

Miniproject08_Attempt2.R

Noah

2022-03-05

```
wisc.df <- read.csv("WisconsinCancer.csv", row.names=1)
wisc.data <- wisc.df[,-1]
diagnosis <- factor(wisc.data[,1])
colMeans(wisc.data[,,-1])
```

##	texture_mean	perimeter_mean	area_mean
##	1.928965e+01	9.196903e+01	6.548891e+02
##	smoothness_mean	compactness_mean	concavity_mean
##	9.636028e-02	1.043410e-01	8.879932e-02
##	concave.points_mean	symmetry_mean	fractal_dimension_mean
##	4.891915e-02	1.811619e-01	6.279761e-02
##	radius_se	texture_se	perimeter_se
##	4.051721e-01	1.216853e+00	2.866059e+00
##	area_se	smoothness_se	compactness_se
##	4.033708e+01	7.040979e-03	2.547814e-02
##	concavity_se	concave.points_se	symmetry_se
##	3.189372e-02	1.179614e-02	2.054230e-02
##	fractal_dimension_se	radius_worst	texture_worst
##	3.794904e-03	1.626919e+01	2.567722e+01
##	perimeter_worst	area_worst	smoothness_worst
##	1.072612e+02	8.805831e+02	1.323686e-01
##	compactness_worst	concavity_worst	concave.points_worst
##	2.542650e-01	2.721885e-01	1.146062e-01
##	symmetry_worst	fractal_dimension_worst	
##	2.900756e-01	8.394582e-02	

```
apply(wisc.data,2,sd)
```

##	radius_mean	texture_mean	perimeter_mean
##	3.524049e+00	4.301036e+00	2.429898e+01
##	area_mean	smoothness_mean	compactness_mean
##	3.519141e+02	1.406413e-02	5.281276e-02
##	concavity_mean	concave.points_mean	symmetry_mean
##	7.971981e-02	3.880284e-02	2.741428e-02
##	fractal_dimension_mean	radius_se	texture_se
##	7.060363e-03	2.773127e-01	5.516484e-01
##	perimeter_se	area_se	smoothness_se
##	2.021855e+00	4.549101e+01	3.002518e-03
##	compactness_se	concavity_se	concave.points_se
##	1.790818e-02	3.018606e-02	6.170285e-03
##	symmetry_se	fractal_dimension_se	radius_worst

```
##          8.266372e-03          2.646071e-03          4.833242e+00
##          texture_worst          perimeter_worst          area_worst
##          6.146258e+00          3.360254e+01          5.693570e+02
##          smoothness_worst          compactness_worst          concavity_worst
##          2.283243e-02          1.573365e-01          2.086243e-01
##          concave.points_worst          symmetry_worst          fractal_dimension_worst
##          6.573234e-02          6.186747e-02          1.806127e-02
```

```
wisc.pr <- prcomp(wisc.data[,-1])
summary(wisc.pr)
```

```
## Importance of components:
##          PC1          PC2          PC3          PC4          PC5          PC6          PC7
## Standard deviation    666.161  85.49544  26.52784  7.38816  6.31187  1.71854  1.347
## Proportion of Variance  0.982  0.01618  0.00156  0.00012  0.00009  0.00001  0.000
## Cumulative Proportion  0.982  0.99822  0.99978  0.99990  0.99999  0.99999  1.000
##          PC8          PC9          PC10          PC11          PC12          PC13          PC14          PC15
## Standard deviation    0.6079  0.3713  0.2867  0.1618  0.0597  0.0471  0.039  0.02855
## Proportion of Variance 0.0000  0.0000  0.0000  0.0000  0.0000  0.0000  0.000  0.00000
## Cumulative Proportion 1.0000  1.0000  1.0000  1.0000  1.0000  1.0000  1.000  1.00000
##          PC16          PC17          PC18          PC19          PC20          PC21
## Standard deviation    0.01981  0.01722  0.01381  0.01344  0.009507  0.007602
## Proportion of Variance 0.00000  0.00000  0.00000  0.00000  0.000000  0.000000
## Cumulative Proportion 1.00000  1.00000  1.00000  1.00000  1.000000  1.000000
##          PC22          PC23          PC24          PC25          PC26          PC27
## Standard deviation    0.005918  0.005557  0.004023  0.003535  0.001925  0.001698
## Proportion of Variance 0.000000  0.000000  0.000000  0.000000  0.000000  0.000000
## Cumulative Proportion 1.000000  1.000000  1.000000  1.000000  1.000000  1.000000
##          PC28          PC29
## Standard deviation    0.001416  0.0008387
## Proportion of Variance 0.000000  0.0000000
## Cumulative Proportion 1.000000  1.0000000
```

```
biplot(wisc.pr)
```

```
## Warning in arrows(0, 0, y[, 1L] * 0.8, y[, 2L] * 0.8, col = col[2L], length =
## arrow.len): zero-length arrow is of indeterminate angle and so skipped
```

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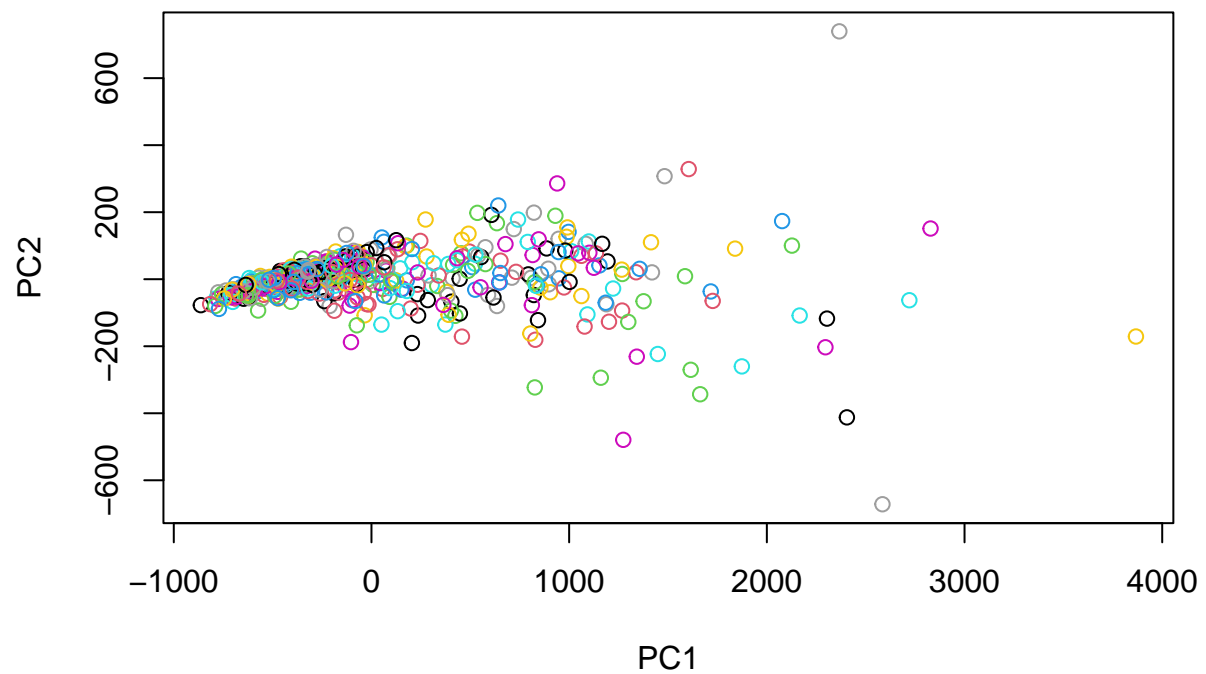
## Warning in arrows(0, 0, y[, 1L] * 0.8, y[, 2L] * 0.8, col = col[2L], length =
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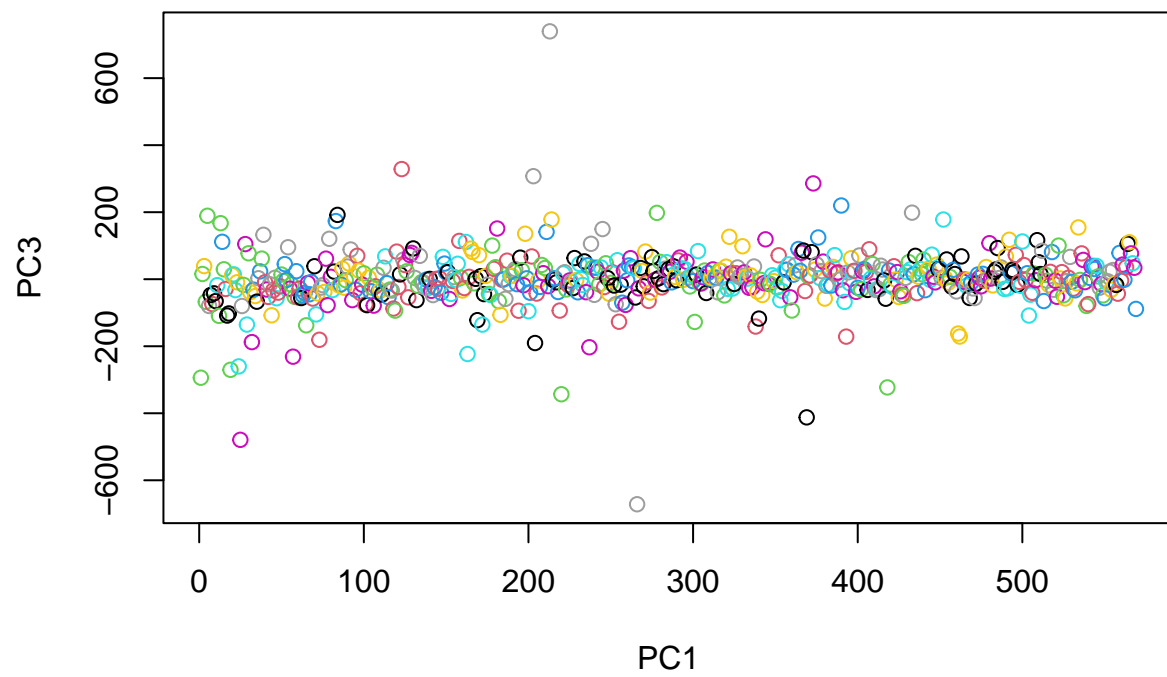
```

```
plot(wisc.pr$x[, 2 ], col = diagnosis,  
      xlab = "PC1", ylab = "PC3")
```

```
df <- as.data.frame(wisc.pr$x)  
df$diagnosis <- diagnosis
```

```
library(ggplot2)
```



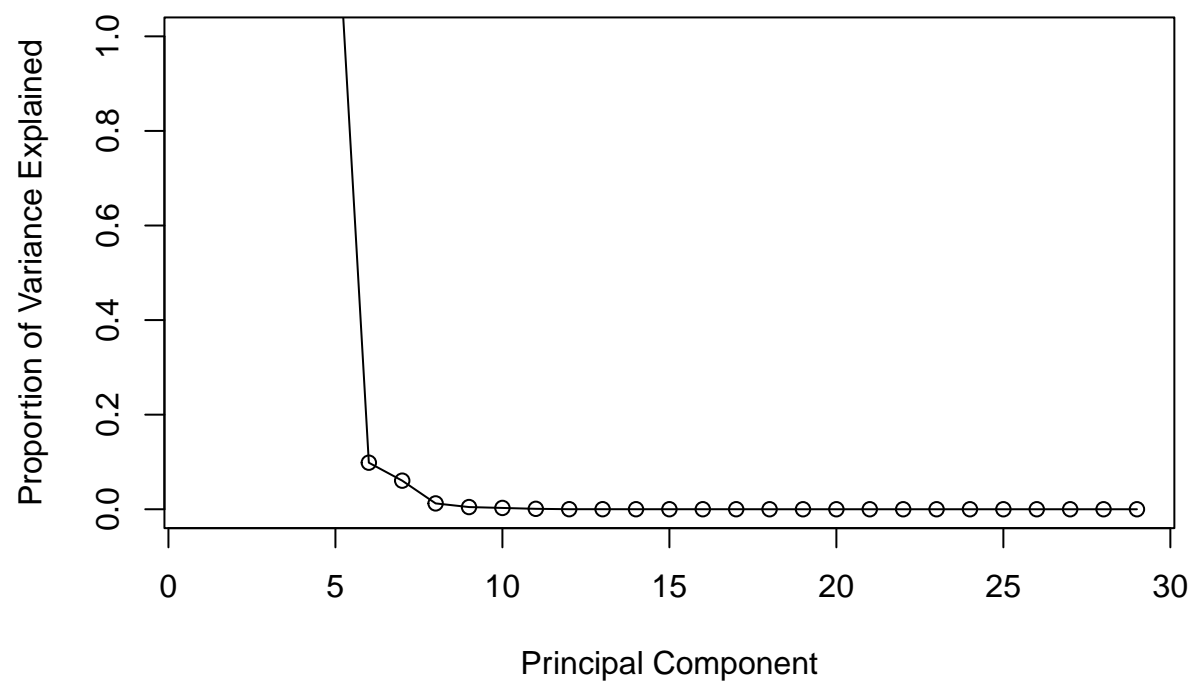
```
ggplot(df) +  
  aes(PC1, PC2, col= diagnosis) +  
  geom_point()
```

11.75	●	12.16	●	12.47	●	12.86	●	13.17	●	13.53	●	13.82	●	14.27	●	14.69	●	15.06
11.76	●	12.18	●	12.49	●	12.87	●	13.2	●	13.54	●	13.85	●	14.29	●	14.71	●	15.07
11.8	●	12.19	●	12.54	●	12.88	●	13.21	●	13.56	●	13.86	●	14.34	●	14.74	●	15.1
11.81	●	12.2	●	12.56	●	12.89	●	13.24	●	13.59	●	13.87	●	14.4	●	14.76	●	15.11
11.84	●	12.21	●	12.58	●	12.9	●	13.27	●	13.61	●	13.88	●	14.41	●	14.78	●	15.12
11.85	●	12.22	●	12.62	●	12.91	●	13.28	●	13.62	●	13.9	●	14.42	●	14.8	●	15.13
11.87	●	12.23	●	12.63	●	12.94	●	13.3	●	13.64	●	13.94	●	14.44	●	14.81	●	15.14
11.89	●	12.25	●	12.65	●	12.95	●	13.34	●	13.65	●	13.96	●	14.45	●	14.86	●	15.15
11.9	●	12.27	●	12.67	●	12.96	●	13.37	●	13.66	●	13.98	●	14.47	●	14.87	●	15.16
11.93	●	12.3	●	12.68	●	12.98	●	13.38	●	13.68	●	14.02	●	14.48	●	14.9	●	15.17
11.94	●	12.31	●	12.7	●	12.99	●	13.4	●	13.69	●	14.03	●	14.5	●	14.92	●	15.18
11.95	●	12.32	●	12.72	●	13	●	13.43	●	13.7	●	14.04	●	14.53	●	14.95	●	15.19
11.99	●	12.34	●	12.75	●	13.01	●	13.44	●	13.71	●	14.05	●	14.54	●	14.96	●	15.2
12	●	12.36	●	12.76	●	13.03	●	13.45	●	13.73	●	14.06	●	14.58	●	14.97	●	15.21
12.03	●	12.39	●	12.77	●	13.05	●	13.46	●	13.74	●	14.11	●	14.59	●	14.99	●	15.22
12.04	●	12.4	●	12.78	●	13.08	●	13.47	●	13.75	●	14.19	●	14.6	●	15	●	15.23
12.05	●	12.42	●	12.8	●	13.11	●	13.48	●	13.77	●	14.2	●	14.61	●	15.04	●	15.24
12.06	●	12.43	●	12.81	●	13.14	●	13.49	●	13.78	●	14.22	●	14.62	●	15.05	●	15.25
12.07	●	12.45	●	12.83	●	13.15	●	13.5	●	13.8	●	14.25	●	14.64	●	15.06	●	15.26

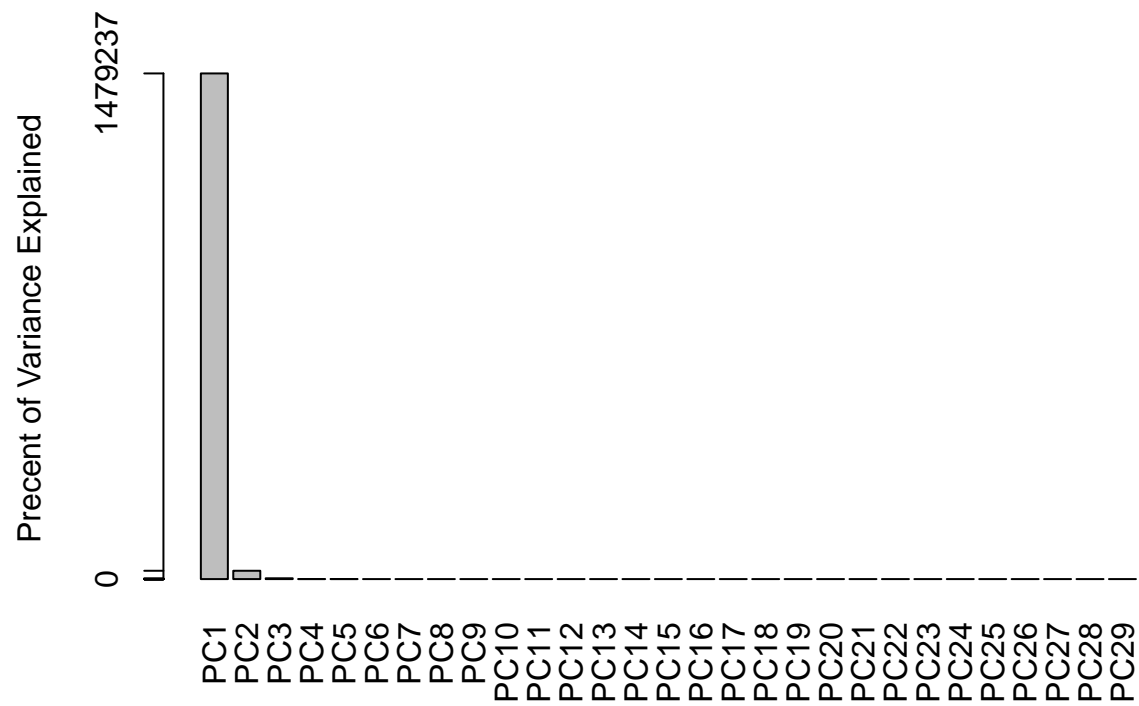
```
pr.var <- wisc.pr$sdev^2
head(pr.var)
```

```
## [1] 4.437711e+05 7.309470e+03 7.037265e+02 5.458498e+01 3.983975e+01
## [6] 2.953395e+00
```

```
pve <- pr.var / 30
plot(pve, xlab = "Principal Component",
     ylab = "Proportion of Variance Explained",
     ylim = c(0, 1), type = "o")
```

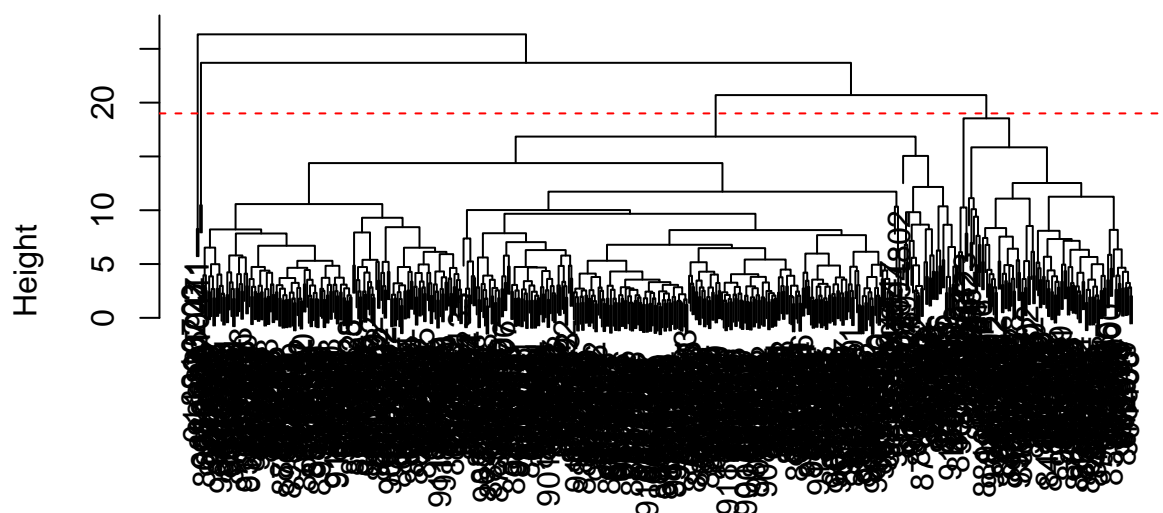


```
barplot(pve, ylab = "Precent of Variance Explained",
        names.arg=paste0("PC",1:length(pve)), las=2, axes = FALSE)
axis(2, at=pve, labels=round(pve,2)*100 )
```

```
data.scaled <- scale(wisc.data[, -1])
data.dist <- dist(data.scaled)
wisc.hclust <- hclust(data.dist, method = "complete")
plot(wisc.hclust)
abline(h = 19, col = "red", lty = 2)
```

Cluster Dendrogram



data.dist
hclust (*, "complete")

```
wisc.hclust.clusters <- cutree(wisc.hclust, k = 4)
table(wisc.hclust.clusters, diagnosis)
```

```
##              diagnosis
## wisc.hclust.clusters 6.981 7.691 7.729 7.76 8.196 8.219 8.571 8.597 8.598 8.618
##              1      0      0      0      0      0      0      0      0      0      0
##              2      1      1      1      1      1      1      1      1      1      1
##              3      0      0      0      0      0      0      0      0      0      0
##              4      0      0      0      0      0      0      0      0      0      0
##              diagnosis
## wisc.hclust.clusters 8.671 8.726 8.734 8.878 8.888 8.95 9 9.029 9.042 9.173
##              1      0      0      0      0      0      0 0      0      0      0
##              2      1      1      1      1      1      1 1      0      1      1
##              3      0      0      0      0      0      0 0      1      0      0
##              4      0      0      0      0      0      0 0      0      0      0
##              diagnosis
## wisc.hclust.clusters 9.268 9.295 9.333 9.397 9.405 9.423 9.436 9.465 9.504
##              1      0      0      0      0      0      0      0      0      0
##              2      1      1      1      1      1      1      1      1      1
##              3      0      0      0      0      0      0      0      0      0
##              4      0      0      0      0      0      0      0      0      0
##              diagnosis
## wisc.hclust.clusters 9.567 9.606 9.667 9.668 9.676 9.683 9.72 9.731 9.738 9.742
##              1      0      0      0      0      0      0      0      0      0
##              2      1      1      1      1      1      1      1      0      1      2
```

```

##          3      0      0      0      0      0      0      0      1      0      0
##          4      0      0      0      0      0      0      0      0      0      0
##          diagnosis
## wisc.hclust.clusters 9.755 9.777 9.787 9.847 9.876 9.904 10.03 10.05 10.08
##          1      0      0      0      0      0      0      0      0      0
##          2      1      1      1      1      2      1      1      1      1
##          3      0      0      0      0      0      0      0      0      0
##          4      0      0      0      0      0      0      0      0      0
##          diagnosis
## wisc.hclust.clusters 10.16 10.17 10.18 10.2 10.25 10.26 10.29 10.32 10.44 10.48
##          1      0      0      0      0      0      0      0      0      0
##          2      1      1      1      1      1      3      1      1      1      2
##          3      0      0      0      0      0      0      0      0      0
##          4      0      0      0      0      0      0      0      0      0
##          diagnosis
## wisc.hclust.clusters 10.49 10.51 10.57 10.6 10.65 10.66 10.71 10.75 10.8 10.82
##          1      0      0      0      0      0      0      0      0      0
##          2      2      2      2      1      1      1      1      1      2      1
##          3      0      0      0      0      0      0      0      0      0
##          4      0      0      0      0      0      0      0      0      0
##          diagnosis
## wisc.hclust.clusters 10.86 10.88 10.9 10.91 10.94 10.95 10.96 10.97 11.04 11.06
##          1      0      0      0      0      0      0      0      0      0
##          2      1      1      1      1      1      1      1      1      2      3
##          3      0      0      0      0      0      0      0      0      0
##          4      0      0      0      0      0      0      0      0      0
##          diagnosis
## wisc.hclust.clusters 11.08 11.13 11.14 11.15 11.16 11.2 11.22 11.25 11.26 11.27
##          1      1      0      0      0      0      0      0      0      0
##          2      1      2      1      1      1      1      2      1      2      2
##          3      0      0      0      0      0      0      0      0      0
##          4      0      0      0      0      0      0      0      0      0
##          diagnosis
## wisc.hclust.clusters 11.28 11.29 11.3 11.31 11.32 11.33 11.34 11.36 11.37 11.41
##          1      0      0      0      0      0      0      0      0      0
##          2      1      1      1      1      1      1      2      1      1      2
##          3      0      0      0      0      0      0      0      0      0
##          4      0      0      0      0      0      0      0      0      0
##          diagnosis
## wisc.hclust.clusters 11.42 11.43 11.45 11.46 11.47 11.49 11.5 11.51 11.52 11.54
##          1      1      0      0      0      0      0      0      0      0
##          2      0      2      1      1      1      1      1      1      2      2
##          3      0      0      0      0      0      0      0      0      0
##          4      0      0      0      0      0      0      0      0      0
##          diagnosis
## wisc.hclust.clusters 11.57 11.6 11.61 11.62 11.63 11.64 11.66 11.67 11.68 11.69
##          1      0      0      0      0      0      0      0      0      0
##          2      1      3      1      1      1      1      1      1      1
##          3      0      0      0      0      0      0      0      0      0
##          4      0      0      0      0      0      0      0      0      0
##          diagnosis
## wisc.hclust.clusters 11.7 11.71 11.74 11.75 11.76 11.8 11.81 11.84 11.85 11.87
##          1      0      0      0      0      0      0      0      1      0      0
##          2      1      3      2      2      2      2      1      1      1      1

```

```

##          3    0    0    0    0    0    0    0    0    0    0
##          4    0    0    0    0    0    0    0    0    0    0
##          diagnosis
## wisc.hclust.clusters 11.89 11.9 11.93 11.94 11.95 11.99 12 12.03 12.04 12.05
##          1    0    0    0    0    0    0    0    0    0    0
##          2    3    1    2    2    1    1    2    1    1    2
##          3    0    0    0    0    0    0    0    0    0    0
##          4    0    0    0    0    0    0    0    0    0    0
##          diagnosis
## wisc.hclust.clusters 12.06 12.07 12.1 12.16 12.18 12.19 12.2 12.21 12.22 12.23
##          1    0    0    0    0    0    0    0    0    0    0
##          2    2    1    1    1    3    1    1    2    1    1
##          3    0    0    0    0    0    0    0    0    0    0
##          4    0    0    0    0    0    0    0    0    0    0
##          diagnosis
## wisc.hclust.clusters 12.25 12.27 12.3 12.31 12.32 12.34 12.36 12.39 12.4 12.42
##          1    0    0    0    0    0    1    0    0    0    0
##          2    2    2    2    1    1    3    2    1    1    1
##          3    0    0    0    0    0    0    0    0    0    0
##          4    0    0    0    0    0    0    0    0    0    0
##          diagnosis
## wisc.hclust.clusters 12.43 12.45 12.46 12.47 12.49 12.54 12.56 12.58 12.62
##          1    0    1    1    0    0    0    0    0    0
##          2    1    1    2    2    1    2    1    1    2
##          3    0    0    0    0    0    0    0    0    0
##          4    0    0    0    0    0    0    0    0    0
##          diagnosis
## wisc.hclust.clusters 12.63 12.65 12.67 12.68 12.7 12.72 12.75 12.76 12.77 12.78
##          1    0    0    0    1    0    0    0    0    0    0
##          2    1    1    1    0    1    2    1    2    3    1
##          3    0    0    0    0    0    0    0    0    0    0
##          4    0    0    0    0    0    0    0    0    0    0
##          diagnosis
## wisc.hclust.clusters 12.8 12.81 12.83 12.85 12.86 12.87 12.88 12.89 12.9 12.91
##          1    0    0    1    0    0    0    0    0    0    0
##          2    1    1    1    1    2    2    2    3    1    1
##          3    0    0    0    0    0    0    0    0    0    0
##          4    0    0    0    0    0    0    0    0    0    0
##          diagnosis
## wisc.hclust.clusters 12.94 12.95 12.96 12.98 12.99 13 13.01 13.03 13.05 13.08
##          1    0    0    0    0    0    1    0    0    0    0
##          2    1    1    1    1    1    2    1    1    3    1
##          3    0    0    0    0    0    0    0    0    0    0
##          4    0    0    0    0    0    0    0    0    0    0
##          diagnosis
## wisc.hclust.clusters 13.11 13.14 13.15 13.16 13.17 13.2 13.21 13.24 13.27 13.28
##          1    1    0    0    0    1    0    0    0    0    1
##          2    1    1    1    1    2    2    2    1    2    1
##          3    0    0    0    0    0    0    0    0    0    0
##          4    0    0    0    0    0    0    0    0    0    0
##          diagnosis
## wisc.hclust.clusters 13.3 13.34 13.37 13.38 13.4 13.43 13.44 13.45 13.46 13.47
##          1    0    1    0    0    1    0    0    0    0    0
##          2    1    0    1    1    1    1    1    1    2    1

```

```

##          3    0    0    0    0    0    0    0    0    0    0
##          4    0    0    0    0    0    0    0    0    0    0
##          diagnosis
## wisc.hclust.clusters 13.48 13.49 13.5 13.51 13.53 13.54 13.56 13.59 13.61 13.62
##          1    0    0    0    0    0    0    0    0    0    0
##          2    1    1    1    1    1    1    1    2    2    1
##          3    0    0    0    0    0    0    0    0    0    0
##          4    0    0    0    0    0    0    0    0    0    0
##          diagnosis
## wisc.hclust.clusters 13.64 13.65 13.66 13.68 13.69 13.7 13.71 13.73 13.74 13.75
##          1    0    0    0    0    0    0    1    1    0    0
##          2    2    1    2    1    1    1    1    0    1    1
##          3    0    0    0    0    0    0    0    0    0    0
##          4    0    0    0    0    0    0    0    0    0    0
##          diagnosis
## wisc.hclust.clusters 13.77 13.78 13.8 13.81 13.82 13.85 13.86 13.87 13.88 13.9
##          1    0    0    0    1    1    0    1    0    0    0
##          2    2    1    1    0    0    3    0    2    1    2
##          3    0    0    0    0    0    0    0    0    0    0
##          4    0    0    0    0    0    0    0    0    0    0
##          diagnosis
## wisc.hclust.clusters 13.94 13.96 13.98 14.02 14.03 14.04 14.05 14.06 14.11
##          1    0    0    0    0    0    0    0    0    0
##          2    1    1    1    1    1    1    1    1    1
##          3    0    0    0    0    0    0    0    0    0
##          4    0    0    0    0    0    0    0    0    0
##          diagnosis
## wisc.hclust.clusters 14.19 14.2 14.22 14.25 14.26 14.27 14.29 14.34 14.4 14.41
##          1    0    0    1    1    0    0    0    0    0
##          2    1    1    1    1    2    1    1    1    1
##          3    0    0    0    0    0    0    0    0    0
##          4    0    0    0    0    0    0    0    0    0
##          diagnosis
## wisc.hclust.clusters 14.42 14.44 14.45 14.47 14.48 14.5 14.53 14.54 14.58 14.59
##          1    0    0    0    0    0    0    0    1    1    0
##          2    2    1    1    1    1    1    2    0    1    1
##          3    0    0    0    0    0    0    0    0    0    0
##          4    0    0    0    0    0    0    0    0    0    0
##          diagnosis
## wisc.hclust.clusters 14.6 14.61 14.62 14.64 14.68 14.69 14.71 14.74 14.76 14.78
##          1    0    0    0    0    0    0    1    0    0    1
##          2    1    1    1    2    1    1    0    1    1    0
##          3    0    0    0    0    0    0    0    0    0    0
##          4    0    0    0    0    0    0    0    0    0    0
##          diagnosis
## wisc.hclust.clusters 14.8 14.81 14.86 14.87 14.9 14.92 14.95 14.96 14.97 14.99
##          1    0    0    0    1    1    0    1    0    0    0
##          2    1    1    2    1    0    1    1    1    2    2
##          3    0    0    0    0    0    0    0    0    0    0
##          4    0    0    0    0    0    0    0    0    0    0
##          diagnosis
## wisc.hclust.clusters 15 15.04 15.05 15.06 15.08 15.1 15.12 15.13 15.19 15.22
##          1 0    0    0    1    0    0    0    0    0    1
##          2 1    1    1    0    1    2    1    1    1    0

```

```

##          3 0    0    0    0    0    0    0    0    0    0
##          4 0    0    0    0    0    0    0    0    0    0
##          diagnosis
## wisc.hclust.clusters 15.27 15.28 15.3 15.32 15.34 15.37 15.46 15.49 15.5 15.53
##          1    0    0    1    1    0    0    2    1    1    1
##          2    1    1    0    0    1    1    1    0    0    0
##          3    0    0    0    0    0    0    0    0    0    0
##          4    0    0    0    0    0    0    0    0    0    0
##          diagnosis
## wisc.hclust.clusters 15.61 15.66 15.7 15.71 15.73 15.75 15.78 15.85 16.02 16.03
##          1    0    1    0    0    0    1    2    0    0    1
##          2    1    0    1    1    1    1    0    1    1    0
##          3    0    0    0    0    0    0    0    0    0    0
##          4    0    0    0    0    0    0    0    0    0    0
##          diagnosis
## wisc.hclust.clusters 16.07 16.11 16.13 16.14 16.16 16.17 16.24 16.25 16.26
##          1    0    0    2    0    0    0    1    0    0
##          2    1    1    0    1    1    1    0    1    1
##          3    0    0    0    0    0    0    0    0    0
##          4    0    0    0    0    0    0    0    0    0
##          diagnosis
## wisc.hclust.clusters 16.27 16.3 16.35 16.46 16.5 16.6 16.65 16.69 16.74 16.78
##          1    0    0    1    1    0    0    0    0    1    0
##          2    1    1    0    0    1    1    1    1    0    1
##          3    0    0    0    0    0    0    0    0    0    0
##          4    0    0    0    0    0    0    0    0    0    0
##          diagnosis
## wisc.hclust.clusters 16.84 17.01 17.02 17.05 17.06 17.08 17.14 17.19 17.2 17.27
##          1    0    0    1    1    0    0    1    0    1    0
##          2    1    1    0    0    1    1    0    1    0    1
##          3    0    0    0    0    0    0    0    0    0    0
##          4    0    0    0    0    0    0    0    0    0    0
##          diagnosis
## wisc.hclust.clusters 17.29 17.3 17.35 17.42 17.46 17.47 17.54 17.57 17.6 17.68
##          1    1    0    0    0    0    1    0    0    1    0
##          2    0    1    1    1    1    0    1    1    0    1
##          3    0    0    0    0    0    0    0    0    0    0
##          4    0    0    0    0    0    0    0    0    0    0
##          diagnosis
## wisc.hclust.clusters 17.75 17.85 17.91 17.93 17.95 17.99 18.01 18.03 18.05
##          1    0    0    1    0    0    1    0    0    1
##          2    1    1    0    1    1    1    1    1    0
##          3    0    0    0    0    0    0    0    0    0
##          4    0    0    0    0    0    0    0    0    0
##          diagnosis
## wisc.hclust.clusters 18.08 18.22 18.25 18.31 18.45 18.46 18.49 18.61 18.63
##          1    0    0    0    0    0    0    0    0    1
##          2    1    2    1    2    1    1    1    1    0
##          3    0    0    0    0    0    0    0    0    0
##          4    0    0    0    0    0    0    0    0    0
##          diagnosis
## wisc.hclust.clusters 18.65 18.66 18.77 18.81 18.82 18.94 19 19.02 19.07 19.1
##          1    1    0    0    0    0    0 0    0    1    1
##          2    0    1    1    1    1    1 1    1    0    0

```

```

##          3      0      0      0      0      0      0      0      0      0      0
##          4      0      0      0      0      0      0      0      0      0      0
##          diagnosis
## wisc.hclust.clusters 19.16 19.17 19.18 19.19 19.21 19.27 19.4 19.44 19.45 19.53
##          1      1      1      0      1      0      1      0      1      0      1
##          2      0      0      1      0      1      0      2      0      1      1
##          3      0      0      0      0      0      0      0      0      0      0
##          4      0      0      0      0      0      0      0      0      0      0
##          diagnosis
## wisc.hclust.clusters 19.55 19.59 19.68 19.69 19.73 19.79 19.8 19.81 19.89 20.09
##          1      2      1      1      1      1      1      1      0      0      1
##          2      0      1      0      0      0      0      0      1      1      0
##          3      0      0      0      0      0      0      0      0      0      0
##          4      0      0      0      0      0      0      0      0      0      0
##          diagnosis
## wisc.hclust.clusters 20.13 20.16 20.18 20.2 20.26 20.29 20.31 20.34 20.44 20.47
##          1      0      0      2      1      1      0      0      1      0      1
##          2      1      1      0      0      0      1      1      0      1      0
##          3      0      0      0      0      0      0      0      0      0      0
##          4      0      0      0      0      0      0      0      0      0      0
##          diagnosis
## wisc.hclust.clusters 20.48 20.51 20.55 20.57 20.58 20.59 20.6 20.64 20.73 20.92
##          1      0      0      1      0      1      1      1      0      1      1
##          2      1      1      0      1      0      0      0      1      0      0
##          3      0      0      0      0      0      0      0      0      0      0
##          4      0      0      0      0      0      0      0      0      0      0
##          diagnosis
## wisc.hclust.clusters 20.94 21.09 21.1 21.16 21.37 21.56 21.61 21.71 21.75 22.01
##          1      1      1      0      0      0      1      1      1      1      1
##          2      0      0      1      1      1      0      0      0      0      0
##          3      0      0      0      0      0      0      0      0      0      0
##          4      0      0      0      0      0      0      0      0      0      0
##          diagnosis
## wisc.hclust.clusters 22.27 23.09 23.21 23.27 23.29 23.51 24.25 24.63 25.22
##          1      1      1      1      0      1      1      1      1      1
##          2      0      0      0      1      0      0      0      0      0
##          3      0      0      0      0      0      0      0      0      0
##          4      0      0      0      0      0      0      0      0      0
##          diagnosis
## wisc.hclust.clusters 25.73 27.22 27.42 28.11
##          1      1      1      0      0
##          2      0      0      0      0
##          3      0      0      0      0
##          4      0      0      1      1

```

```

grps <- cutree(wisc.hclust, k=2)
table(grps)

```

```

## grps
##    1    2
## 567    2

```

```
table(grps, diagnosis)
```

```
##      diagnosis
## grps 6.981 7.691 7.729 7.76 8.196 8.219 8.571 8.597 8.598 8.618 8.671 8.726
##    1    1    1    1    1    1    1    1    1    1    1    1
##    2    0    0    0    0    0    0    0    0    0    0    0
##      diagnosis
## grps 8.734 8.878 8.888 8.95 9 9.029 9.042 9.173 9.268 9.295 9.333 9.397 9.405
##    1    1    1    1    1 1    1    1    1    1    1    1    1
##    2    0    0    0    0 0    0    0    0    0    0    0    0
##      diagnosis
## grps 9.423 9.436 9.465 9.504 9.567 9.606 9.667 9.668 9.676 9.683 9.72 9.731
##    1    1    1    1    1    1    1    1    1    1    1    1
##    2    0    0    0    0    0    0    0    0    0    0    0
##      diagnosis
## grps 9.738 9.742 9.755 9.777 9.787 9.847 9.876 9.904 10.03 10.05 10.08 10.16
##    1    1    2    1    1    1    1    2    1    1    1    1
##    2    0    0    0    0    0    0    0    0    0    0    0
##      diagnosis
## grps 10.17 10.18 10.2 10.25 10.26 10.29 10.32 10.44 10.48 10.49 10.51 10.57
##    1    1    1    1    1    3    1    1    1    2    2    2
##    2    0    0    0    0    0    0    0    0    0    0    0
##      diagnosis
## grps 10.6 10.65 10.66 10.71 10.75 10.8 10.82 10.86 10.88 10.9 10.91 10.94 10.95
##    1    1    1    1    1    1    2    1    1    1    1    1    1
##    2    0    0    0    0    0    0    0    0    0    0    0    0
##      diagnosis
## grps 10.96 10.97 11.04 11.06 11.08 11.13 11.14 11.15 11.16 11.2 11.22 11.25
##    1    1    1    2    3    2    2    1    1    1    1    2
##    2    0    0    0    0    0    0    0    0    0    0    0
##      diagnosis
## grps 11.26 11.27 11.28 11.29 11.3 11.31 11.32 11.33 11.34 11.36 11.37 11.41
##    1    2    2    1    1    1    1    1    1    2    1    2
##    2    0    0    0    0    0    0    0    0    0    0    0
##      diagnosis
## grps 11.42 11.43 11.45 11.46 11.47 11.49 11.5 11.51 11.52 11.54 11.57 11.6
##    1    1    2    1    1    1    1    1    1    2    2    1
##    2    0    0    0    0    0    0    0    0    0    0    0
##      diagnosis
## grps 11.61 11.62 11.63 11.64 11.66 11.67 11.68 11.69 11.7 11.71 11.74 11.75
##    1    1    1    1    1    1    1    1    1    1    3    2
##    2    0    0    0    0    0    0    0    0    0    0    0
##      diagnosis
## grps 11.76 11.8 11.81 11.84 11.85 11.87 11.89 11.9 11.93 11.94 11.95 11.99 12
##    1    2    2    1    2    1    1    3    1    2    2    1    2
##    2    0    0    0    0    0    0    0    0    0    0    0    0
##      diagnosis
## grps 12.03 12.04 12.05 12.06 12.07 12.1 12.16 12.18 12.19 12.2 12.21 12.22
##    1    1    1    2    2    1    1    1    3    1    1    2
##    2    0    0    0    0    0    0    0    0    0    0    0
##      diagnosis
## grps 12.23 12.25 12.27 12.3 12.31 12.32 12.34 12.36 12.39 12.4 12.42 12.43
##    1    1    2    2    2    1    1    4    2    1    1    1
```



```

##      2      0      0      0      0      0      0      0      0      0      0      0      0
##      diagnosis
## grps 12.45 12.46 12.47 12.49 12.54 12.56 12.58 12.62 12.63 12.65 12.67 12.68
##      1      2      3      2      1      2      1      1      2      1      1      1      1
##      2      0      0      0      0      0      0      0      0      0      0      0      0
##      diagnosis
## grps 12.7 12.72 12.75 12.76 12.77 12.78 12.8 12.81 12.83 12.85 12.86 12.87
##      1      1      2      1      2      3      1      1      1      2      1      2      2
##      2      0      0      0      0      0      0      0      0      0      0      0      0
##      diagnosis
## grps 12.88 12.89 12.9 12.91 12.94 12.95 12.96 12.98 12.99 13 13.01 13.03 13.05
##      1      2      3      1      1      1      1      1      1      1      3      1      3
##      2      0      0      0      0      0      0      0      0      0      0      0      0
##      diagnosis
## grps 13.08 13.11 13.14 13.15 13.16 13.17 13.2 13.21 13.24 13.27 13.28 13.3
##      1      1      2      1      1      1      3      2      2      1      2      2      1
##      2      0      0      0      0      0      0      0      0      0      0      0      0
##      diagnosis
## grps 13.34 13.37 13.38 13.4 13.43 13.44 13.45 13.46 13.47 13.48 13.49 13.5
##      1      1      1      1      2      1      1      1      2      1      1      1      1
##      2      0      0      0      0      0      0      0      0      0      0      0      0
##      diagnosis
## grps 13.51 13.53 13.54 13.56 13.59 13.61 13.62 13.64 13.65 13.66 13.68 13.69
##      1      1      1      1      1      2      2      1      2      1      2      1      1
##      2      0      0      0      0      0      0      0      0      0      0      0      0
##      diagnosis
## grps 13.7 13.71 13.73 13.74 13.75 13.77 13.78 13.8 13.81 13.82 13.85 13.86
##      1      1      2      1      1      1      2      1      1      1      1      3      1
##      2      0      0      0      0      0      0      0      0      0      0      0      0
##      diagnosis
## grps 13.87 13.88 13.9 13.94 13.96 13.98 14.02 14.03 14.04 14.05 14.06 14.11
##      1      2      1      2      1      1      1      1      1      1      1      1      1
##      2      0      0      0      0      0      0      0      0      0      0      0      0
##      diagnosis
## grps 14.19 14.2 14.22 14.25 14.26 14.27 14.29 14.34 14.4 14.41 14.42 14.44
##      1      1      1      2      2      2      1      1      1      1      1      2      1
##      2      0      0      0      0      0      0      0      0      0      0      0      0
##      diagnosis
## grps 14.45 14.47 14.48 14.5 14.53 14.54 14.58 14.59 14.6 14.61 14.62 14.64
##      1      1      1      1      1      2      1      2      1      1      1      1      2
##      2      0      0      0      0      0      0      0      0      0      0      0      0
##      diagnosis
## grps 14.68 14.69 14.71 14.74 14.76 14.78 14.8 14.81 14.86 14.87 14.9 14.92
##      1      1      1      1      1      1      1      1      1      2      2      1      1
##      2      0      0      0      0      0      0      0      0      0      0      0      0
##      diagnosis
## grps 14.95 14.96 14.97 14.99 15 15.04 15.05 15.06 15.08 15.1 15.12 15.13 15.19
##      1      2      1      2      2      1      1      1      1      1      2      1      1
##      2      0      0      0      0      0      0      0      0      0      0      0      0
##      diagnosis
## grps 15.22 15.27 15.28 15.3 15.32 15.34 15.37 15.46 15.49 15.5 15.53 15.61
##      1      1      1      1      1      1      1      1      3      1      1      1      1
##      2      0      0      0      0      0      0      0      0      0      0      0      0
##      diagnosis

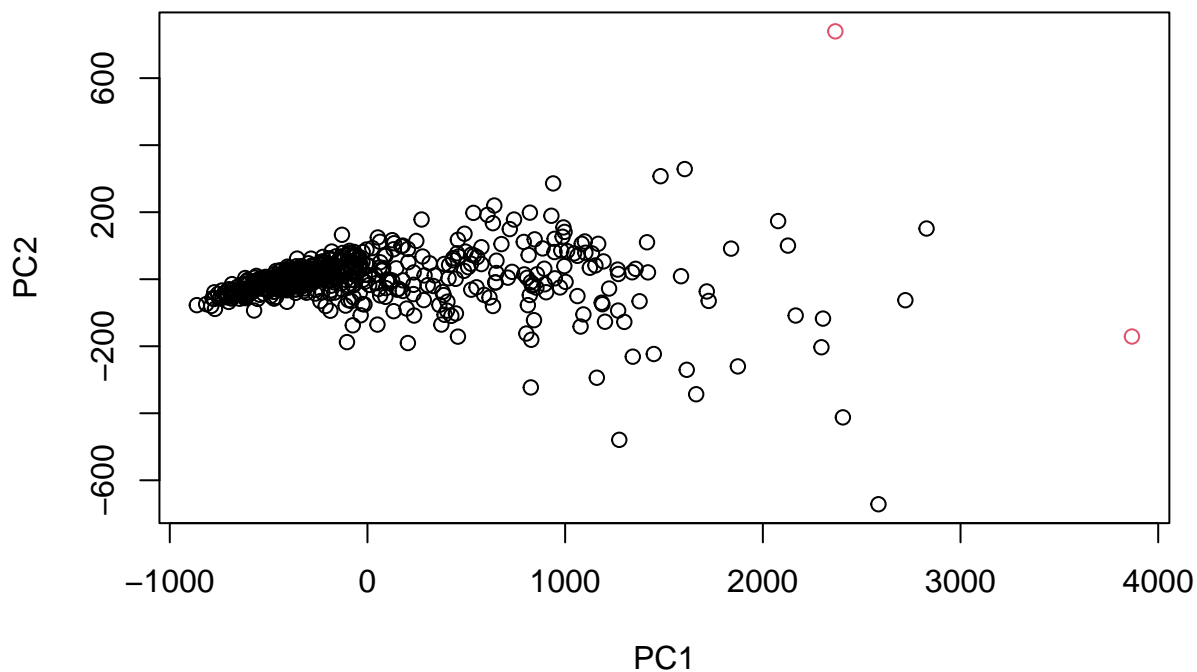
```

```

## grps 15.66 15.7 15.71 15.73 15.75 15.78 15.85 16.02 16.03 16.07 16.11 16.13
## 1 1 1 1 1 2 2 1 1 1 1 2
## 2 0 0 0 0 0 0 0 0 0 0 0
## diagnosis
## grps 16.14 16.16 16.17 16.24 16.25 16.26 16.27 16.3 16.35 16.46 16.5 16.6 16.65
## 1 1 1 1 1 1 1 1 1 1 1 1 1
## 2 0 0 0 0 0 0 0 0 0 0 0 0
## diagnosis
## grps 16.69 16.74 16.78 16.84 17.01 17.02 17.05 17.06 17.08 17.14 17.19 17.2
## 1 1 1 1 1 1 1 1 1 1 1 1
## 2 0 0 0 0 0 0 0 0 0 0 0
## diagnosis
## grps 17.27 17.29 17.3 17.35 17.42 17.46 17.47 17.54 17.57 17.6 17.68 17.75
## 1 1 1 1 1 1 1 1 1 1 1 1
## 2 0 0 0 0 0 0 0 0 0 0 0
## diagnosis
## grps 17.85 17.91 17.93 17.95 17.99 18.01 18.03 18.05 18.08 18.22 18.25 18.31
## 1 1 1 1 1 2 1 1 1 1 2 1 2
## 2 0 0 0 0 0 0 0 0 0 0 0 0
## diagnosis
## grps 18.45 18.46 18.49 18.61 18.63 18.65 18.66 18.77 18.81 18.82 18.94 19 19.02
## 1 1 1 1 1 1 1 1 1 1 1 1 1
## 2 0 0 0 0 0 0 0 0 0 0 0 0
## diagnosis
## grps 19.07 19.1 19.16 19.17 19.18 19.19 19.21 19.27 19.4 19.44 19.45 19.53
## 1 1 1 1 1 1 1 1 1 2 1 1 2
## 2 0 0 0 0 0 0 0 0 0 0 0 0
## diagnosis
## grps 19.55 19.59 19.68 19.69 19.73 19.79 19.8 19.81 19.89 20.09 20.13 20.16
## 1 2 2 1 1 1 1 1 1 1 1 1
## 2 0 0 0 0 0 0 0 0 0 0 0
## diagnosis
## grps 20.18 20.2 20.26 20.29 20.31 20.34 20.44 20.47 20.48 20.51 20.55 20.57
## 1 2 1 1 1 1 1 1 1 1 1 1
## 2 0 0 0 0 0 0 0 0 0 0 0
## diagnosis
## grps 20.58 20.59 20.6 20.64 20.73 20.92 20.94 21.09 21.1 21.16 21.37 21.56
## 1 1 1 1 1 1 1 1 1 1 1 1
## 2 0 0 0 0 0 0 0 0 0 0 0
## diagnosis
## grps 21.61 21.71 21.75 22.01 22.27 23.09 23.21 23.27 23.29 23.51 24.25 24.63
## 1 1 1 1 1 1 1 1 1 1 1 1
## 2 0 0 0 0 0 0 0 0 0 0 0
## diagnosis
## grps 25.22 25.73 27.22 27.42 28.11
## 1 1 1 1 0 0
## 2 0 0 0 1 1

```

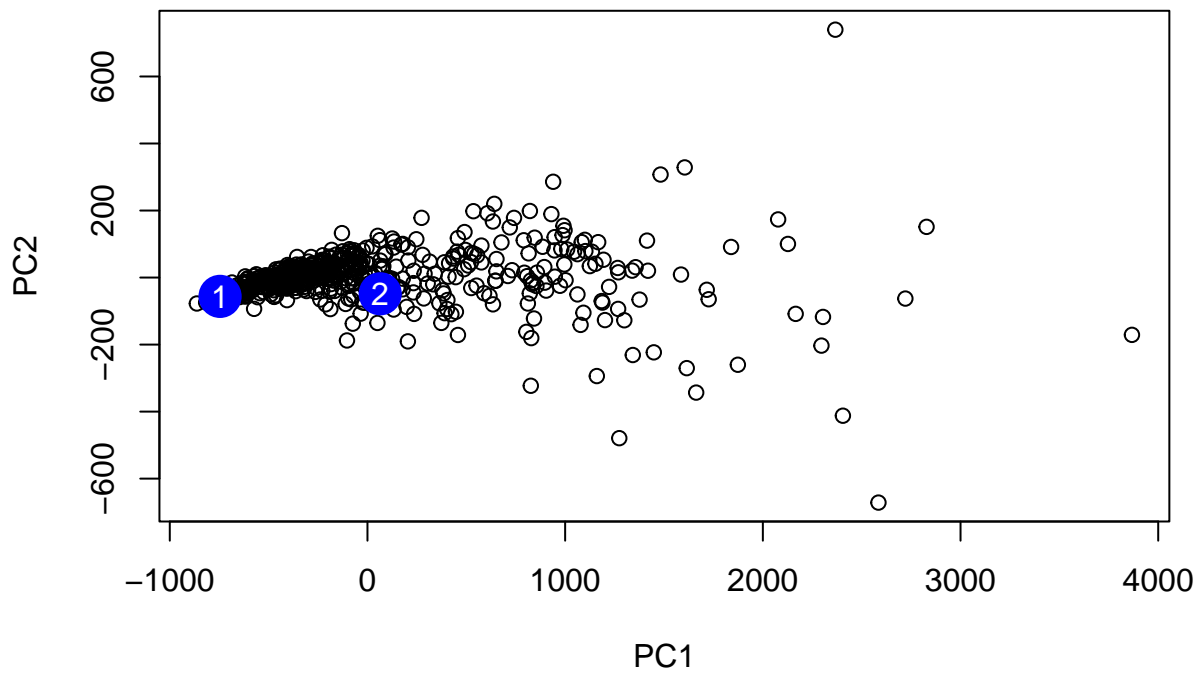
```
plot(wisc.pr$x[,1:2], col=grps)
```



```
url <- "https://tinyurl.com/new-samples-CSV"
new <- read.csv(url)
npc <- predict(wisc.pr, newdata=new)
npc
```

```
##          PC1          PC2          PC3          PC4          PC5          PC6          PC7
## [1,] -745.58228 -56.15114 -21.14308 -3.269575  9.342062  2.265565 -1.156815
## [2,]  64.40863 -48.46994 -15.93951 12.065347 -4.676632 -1.080107 -0.277376
##          PC8          PC9          PC10          PC11          PC12          PC13
## [1,]  0.7563343 -0.1032757  0.0688537 -0.09202633  0.06375044 -0.01669926
## [2,]  0.7260377  0.2276038 -0.7097099 -0.30447319 -0.01883250  0.11744318
##          PC14          PC15          PC16          PC17          PC18
## [1,] -0.023223923  0.009950655 -0.008509815 -0.01948388 -0.007793169
## [2,] -0.007291041 -0.018657551 -0.011000041 -0.02647465  0.000611957
##          PC19          PC20          PC21          PC22          PC23
## [1,] -0.031830403 -0.005101987  0.01845962 -0.0005420926  0.006199072
## [2,]  0.009654825 -0.018181800  0.01048644  0.0088785987 -0.007183325
##          PC24          PC25          PC26          PC27          PC28
## [1,]  0.005901854 -0.005404442 -0.0001697478 -0.003258087 -0.0012304929
## [2,] -0.003020084 -0.003319119 -0.0021016368  0.002101994 -0.0009562871
##          PC29
## [1,]  0.0009908106
## [2,]  0.0008586404
```

```
plot(wisc.pr$x[,1:2])
points(npc[,1], npc[,2], col="blue", pch=16, cex=3)
text(npc[,1], npc[,2], c(1,2), col="white")
```



```
sessionInfo()
```

```
## R version 4.1.2 (2021-11-01)
## Platform: x86_64-w64-mingw32/x64 (64-bit)
## Running under: Windows 10 x64 (build 19043)
##
## Matrix products: default
##
## locale:
## [1] LC_COLLATE=English_United States.1252
## [2] LC_CTYPE=English_United States.1252
## [3] LC_MONETARY=English_United States.1252
## [4] LC_NUMERIC=C
## [5] LC_TIME=English_United States.1252
##
## attached base packages:
## [1] stats      graphics  grDevices  utils      datasets  methods   base
##
## other attached packages:
## [1] ggplot2_3.3.5
##
```

```
## loaded via a namespace (and not attached):
## [1] pillar_1.7.0      compiler_4.1.2    highr_0.9         tools_4.1.2
## [5] digest_0.6.29     evaluate_0.15     lifecycle_1.0.1   tibble_3.1.6
## [9] gtable_0.3.0      pkgconfig_2.0.3   rlang_1.0.2       cli_3.1.1
## [13] rstudioapi_0.13   yaml_2.3.5        xfun_0.30         fastmap_1.1.0
## [17] withr_2.5.0       stringr_1.4.0     dplyr_1.0.8       knitr_1.37
## [21] generics_0.1.2    vctrs_0.3.8       grid_4.1.2        tidyselect_1.1.2
## [25] glue_1.6.2        R6_2.5.1          fansi_1.0.2       rmarkdown_2.12
## [29] farver_2.1.0      purrr_0.3.4       magrittr_2.0.2    scales_1.1.1
## [33] ellipsis_0.3.2    htmltools_0.5.2   colorspace_2.0-3  labeling_0.4.2
## [37] utf8_1.2.2        stringi_1.7.6     munsell_0.5.0     crayon_1.5.0
```