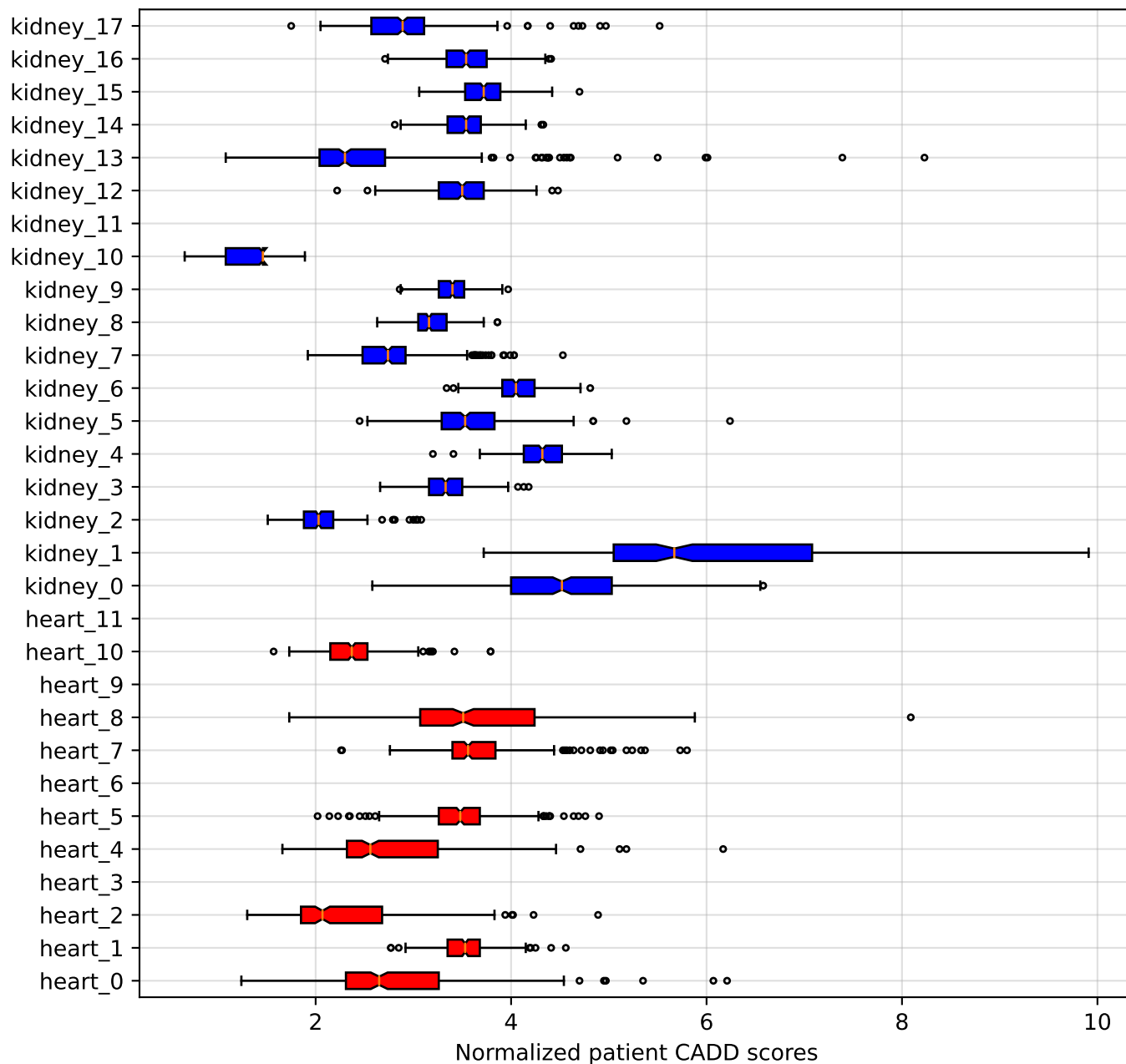
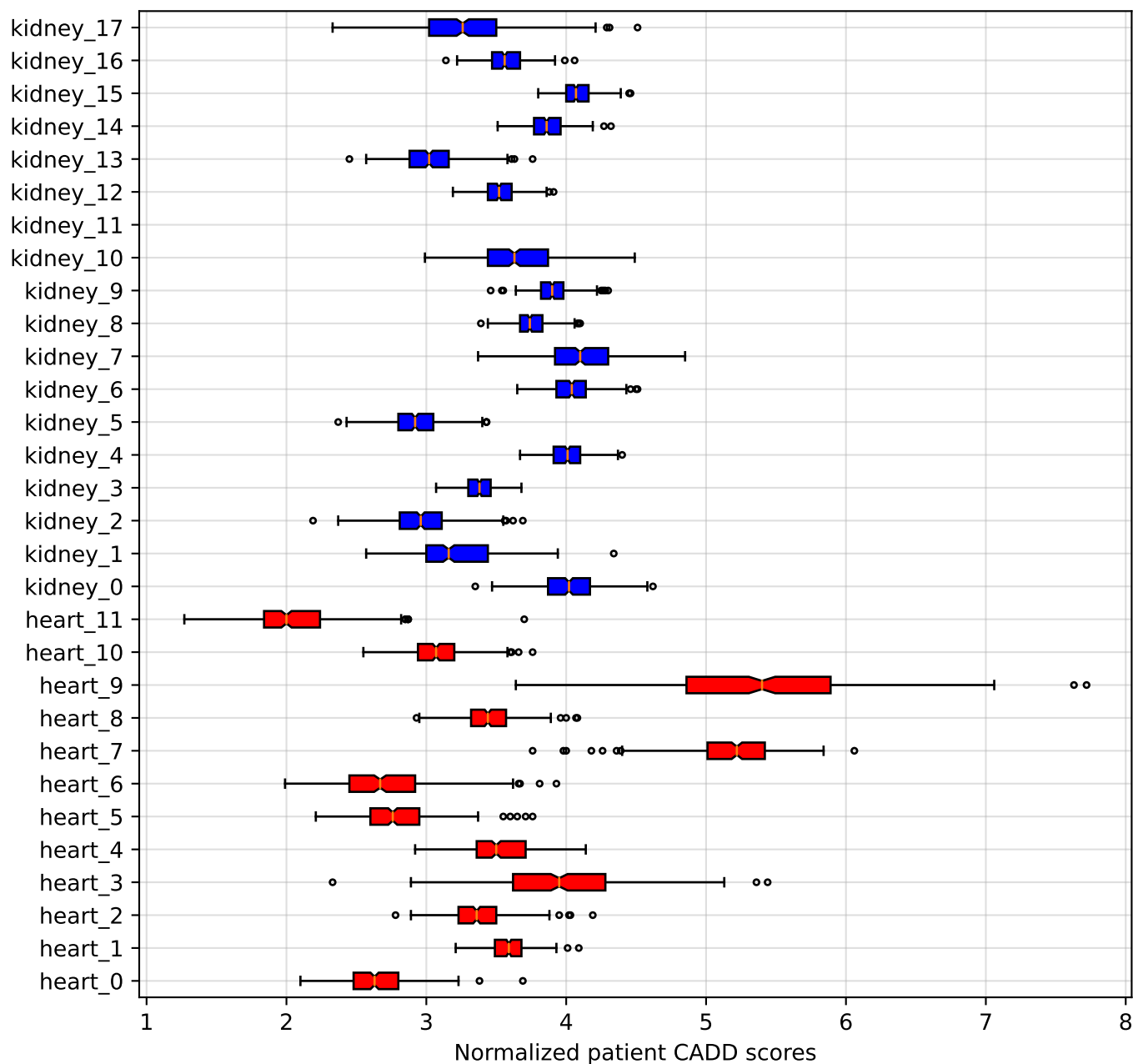


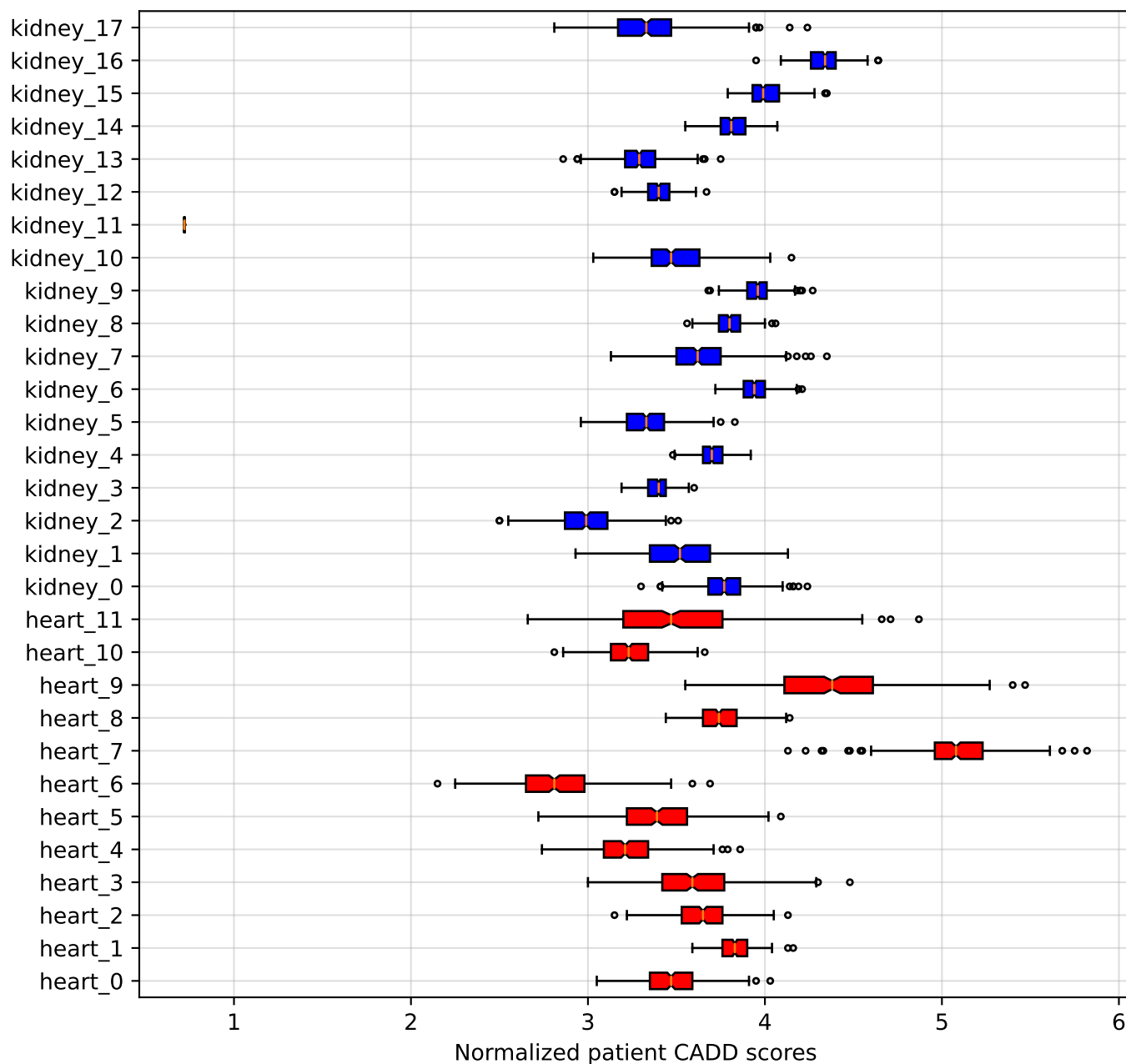
Top 1.0% smallest p-values of marker genes from each cluster  
were used to filter KUTD cohort



Top 5.0% smallest p-values of marker genes from each cluster  
were used to filter KUTD cohort



Top 10.0% smallest p-values of marker genes from each cluster  
were used to filter KUTD cohort



Top 25.0% smallest p-values of marker genes from each cluster  
were used to filter KUTD cohort

