In [1]:
```
import pandas as pd
from sklearn.metrics import mean_squared_error

data = pd.read_csv("/Users/user/documents/Class/BMEN415/GitHub/BMEN415FinalProject/Dat

#Converted the group to dummy variables
data.loc[data.Group=='Nondemented', 'Group'] = 0
data.loc[data.Group=='Demented', 'Group'] = 1
data.loc[data.Group=='Converted', 'Group'] = 2

#Converted the M/F to dummy variables
data.loc[data.Sex=='M', 'Sex'] = 0
data.loc[data.Sex=='F', 'Sex'] = 1

#ToDo
#Fill in SES and Mini Mental State missing sample values
#Use the mean of that category for the value
# Get rid of SES data

data
```

Out[1]:

	Subject ID	MRI ID	Group	Visit	MR Delay	Sex	Hand	Age	EDUC	SES	Mini Mental State	Clinica Dementi Ratin
0	OAS2_0001	OAS2_0001_MR1	0	1	0	0	R	87	14	2.0	27.0	0.
1	OAS2_0001	OAS2_0001_MR2	0	2	457	0	R	88	14	2.0	30.0	0.
2	OAS2_0002	OAS2_0002_MR1	1	1	0	0	R	75	12	NaN	23.0	0.
3	OAS2_0002	OAS2_0002_MR2	1	2	560	0	R	76	12	NaN	28.0	0.
4	OAS2_0002	OAS2_0002_MR3	1	3	1895	0	R	80	12	NaN	22.0	0.
...	...	...	...	...	...	...	...	...	...	...	...	.
368	OAS2_0185	OAS2_0185_MR2	1	2	842	0	R	82	16	1.0	28.0	0.
369	OAS2_0185	OAS2_0185_MR3	1	3	2297	0	R	86	16	1.0	26.0	0.
370	OAS2_0186	OAS2_0186_MR1	0	1	0	1	R	61	13	2.0	30.0	0.
371	OAS2_0186	OAS2_0186_MR2	0	2	763	1	R	63	13	2.0	30.0	0.
372	OAS2_0186	OAS2_0186_MR3	0	3	1608	1	R	65	13	2.0	30.0	0.

373 rows × 15 columns

In [2]:
```
import numpy as np
from sklearn.model_selection import train_test_split
```

In [3]:
```
#PCA might be a good technique to select predictors

#note that PCA performs best when data is normalized (range b/w 0 and 1)

#It is possible to use categorical and continuous predictors
#for a regression problem. My understanding is you need to make
#dummy variables for the binary predictors.

#Variables that we will need to deal with:
# Hand, Visit, Subject ID, MRI ID
```

In [4]:
```
#Attempting PCA on data
#Hand is completely useless as it is identical for all samples
data_drop = data.drop(['Hand','Visit','Subject ID','MRI ID'], axis = 1) #axis = 1 mea

#get rid of row 360 and 359 bc they are missing alot of data (both SES and MMS)
data_drop = data_drop.drop([360, 359])


#delete all data points that dont have SES in them (this is where they have NaN)
data_drop = data_drop.dropna()

#dementia status is what we want to predict - change this to single target
group = data_drop[['Group']]


data_drop = data_drop.drop(['Group'], axis = 1) #axis = 1 means to drop column not ro
```

In [5]:
```
# give it a label type
group = group.astype('int')
```

In [6]:
```
#get a list of columns in pandas object
names_of_data = data_drop.columns.tolist()

#shuffle = false prevents data split being different everytime
X_train, X_test, y_train, y_test = train_test_split(data_drop, group, test_size=0.2, s

#split test into validate and test, again making sure the data is always the same for
X_train, X_val, y_train, y_val = train_test_split(X_train, y_train, test_size=0.05, sh

#Normalizing the data
from sklearn.preprocessing import StandardScaler
sc = StandardScaler()
X_train = sc.fit_transform(X_train)
X_test = sc.transform(X_test)

#running the actual PCA
from sklearn.decomposition import PCA

pca = PCA()
X_train = pca.fit_transform(X_train)
X_test = pca.transform(X_test)

#relief f algorithm - sorting features
```

In [7]:
```
data_drop.loc[355]
```

Out[7]:
```
MR Delay                             652
Sex                                    0
Age                                   81
EDUC                                  20
SES                                    1
Mini Mental State                     26
Clinical Dementia Rating             0.5
Estimated total Intracranial Volume 1556
Normalize Whole Brain Volume        0.691
Atlas Scaling Factor                1.128
Name: 355, dtype: object
```

In [8]:
```
data_drop.loc[354]
```

Out[8]:
```
MR Delay                             0
Sex                                  0
Age                                 79
EDUC                                20
SES                                  1
Mini Mental State                   26
Clinical Dementia Rating           0.5
Estimated total Intracranial Volume 1548
Normalize Whole Brain Volume       0.711
Atlas Scaling Factor               1.134
Name: 354, dtype: object
```

In [9]:
```
explained_variance = pca.explained_variance_ratio_
print(len(explained_variance))
print(explained_variance)
```

```
10
[0.28391839 0.21461209 0.15304569 0.1266165  0.08687837 0.05114889
 0.03240389 0.02838197 0.02191061 0.00108361]
```

From now on is Multi Layer Perceptron

In [ ]:
```
from sklearn.neural_network import MLPClassifier
MLPAlz = MLPClassifier(random_state = 1, max_iter = 100000, hidden_layer_sizes=(256,5
history = MLPAlz.fit(X_train,y_train.values.ravel())
```

In [ ]:
```
predictedMLPAlz = MLPAlz.predict(X_test)
predictedMLPAlz
```

In [ ]:
```
from sklearn.metrics import classification_report, confusion_matrix, accuracy_score
print(confusion_matrix(y_test, predictedMLPAlz))
print(classification_report(y_test, predictedMLPAlz))
results = classification_report(y_test, predictedMLPAlz)

accuracy_score(y_test, predictedMLPAlz)
```