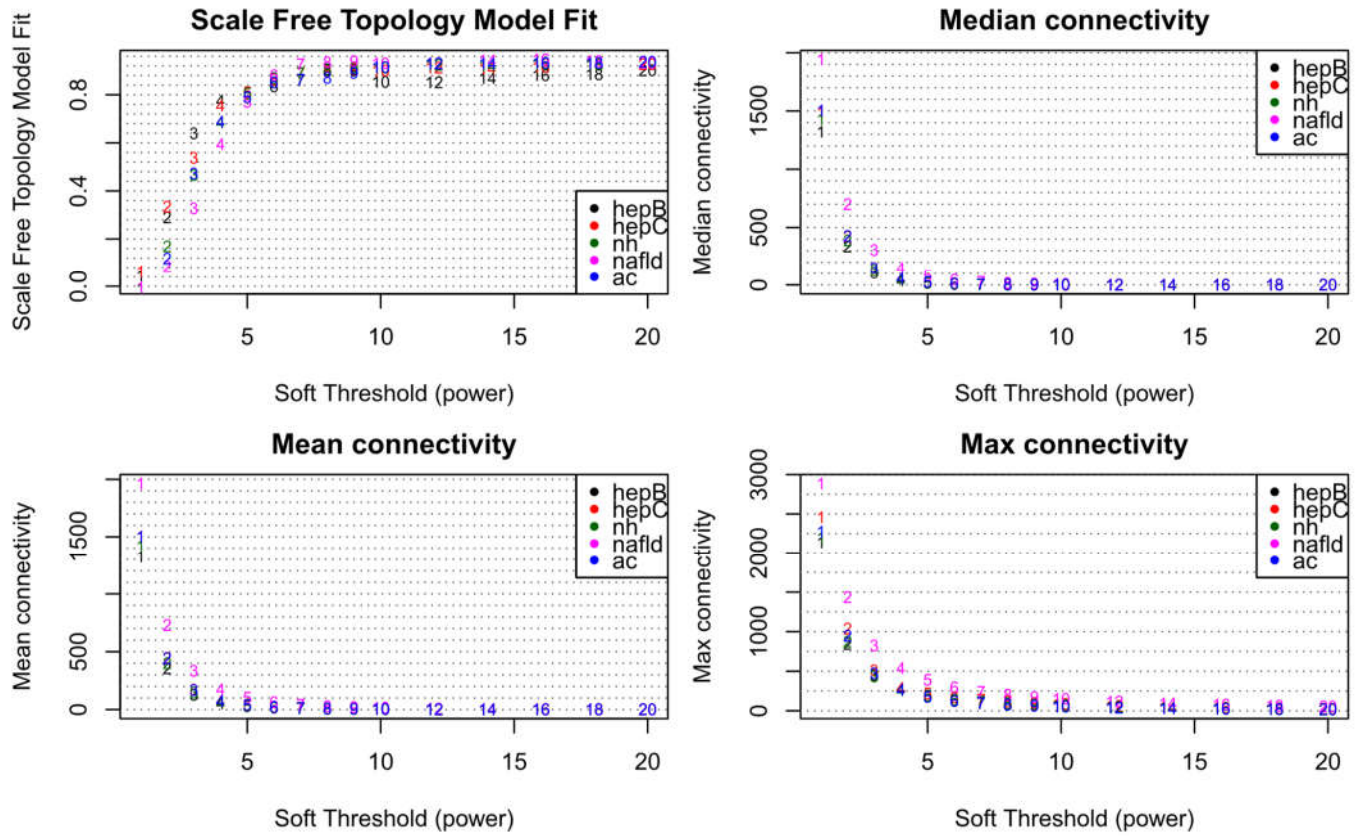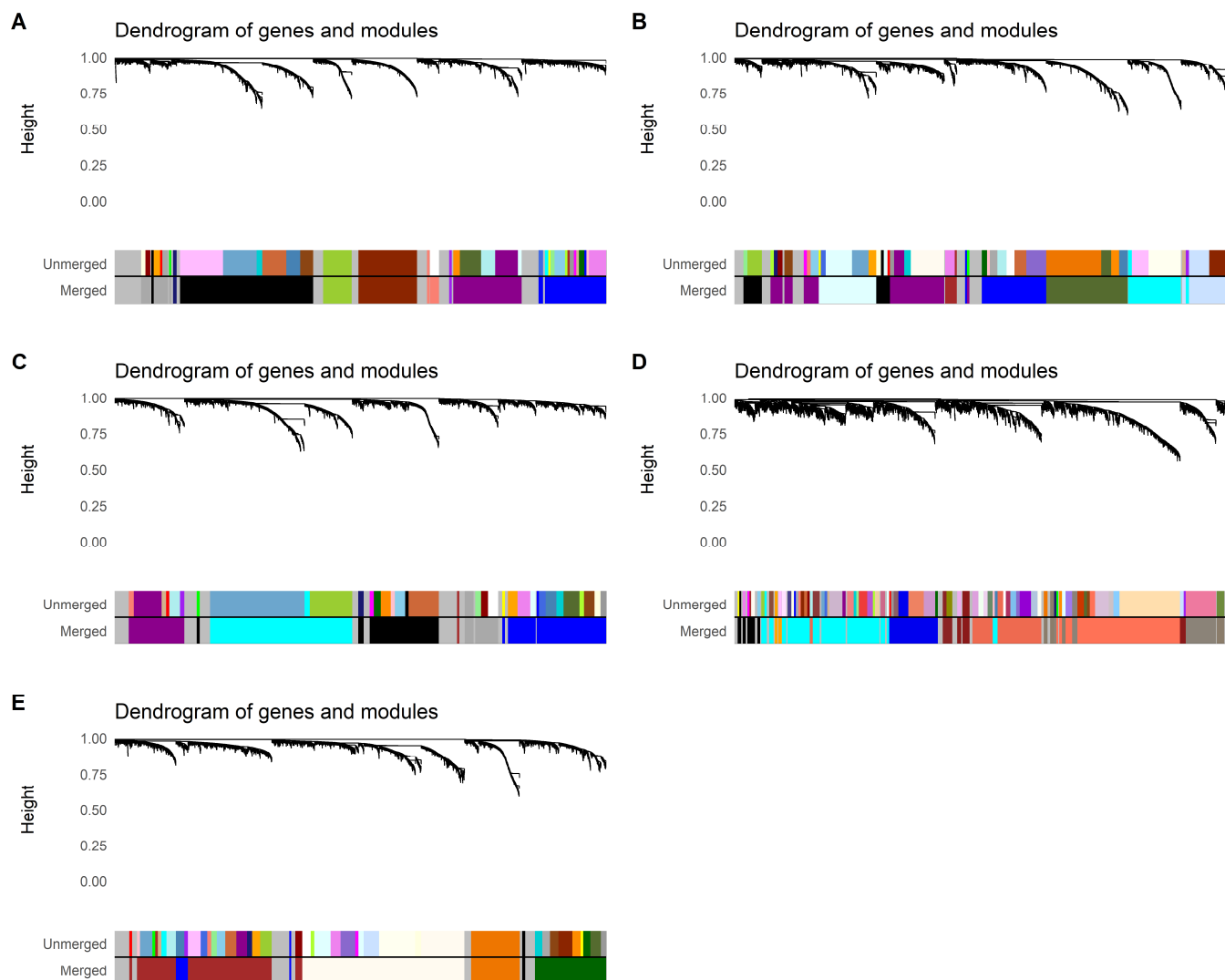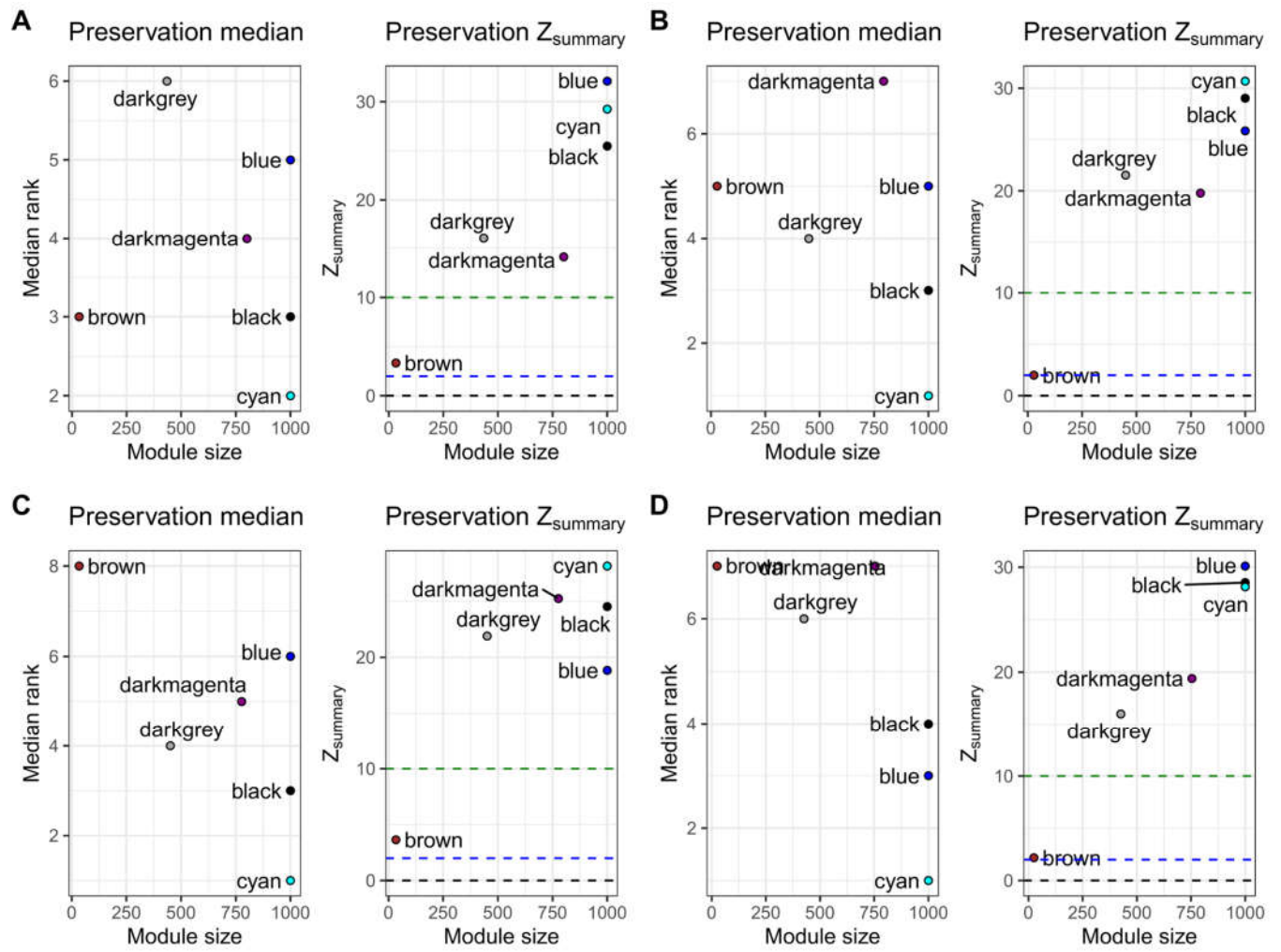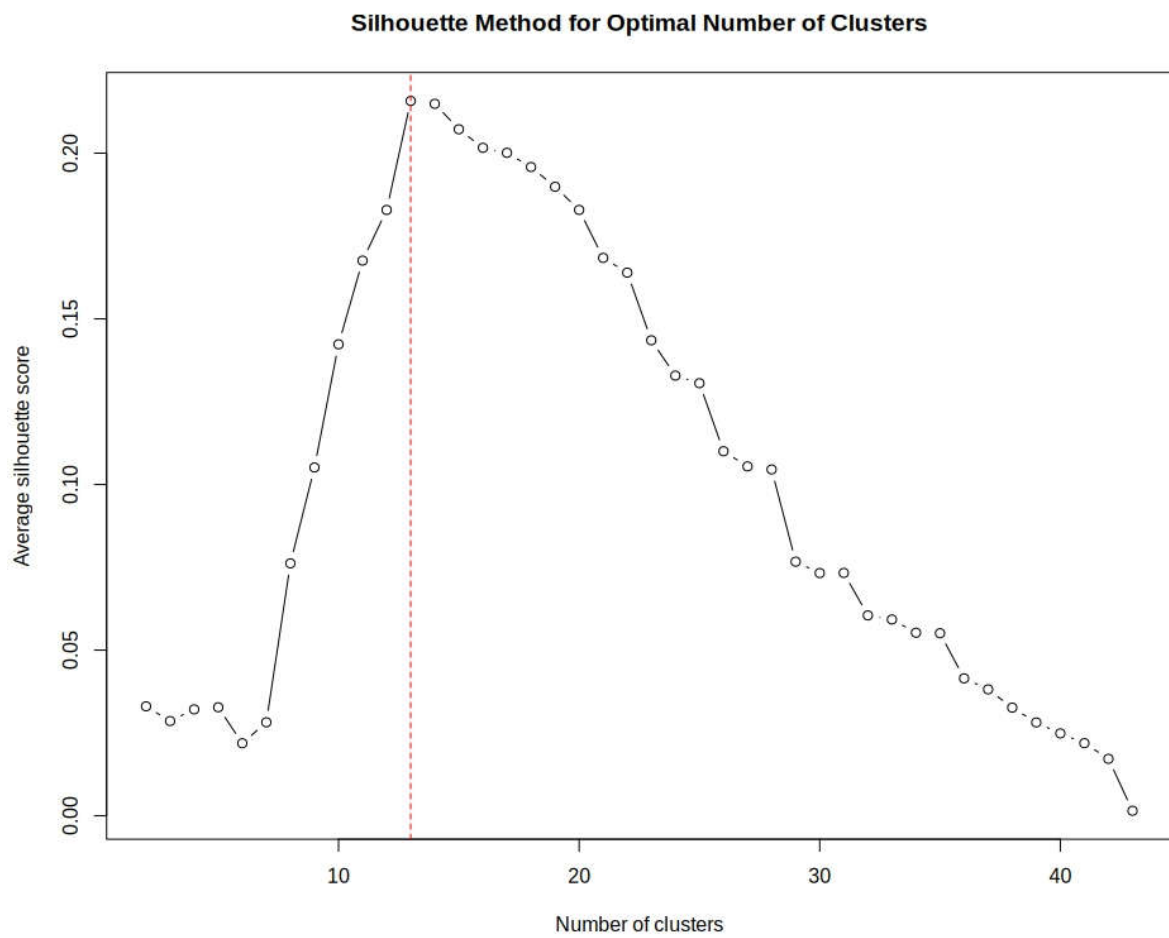# Supplementary Figures

Figure S1. Scale independence and mean connectivity results for the gene expression of each examined etiology of HCC.

Figure S2. Network Construction Dendrogram Results per Etiology with **(A)** as Alcohol Consumption **(B)** Hepatitis B, **(C)** Hepatitis C, **(D)** NAFLD, **(E)** No History.

Figure S3. Module preservation results with No History dataset as reference and etiologies **(A)** Alcohol Consumption **(B)** Hepatitis B, **(C)** Hepatitis C, **(D)** NAFLD as the test set.

Figure S4. Silhouette method results to determine optimal number of clusters for hierarchical clustering.