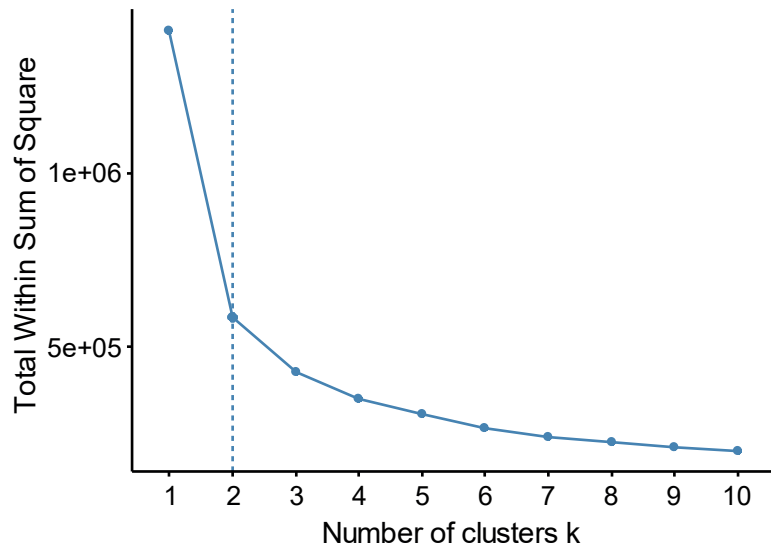


a) **Elbow method** (Optimal number of clusters for ChIPseq heatmap shown in Figure 2d)



b) **Silhouette method** (Optimal number of clusters for ChIPseq heatmap shown in Figure 2d)

