Class08: Machine Learning (Mini Project)

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Breast Cancer Project

Today we are going to explore some data from the University of Wisconsin Cancer Center on Breast biopsy data.

```
wisc.data<-read.csv("WisconsinCancer.csv", row.names=1)
head(wisc.data)</pre>
```

	diagnosis	radius_mean	${\tt texture_mean}$	<pre>perimeter_mean</pre>	area_mean	L
842302	M	17.99	10.38	122.80	1001.0)
842517	M	20.57	17.77	132.90	1326.0)
84300903	M	19.69	21.25	130.00	1203.0)
84348301	M	11.42	20.38	77.58	386.1	
84358402	M	20.29	14.34	135.10	1297.0)
843786	M	12.45	15.70	82.57	477.1	
	smoothness	s_mean compa	ctness_mean co	oncavity_mean co	oncave.poi	nts_mean
842302	0	.11840	0.27760	0.3001		0.14710
842517	0	.08474	0.07864	0.0869		0.07017
84300903	0	.10960	0.15990	0.1974		0.12790
84348301	0	. 14250	0.28390	0.2414		0.10520
84358402	0	.10030	0.13280	0.1980		0.10430
843786	0	. 12780	0.17000	0.1578		0.08089
symmetry_mean fractal_dimension_mean radius_se texture_se perimeter_se						
842302	0.2	2419	0.0787	1.0950	0.9053	8.589
842517	0.1	1812	0.0566	0.5435	0.7339	3.398
84300903	0.2	2069	0.0599	0.7456	0.7869	4.585
84348301	0.2	2597	0.0974	0.4956	1.1560	3.445
84358402	0.1	1809	0.0588	0.7572	0.7813	5.438
843786	0.2	2087	0.0761	.3 0.3345	0.8902	2.217
	area_se sm	noothness_se	compactness_s	se concavity_se	concave.p	oints_se
842302	153.40	0.006399	0.0490	0.05373		0.01587

842517	74.08	0.005225	0.01308	0.01860	0.01340
84300903	94.03	0.006150	0.04006	0.03832	0.02058
84348301	27.23	0.009110	0.07458	0.05661	0.01867
84358402	94.44	0.011490	0.02461	0.05688	0.01885
843786	27.19	0.007510	0.03345	0.03672	0.01137
	symmetry_se	fractal_dime	nsion_se rad	ius_worst text	ure_worst
842302	0.03003		0.006193	25.38	17.33
842517	0.01389		0.003532	24.99	23.41
84300903	0.02250		0.004571	23.57	25.53
84348301	0.05963		0.009208	14.91	26.50
84358402	0.01756		0.005115	22.54	16.67
843786	0.02165		0.005082	15.47	23.75
	perimeter_wo	rst area_wor	st smoothnes	s_worst compac	tness_worst
842302	184	.60 2019	0.0	0.1622	0.6656
842517	158	.80 1956	3.0	0.1238	0.1866
84300903	152	.50 1709	0.0	0.1444	0.4245
84348301	98	.87 567	7.7	0.2098	0.8663
84358402	152	.20 1575	5.0	0.1374	0.2050
843786	103	.40 741	6	0.1791	0.5249
	concavity_wo	rst concave.	points_worst	symmetry_wors	t
842302	0.7	119	0.2654	0.460	1
842517	0.2	416	0.1860	0.275	0
84300903	0.4	504	0.2430	0.361	3
84348301	0.6	869	0.2575	0.663	8
84358402	0.4	000	0.1625	0.236	4
843786	0.5	355	0.1741	0.398	5
	<pre>fractal_dime</pre>	nsion_worst			
842302		0.11890			
842517		0.08902			
84300903		0.08758			
84348301		0.17300			
84358402		0.07678			
843786		0.12440			

Q1. How many patient samples are in this dataset?

```
nrow(wisc.data)
```

[1] 569

There are 569 patients in this dataset.

Q2. How many cancer(M) and non cancer (B) samples are there?

```
table(wisc.data$diagnosis)
```

```
B M
357 212
```

There are 357 benign and 212 malignant cells.

Save the diagnosis for later use as a reference to compare how well we do with PCA etc.

```
diagnosis <- as.factor(wisc.data$diagnosis)
#diagnosis</pre>
```

Now exclude the diagnosis column from the data.

```
wisc <- wisc.data[,-1]
```

Q3. How many 'dimensions', 'variables', 'columns' are there in this dataset?

```
ncol(wisc)
```

[1] 30

We have 30 variables in this dataset.

Performing Principal Component Analysis(PCA)

To perform PCA in R, we can use the prcomp() function. It takes as input a numerical dataset and optional scale=FALSE/TRUE argument.

We generally always want to set scale=TRUE but let's make sure by checking if the mean and standard deviation values are different across these 30 columns.

```
# Check column means and standard deviations
round(colMeans(wisc))
```

radius_mean	texture_mean	perimeter_mean
14	19	92
area_mean	${\tt smoothness_mean}$	compactness_mean
655	0	0
concavity_mean	concave.points_mean	symmetry_mean
0	0	0
fractal_dimension_mean	radius_se	texture_se
0	0	1
perimeter_se	area_se	smoothness_se
3	40	0
compactness_se	concavity_se	concave.points_se
0	0	0
symmetry_se	fractal_dimension_se	radius_worst
0	0	16
texture_worst	perimeter_worst	area_worst
26	107	881
${\tt smoothness_worst}$	compactness_worst	${\tt concavity_worst}$
0	0	0
concave.points_worst	symmetry_worst	${\tt fractal_dimension_worst}$
0	0	0

round(apply(wisc,2,sd))

radius_mean	texture_mean	perimeter_mean
4	4	24
area_mean	smoothness_mean	compactness_mean
352	0	0
${\tt concavity_mean}$	concave.points_mean	$\operatorname{symmetry_mean}$
0	0	0
${\tt fractal_dimension_mean}$	radius_se	texture_se
0	0	1
perimeter_se	area_se	${\tt smoothness_se}$
2	45	0
compactness_se	concavity_se	concave.points_se
0	0	0
symmetry_se	fractal_dimension_se	${\tt radius_worst}$
0	0	5
texture_worst	perimeter_worst	area_worst
6	34	569
${\tt smoothness_worst}$	compactness_worst	concavity_worst
0	0	0

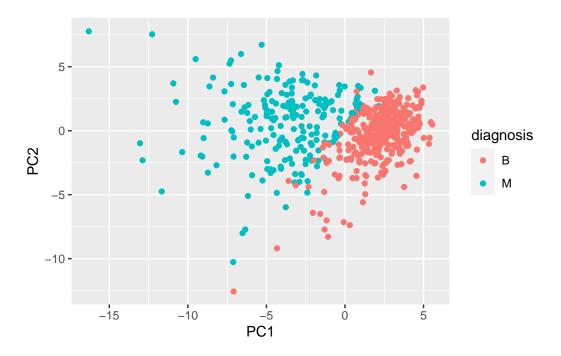
```
symmetry_worst fractal_dimension_worst
   concave.points_worst
                      0
  pca <- prcomp(wisc,scale=TRUE)</pre>
  summary(pca)
Importance of components:
                                                                  PC6
                                 PC2
                                         PC3
                                                  PC4
                                                          PC5
                          PC1
                                                                          PC7
                       3.6444 2.3857 1.67867 1.40735 1.28403 1.09880 0.82172
Standard deviation
Proportion of Variance 0.4427 0.1897 0.09393 0.06602 0.05496 0.04025 0.02251
Cumulative Proportion 0.4427 0.6324 0.72636 0.79239 0.84734 0.88759 0.91010
                           PC8
                                         PC10
                                                PC11
                                                         PC12
                                  PC9
                                                                 PC13
Standard deviation
                       0.69037 0.6457 0.59219 0.5421 0.51104 0.49128 0.39624
Proportion of Variance 0.01589 0.0139 0.01169 0.0098 0.00871 0.00805 0.00523
Cumulative Proportion 0.92598 0.9399 0.95157 0.9614 0.97007 0.97812 0.98335
                          PC15
                                  PC16
                                          PC17
                                                   PC18
                                                           PC19
                                                                   PC20
                                                                          PC21
Standard deviation
                       0.30681 0.28260 0.24372 0.22939 0.22244 0.17652 0.1731
Proportion of Variance 0.00314 0.00266 0.00198 0.00175 0.00165 0.00104 0.0010
Cumulative Proportion 0.98649 0.98915 0.99113 0.99288 0.99453 0.99557 0.9966
                                  PC23
                                         PC24
                                                  PC25
                                                          PC26
                          PC22
                                                                  PC27
                                                                          PC28
Standard deviation
                       0.16565 0.15602 0.1344 0.12442 0.09043 0.08307 0.03987
Proportion of Variance 0.00091 0.00081 0.0006 0.00052 0.00027 0.00023 0.00005
Cumulative Proportion
                       0.99749 0.99830 0.9989 0.99942 0.99969 0.99992 0.99997
                          PC29
                                  PC30
Standard deviation
                       0.02736 0.01153
Proportion of Variance 0.00002 0.00000
Cumulative Proportion 1.00000 1.00000
  attributes(pca)
$names
[1] "sdev"
               "rotation" "center"
                                      "scale"
$class
[1] "prcomp"
  plot(pca$x[,1], pca$x[,2], col=diagnosis)
```

```
Sca$x[, 1]
```

```
library(ggplot2)

x<- as.data.frame(pca$x)

ggplot(x)+
  aes(PC1, PC2, col=diagnosis)+
  geom_point()</pre>
```



Q4. How much varience is captured in the top 3 PCs?

They capture 76.2% of the total variance.

Q9. For the first principal component, what is the component of the loading vector (i.e. wisc.pr\$rotation[,1]) for the feature concave.points_mean? This tells us how much this original feature contributes to the first PC

```
pca$rotation["concave.points_mean",1]
```

[1] -0.2608538

attributes(pca)

```
$names
```

[1] "sdev" "rotation" "center" "scale" "x"

\$class

[1] "prcomp"

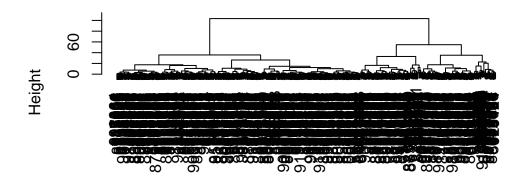
Combining PCA results with clustering

We can use our new PCA variables (i.e. the scores along the PCs contained in the pca\$x) as input for other methods such as clustering.

```
#Hclust needs a distance matrix as input
d <- dist(pca$x[,1:3])

hc <- hclust(d, method="ward.D2")
plot(hc)</pre>
```

Cluster Dendrogram



d hclust (*, "ward.D2")

To get our cluster membership vector we can use the cutree() function and specify a heigt(h) or number of groups(k).

```
grps <- cutree(hc, h=80)
table(grps)

grps
    1      2
203 366</pre>
```

I want to find out how many diagnosis "M" and "B" are in each grp?

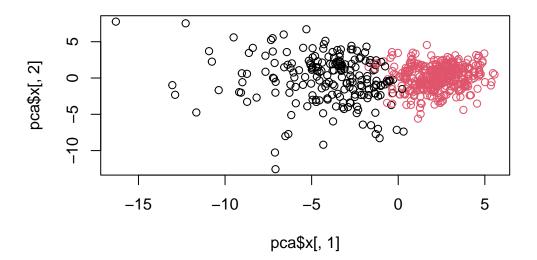
```
table(diagnosis,grps)
```

```
grps
diagnosis 1 2
B 24 333
M 179 33
```

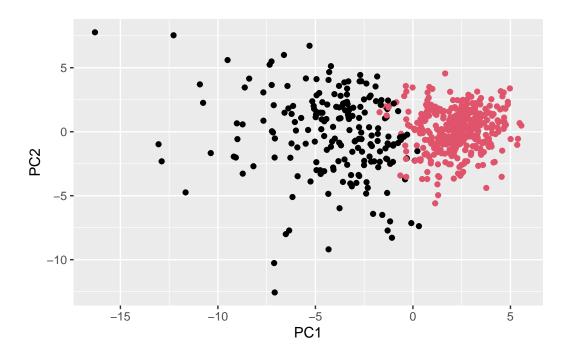
In group 1, there are 24 benign and 179 malignant cases. In group 2, 333 cases were benign and 33 were malignant.

We can also plot our results using our clustering vector grps.

```
plot(pca$x[,1], pca$x[,2], col=grps)
```



```
ggplot(x)+
  aes(PC1, PC2)+
  geom_point(col=grps)
```



Q15. What is the sensitivity and specificity of our current results?

```
table(wisc.data$diagnosis)
```

```
B M
357 212

table(diagnosis,grps)

grps
diagnosis 1 2
B 24 333
M 179 33

Sensitivity= TP/(TP+FN)

Sen<-179/(179+33)
Sen
```

[1] 0.8443396 Specificity = TN/(TN+FN) Spec <- 333/(333+33) Spec

The specificity is 90.98% and the sensitivity is 84.43%.

Prediction

[1] 0.9098361

```
url <- "new_samples.csv"</pre>
  url <- "https://tinyurl.com/new-samples-CSV"</pre>
  new <- read.csv(url)</pre>
  npc <- predict(pca, newdata=new)</pre>
  npc
           PC1
                     PC2
                                PC3
                                            PC4
                                                      PC5
                                                                  PC6
                                                                             PC7
[1,] 2.576616 -3.135913 1.3990492 -0.7631950 2.781648 -0.8150185 -0.3959098
[2,] -4.754928 -3.009033 -0.1660946 -0.6052952 -1.140698 -1.2189945
                                                                      0.8193031
            PC8
                      PC9
                                PC10
                                           PC11
                                                     PC12
                                                                PC13
                                                                         PC14
[1,] -0.2307350 0.1029569 -0.9272861 0.3411457 0.375921 0.1610764 1.187882
[2,] -0.3307423 0.5281896 -0.4855301 0.7173233 -1.185917 0.5893856 0.303029
          PC15
                     PC16
                                  PC17
                                              PC18
                                                          PC19
                                                                      PC20
[1,] 0.3216974 -0.1743616 -0.07875393 -0.11207028 -0.08802955 -0.2495216
[2,] 0.1299153 0.1448061 -0.40509706 0.06565549 0.25591230 -0.4289500
                      PC22
                                  PC23
                                             PC24
                                                         PC25
                                                                       PC26
           PC21
[1,] 0.1228233 0.09358453 0.08347651 0.1223396 0.02124121 0.078884581
[2,] -0.1224776 0.01732146 0.06316631 -0.2338618 -0.20755948 -0.009833238
             PC27
                         PC28
                                       PC29
                                                    PC30
[1,] 0.220199544 -0.02946023 -0.015620933 0.005269029
[2,] -0.001134152  0.09638361  0.002795349 -0.019015820
  plot(pca$x[,1:2], col=grps)
  points(npc[,1], npc[,2], col="blue", pch=16, cex=3)
  text(npc[,1], npc[,2], c(1,2), col="white")
```

