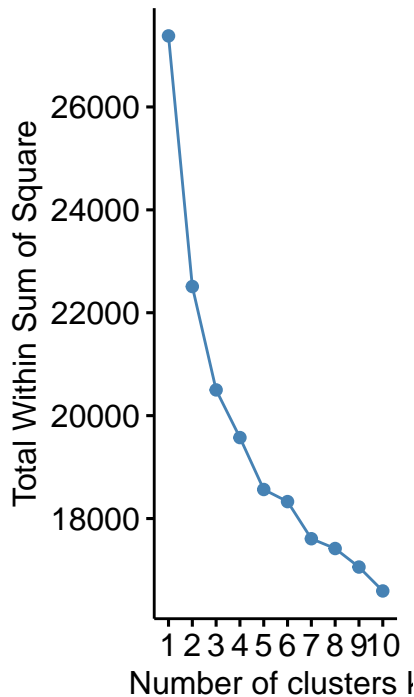
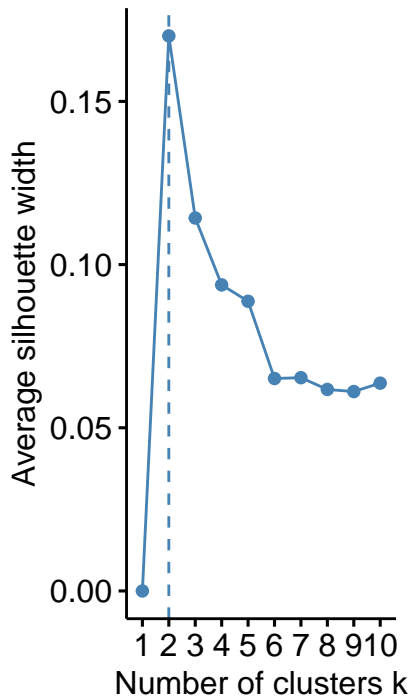


Optimal number of clusters
for K-medoids (50% of genes)

Wss



Silhouette



Gap

