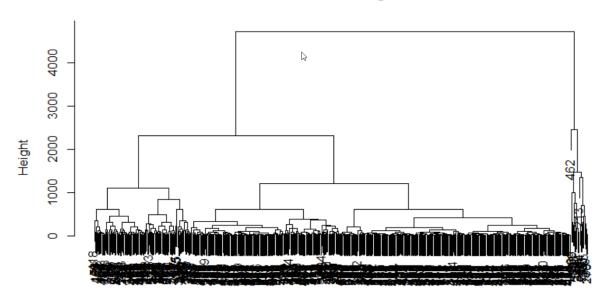


Cluster Dendrogram



cancer_dist hclust (*, "complete")

Assignment 9

```
> View(cancer)
> summary(cancer)
id
Min. : 8670
1st Qu.: 869218
Median : 906024
Mean : 30371831
3rd Qu.: 8813129
Max. : 911320502
                  diagnosis radius_mean
B:357 Min. : 6.981
M:212 Ist Qu.:11.700
Median :13.370
Mean :14.127
3rd Qu.:15.780
                                          texture_mean
Min. : 9.71
1st Qu.:16.17
Median :18.84
Mean :19.29
3rd Qu.:21.80
                                                        perimeter_mean
Min. : 43.79
1st Qu.: 75.17
Median : 86.24
Mean : 91.97
3rd Qu.:104.10
Max. :188.50
                                                                        area_mean
Min. : 143.5
1st Qu.: 420.3
Median : 551.1
Mean : 654.9
3rd Qu.: 782.7
                                                                                       smoothness_mean
Min. :0.05263
1st Qu.:0.08637
Median :0.09587
Mean :0.09636
3rd Qu.:0.10530
              Max. :911.320502 (compactness.mean Min. :0.01938 1st Qu.:0.06492 Median: 0.09263 Median: 0.09263 Median: 0.09263 Median: 0.0434 3rd Qu.:0.13040 Max. :0.34540 texture_se Min. :0.3602 !st Qu.:0.8339 "Ard Qu.:1.4740 Max. :4.8850 (concave.points_se concave.points_se
        :911320502
                                                                               2501.0
                                                                                              :0.16340
                                                                  fractal_dimension_mean
                                                                                 3rd qu.:0.04205
Max. :0.39600
perimeter_wors'
Min. : 50.41
ist qu.:84.11
Median : 97.66
3rd qu.:125.40
Max. :251.20
t symmetry_worst
Min. :0.1565
ist qu.:0.2504
Median :0.2822
Mean :0.2901
3rd qu.:0.3179
Max. :0.6638
3rd Qu.:1084.0 3rd Qu.
Max. :4254.0 Max.
fractal_dimension_worst
Min. :0.05504
1st Qu.:0.07146
Median :0.08004
 Mean :0.08395
3rd Qu.:0.09208
Max. :0.20750
> table(cancer$diagnosis)
B M

357 212

>#TO factor the data set

> cancer<-na.omit(cancer)

> cancer<-cancer[-1]

> kmeans_2<- kmeans(cancer[,-1],2,nstart = 10)

> kmeans_25cluster
           4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31
Console Terminal × Jobs ×
_/
Mean :0.08395
3rd Qu.:0.09208
Max. :0.20750
> table(cancer$diagnosis)
```