**Practical No.7: Implementing Clustering with R(k-means clustering)**

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| newiris <- iris  newiris$Species <- NULL  (kc <- kmeans(newiris,3))  print(kc)  # Compare the Species label with the clustering result.  table (iris$Species,kc$cluster)  plot(newiris[c("Sepal.Length","Sepal.Width")],col=kc$cluster)  points(kc$centers[,c("Sepal.Length","Sepal.Width")],col=1:3,pch=8,cex=2) |

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