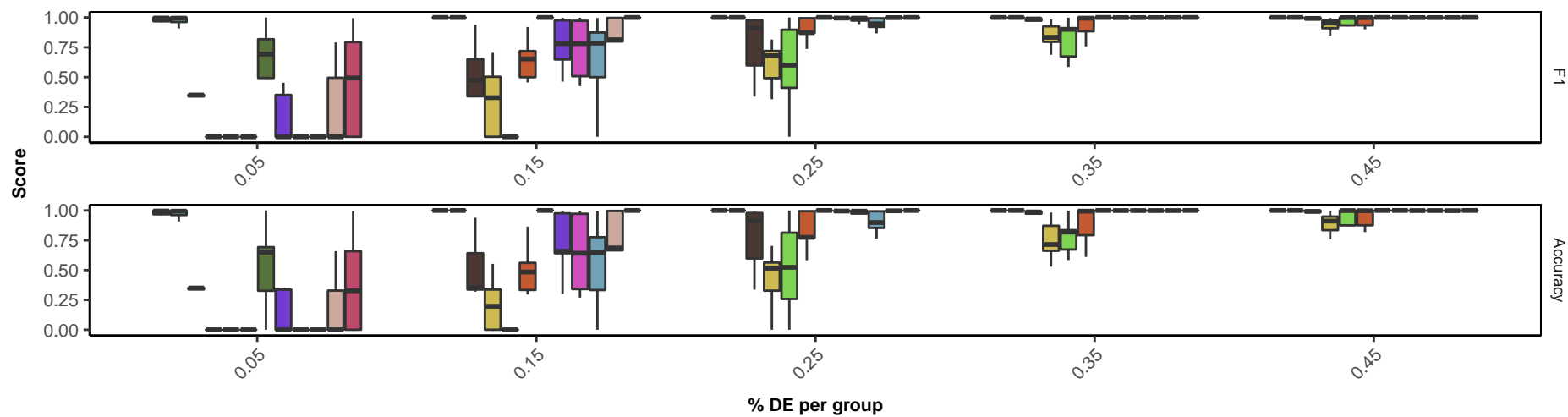
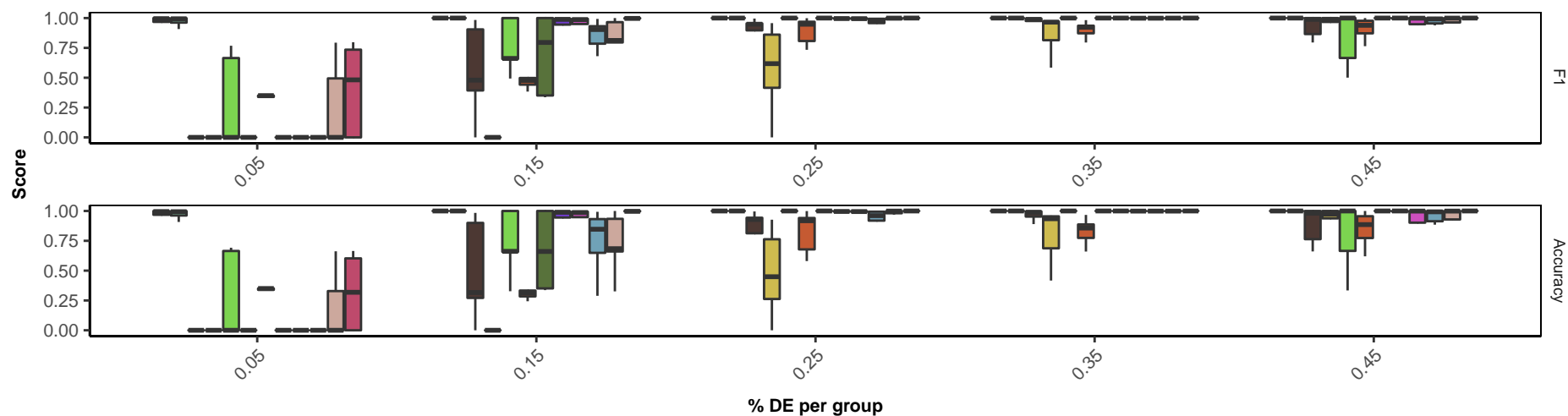


a**b**

Method

cellassign, densitycut, kmeans, SC3, seurat_0.4, seurat_1.2, cellassign_shrinkage, dynamicTreeCut, phenograph, scmap, seurat_0.8, Zheng_cor

c