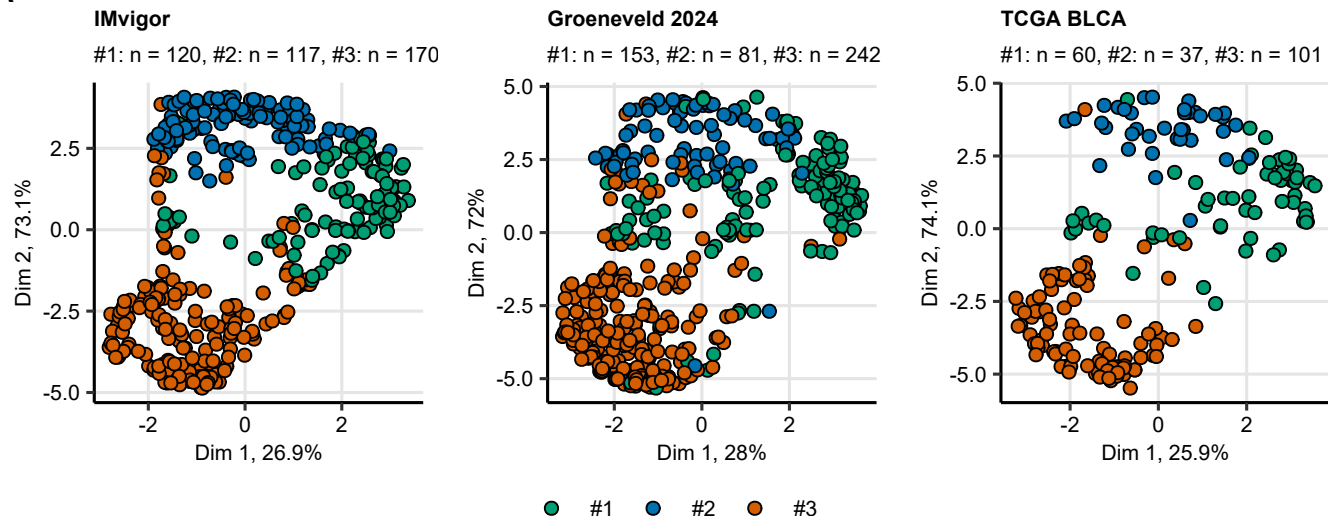
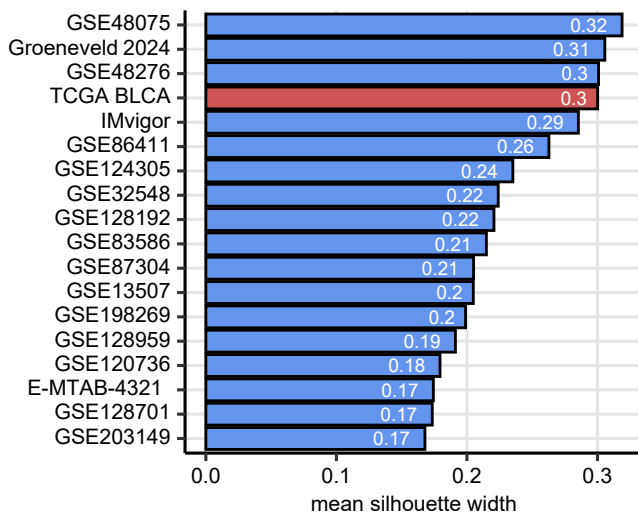
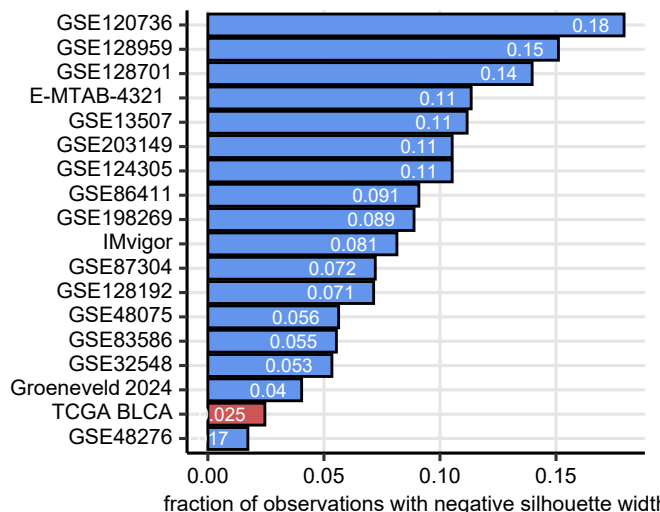
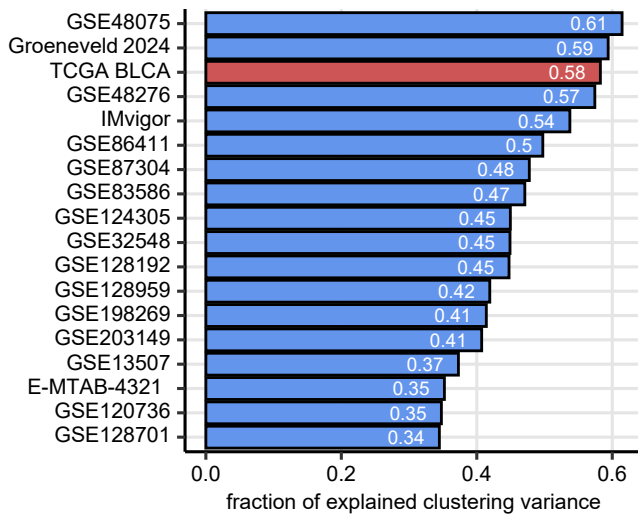


A**B****Cluster separation, transcriptome****Misclassification rate, transcriptome****Explained variance, transcriptome****Neighborhood preservation, transcriptome**