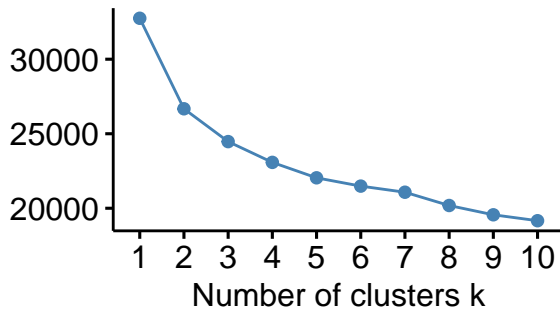Optimal number of clusters for kmeans (75% of genes)