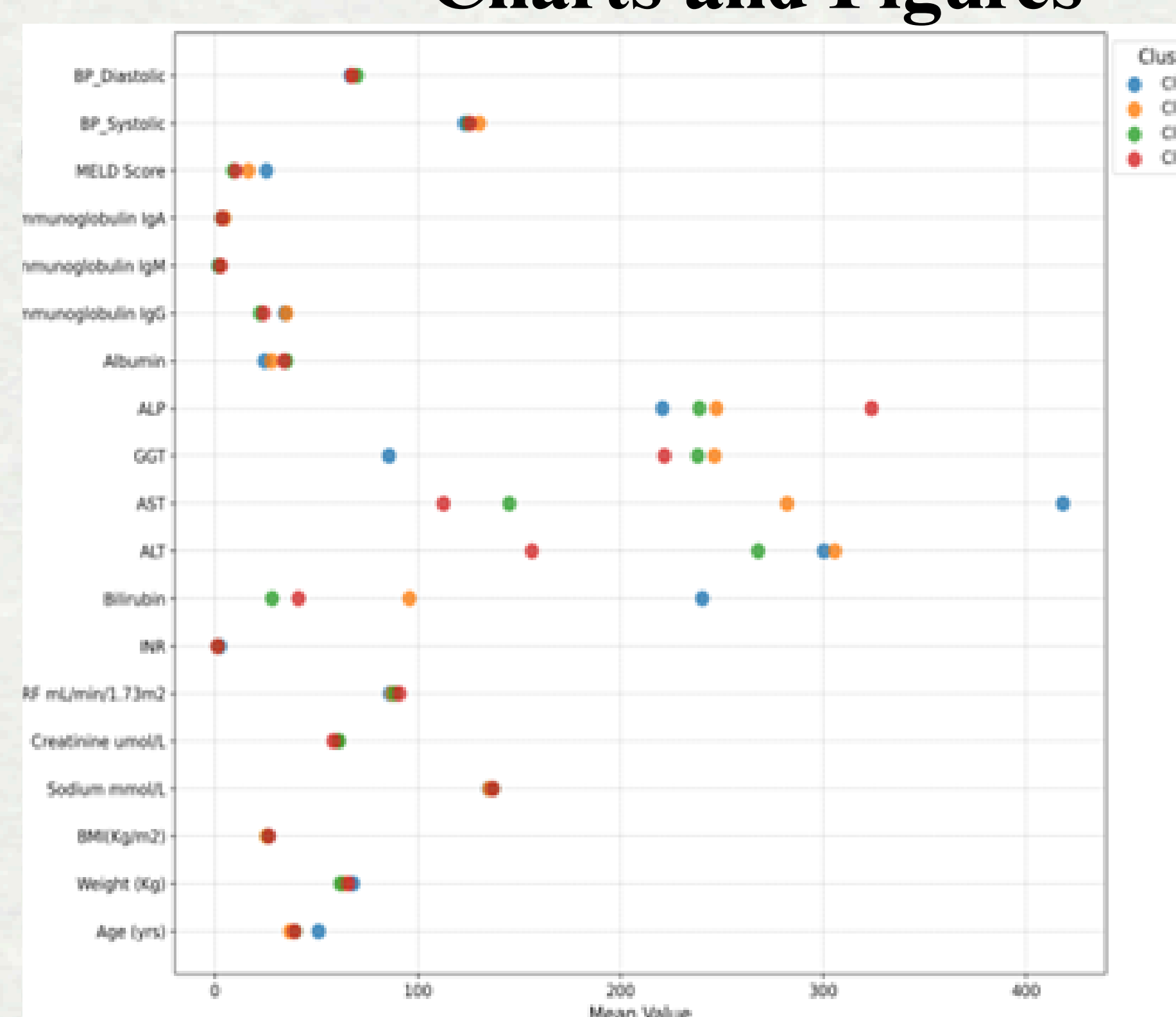


Fig. 2. Distribution of mean values of each cluster

Gender	Female	Female	Female	Female
Status	Alive	Alive	Alive	Alive
Alcoholic	no	no	no	no
Smoking	no	no	no	no
Diabetes Status	no	no	no	no
Hypertension	no	no	no	no
ANA (Antinuclear Antibody)	non-reactive	reactive	reactive	non-reactive
ASMA (Anti-Smooth Muscle Antibody)	non-reactive	non-reactive	non-reactive	non-reactive
Anti-LKM (Liver Kidney Microsomal Antibody)	non-reactive	non-reactive	non-reactive	non-reactive
AMA (Anti-Mitochondrial)	non-reactive	non-reactive	non-reactive	reactive
Hep A	non-reactive	non-reactive	non-reactive	non-reactive
Hep B	non-reactive	non-reactive	non-reactive	non-reactive
HCV	non-reactive	non-reactive	non-reactive	non-reactive
Fibrosis_Stage	F4	F2	F2	F2
Child_Pugh_Score	Class C, 11	Class B, 8	Class A, 6	Class A, 5
	Cluster 0	Cluster 1	Cluster 2	Cluster 3

Fig. 3. Categorical feature analysis of clusters

## Analysis and Result

KMeans achieved the highest combined performance score of 0.969 among the eight clustering algorithms. The elbow method identified four optimal clusters. Feature importance analysis found albumin, the Child–Pugh score, and key immune markers as crucial parameters relevant to cluster formation. Comprehensive cluster analyses revealed distinct patterns of liver disease progression and severity. The identified clusters revealed a spectrum of disease states, ranging from severe to mild, with intermediate groups showing active immune involvement. These subgroup patterns aligned with varying stages of disease progression and severity

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## Conclusion

This study demonstrates the use of machine learning based clustering in identifying distinct patient subgroups in autoimmune liver disease. This clustering framework underscores the potential of clustering techniques to cluster patients to enhance the understanding of disease heterogeneity, improving patient outcomes and providing personalized treatments. This work contributes to biomedical research by effectively utilizing machine learning based clustering to uncover autoimmune liver disease patterns and support more accurate predictions.

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