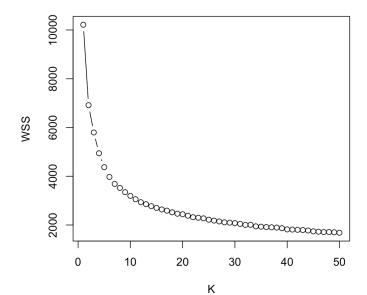
```
1.
(a)
> tree_health_data = tree(readmit30~.-readmit30, healthdata)
> plot(tree_health_data)
> text(tree_health_data,pretty=0)
> tree_health_data
node), split, n, deviance, yval, (yprob)
       * denotes terminal node
                                                                                       comorbidity.score < 121.5
1) root 4382 4702.0 0 ( 0.77225 0.22775 )
  2) comorbidity.score < 121.5 3084 2440.0 0 ( 0.86511 0.13489 )
     4) severity.score < 31.5 2423 1609.0 0 ( 0.89682 0.10318 )
       8) comorbidity.score < 68.5 1337 616.7 0 ( 0.93867 0.06133 ) *
       9) comorbidity.score > 68.5 1086 935.6 0 ( 0.84530 0.15470 ) *
     5) severity.score > 31.5 661 745.1 0 ( 0.74887 0.25113 ) *
                                                                                  severity.score < 31.5comorbidity.score < 167.5
   3) comorbidity.score > 121.5 1298 1786.0 0 ( 0.55162 0.44838 )
                                                                            comorbidity.score < 68.5
     6) comorbidity.score < 167.5 770 1001.0 0 ( 0.64545 0.35455 ) *
                                                                                ò
     7) comorbidity.score > 167.5 528 716.5 1 ( 0.41477 0.58523 ) *
(b)
> cv.health
$size
[1] 5 4 3 2 1
$dev
[1] 4056.280 4131.010 4167.384 4234.041 4704.065
$k
[1]
         -Inf 56.62428 67.67316 85.95055 476.79161
$method
[1] "deviance"
attr(,"class")
[1] "prune"
                   "tree.sequence"
(c)
> healthdata$caretrack = 1
> healthdata[comorbidity.score<68.5 & severity.score<31.5,]$caretrack = 0</pre>
> as.double(sum(healthdata$caretrack==1) / nrow(healthdata))
[1] 0.6948882
(d) The unawareness difference between male and female is about 0.0395, while the accuracy
parity difference is about 0.0745
> # unawareness
> nrow(female_care)/nrow(female) - nrow(care)/nrow(healthdata)
[1] 0.03954762
> # Accuracy Parity
> nrow(female_care)/nrow(female) - nrow(male_care)/nrow(male)
[1] 0.07450459
```

```
2.
(a)
> linReg = lm(SAT_AVG~.-INSTNM, data=clgdata)
> summary(linReg)
Call:
lm(formula = SAT_AVG ~ . - INSTNM, data = clgdata)
Residuals:
    Min
             10 Median
                             30
                                    Max
-254.21 -44.63
                   3.83
                          45.58 326.06
Coefficients:
                 Estimate Std. Error t value Pr(>|t|)
(Intercept)
                6.515e+02 1.165e+01 55.941 < 2e-16 ***
UGDS
                                      0.538 0.59089
                2.019e-04 3.755e-04
                                       0.065 0.94828
COSTT4_A
                3.288e-05 5.068e-04
TUITIONFEE_OUT 1.798e-03 6.596e-04
                                       2.725 0.00653 **
TUITFTE
               -7.138e-04 6.609e-04
                                     -1.080 0.28034
AVGFACSAL
                1.626e-02 1.453e-03
                                     11.197 < 2e-16 ***
PFTFAC
                4.090e+01 9.069e+00
                                      4.510 7.16e-06 ***
C150_4
                4.226e+02 1.962e+01 21.533 < 2e-16 ***
PFTFTUG1_EF
               -1.672e+01 1.375e+01 -1.216 0.22419
Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
Residual standard error: 70.62 on 1127 degrees of freedom
                               Adjusted R-squared: 0.6817
Multiple R-squared: 0.6839,
F-statistic: 304.8 on 8 and 1127 DF, p-value: < 2.2e-16
```

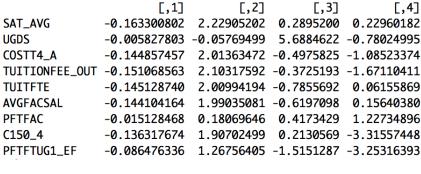
(b) I will choose k=10, because WSS[10] is quite small and k=10 is not large.

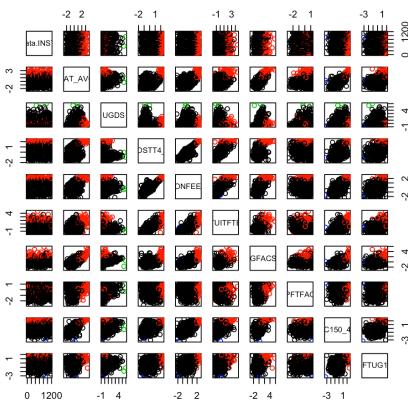
```
> wss
[1] 10215.000 6914.788 5795.651 4942.431 4373.990 3974.122 3684.776 3506.198 3353.539 3200.516 3068.700
[12] 2936.936 2854.959 2774.736 2709.847 2644.923 2590.017 2537.823 2470.561 2429.626 2399.490 2344.324
[23] 2282.184 2256.681 2205.501 2183.340 2133.925 2118.222 2086.069 2071.296 2026.466 2019.438 1977.567
[34] 1963.573 1934.162 1926.669 1892.010 1882.702 1853.873 1832.858 1826.580 1796.540 1790.871 1771.559
[45] 1762.516 1731.407 1718.260 1711.130 1685.122 1672.938

> which.min(wss)
[1] 50
```



## (c) Four centroids are as below

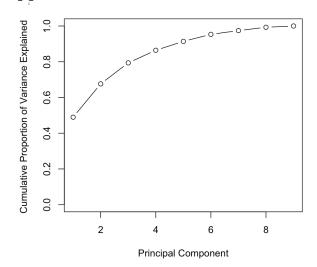




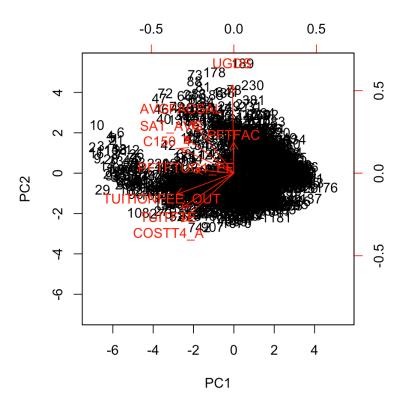
## (d) PVE values and cumulative PVE plot is as below.

> pve

 $\hbox{\tt [1]} \ \ 0.489610289 \ \ 0.186469957 \ \ 0.117469839 \ \ 0.070554030 \ \ 0.049739869 \ \ 0.039294997 \ \ 0.020962470 \ \ 0.018814019 \ \ 0.007084531$ 

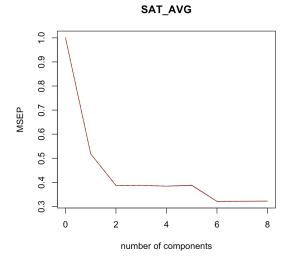


(e) PC1 indicates a linear combination of COSTT4\_A, TUITFTE, TUITIONFEE\_OUT, PFTFTUG1\_EF, C150\_4, SAT\_AVG and AVGFACSAL, and obviously that PFTFTUG1\_EF plays an important part on it. While PC2 indicates another linear combination of UGDS, PFTFTUG1\_EF, C150\_4, SAT\_AVG, AVGFACSAL, COSTT4\_A, TUITFTE and TUITIONFEE\_OUT, and obviously that UGDS, PFTFTUG1\_EF play an important part on it. In addition, PC1 and PC2 are orthogonal.



(f) Below is the summary for PCR, I will choose M=5 because it explains 92% of the variance.

```
> summary(pcr.fit)
       X dimension: 1136 8
        Y dimension: 1136 1
Fit method: svdpc
Number of components considered: 8
VALIDATION: RMSEP
Cross-validated using 5 random segments.
       (Intercept) 1 comps 2 comps
                                      3 comps 4 comps
                                                        5 comps
                                                                           7 comps
                                                                  6 comps
C۷
                     0.7201
                              0.6223
                                       0.6227
                                                0.6199
                                                          0.6230
                                                                   0.5669
                                                                            0.5674
                                                                                     0.5677
adjCV
                 1
                     0.7198
                              0.6221
                                       0.6224
                                                0.6196
                                                         0.6219
                                                                   0.5663
                                                                            0.5668
                                                                                     0.5671
TRAINING: % variance explained
         1 comps 2 comps 3 comps
                                    4 comps
                                             5 comps
                                                      6 comps
                                                               7 comps
                                                                         8 comps
X
           48.09
                                                        97.07
                                                                  99.20
                                                                          100.00
                    67.41
                             80.62
                                      88.49
                                               92.92
SAT_AVG
           48.38
                    61.45
                             61.45
                                      61.79
                                               62.11
                                                        68.35
                                                                  68.38
                                                                           68.39
```



Below is the summary for PLS, I will choose M=6 because it explains 94% of the variance.

> summary(pls.fit)

Data: X dimension: 1136 8 Y dimension: 1136 1

Fit method: kernelpls

Number of components considered: 8

VALIDATION: RMSEP

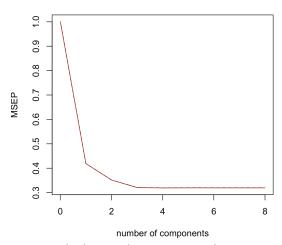
Cross-validated using 5 random segments.

	(Intercept)	1 comps	2 comps	3 comps	4 comps	5 comps	6 comps	7 comps	8 comps
CV	1	0.6474	0.5931	0.5664	0.5649	0.5652	0.5652	0.5652	0.5652
adjCV	1	0.6473	0.5929	0.5658	0.5646	0.5649	0.5648	0.5648	0.5648

TRAINING: % variance explained

	1 comps	2 comps	3 comps	4 comps	5 comps	6 comps	7 comps	8 comps
Χ	46.84	66.53	71.87	79.44	87.68	94.64	97.73	100.00
SAT AVG	58.23	65.10	68.32	68.38	68.39	68.39	68.39	68.39

## SAT\_AVG



PCR works better because it only requires M=5 while PCR asks for M=6