

# MRI and Alzheimer

 Predict Dementia Using Longitudinal MRI data in Nondemented and Demented Older Adults

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### Background

### - What is Dementia?

- Dementia is a general term for a decline in mental ability severe enough to interfere with daily life (e.g. Memory loss). Dementia is not a specific disease. It's an overall term that describe a group of symptoms associated with a decline in memory or other thinking skills severe enough to reduce a person's ability to perform everyday activities.
- Alzheimer's disease accounts for 60% 80% of cases.

### Diagnose Dementia

- There is no one test to determine if someone has dementia.
- Careful medical history, physical examination, laboratory tests, and characteristics changing thinking, day-to-tay function and behavior associated with each type.
- It's hard to determine the exact type of dementia because the symptoms and brain changes of different dementias can overlap.

### Data

### **Data Features (150 subjects)**

- Subject ID
- MRI ID
- Visit
- MR Delay

#### Demographic

- M/F
- · Hand Right hand
- Age (60 98)
- EDUC (years)
- SES Socioeconomic status 1: less than high school grad., 2: high school grad., 3: some college, 4: college grad., 5: beyond college.

#### Clinical

- MMSE Mini-Mental State Examination (commonly used set of questions for screening cognitive function; 0-10 = Severe, 10-20 = Moderate; 20-25 = Mild; 25-30 = Questionably Significant)
- CDR Clinical Dementia Rating (CDR; 0 = nondemented; 0.5 = very mild dementia; 1 = mild dementia; 2 = moderate dementia) (Morris, 1993). All participants with dementia (CDR >0) were diagnosed with probable AD.

#### Derived anatomic volumes

- eTIV Estimated total intracranial volume
- nWBV Normalized whole brain volume
- ASF Atlas scaling factor

### Response to be predicted

• Group - Demented, Nondemented, Converted

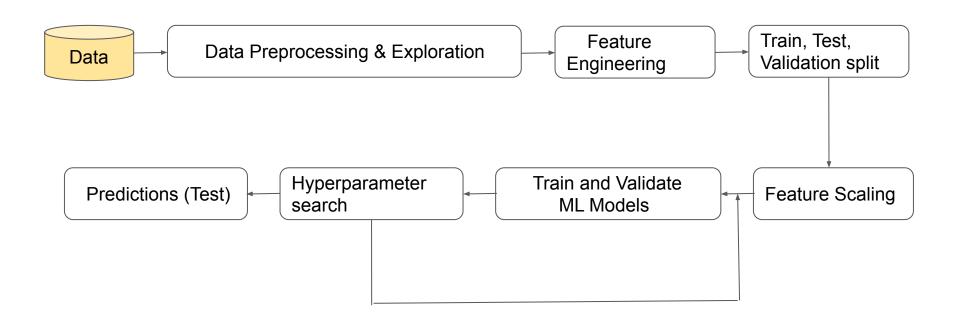
### Data (cont)

```
import numpy as np
import pandas as pd
import seaborn as sns
import matplotlib.pyplot as plt
%matplotlib inline

mri_long = pd.read_csv('oasis_longitudinal.csv', sep = ',')
mri_long.head()
```

10	Subject ID	MRI ID	Group	Visit	MR Delay	M/F	Hand	Age	EDUC	SES	MMSE	CDR	eTIV	nWBV	ASF
0	OAS2_0001	OAS2_0001_MR1	Nondemented	1	0	М	R	87	14	2.0	27.0	0.0	1987	0.696	0.883
1	OAS2_0001	OAS2_0001_MR2	Nondemented	2	457	М	R	88	14	2.0	30.0	0.0	2004	0.681	0.876
2	OAS2_0002	OAS2_0002_MR1	Demented	1	0	М	R	75	12	NaN	23.0	0.5	1678	0.736	1.046
3	OAS2_0002	OAS2_0002_MR2	Demented	2	560	М	R	76	12	NaN	28.0	0.5	1738	0.713	1.010
4	OAS2_0002	OAS2_0002_MR3	Demented	3	1895	М	R	80	12	NaN	22.0	0.5	1698	0.701	1.034

### Workflow



### Methods

### Classification

- Logistic Regression
- Decision Tree Classifier
- Random Forest
- Gradient Boosting
- Naive Bayes
- SVM
- KNN

### **Data Preprocessing**

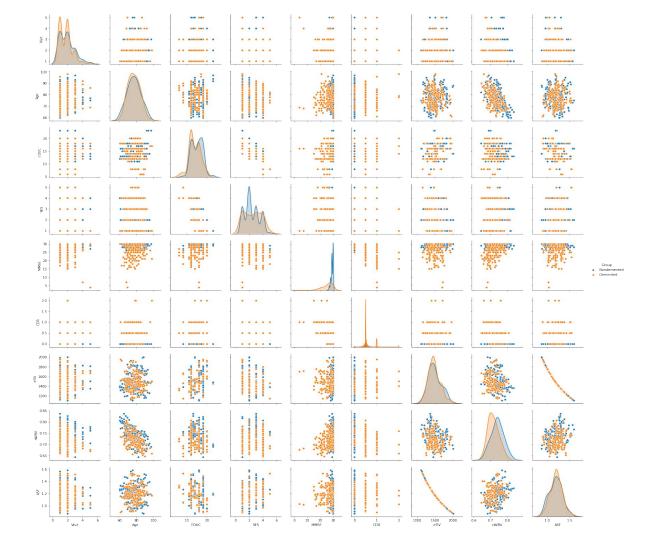
```
# Drop the trivial/unrelated predictors
df = mri long
df = df.drop(['Subject ID', 'MRI ID', 'Hand'], axis = 1)
df['Group'] = df['Group'].replace(['Converted'],['Demented'])
df['Group code'] = LabelEncoder().fit transform(df['Group'])
df['CDR code'] = LabelEncoder().fit transform(df['CDR'])
   Nondemented
                        457
                              M
                                  88
                                        14
                                            2.0
                                                  30.0
                                                        0.0 2004
                                                                  0.681 0.876
                          0
                              M
                                  75
     Demented
                                        12 NaN
                                                  23.0
                                                        0.5 1678
                                                                0.736 1.046
2
                              M
                                  76
3
     Demented
                 2
                        560
                                        12 NaN
                                                  28.0
                                                        0.5 1738
                                                                  0.713 1.010
4
     Demented
                 3
                       1895
                              М
                                  80
                                        12 NaN
                                                  22.0
                                                        0.5 1698
                                                                  0.701 1.034
(373, 12)
```

## **Data Exploration**

Age ~ -nWBV

eTIV ~ -nWBV

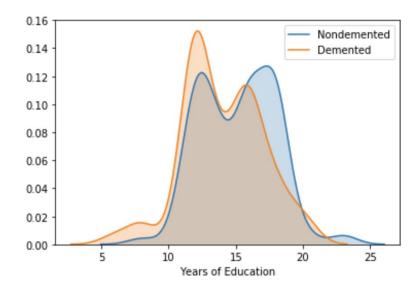
eTIV ~ -ASF (linear)



### Data Exploration (cont)

```
# Years of education
# More Demented people with less years of education
sns.kdeplot(df.EDUC[df.Group=='Nondemented'], label='Nondemented', shade=True)
sns.kdeplot(df.EDUC[df.Group=='Demented'], label='Demented', shade=True)
plt.xlabel('Years of Education')
```

Text(0.5,0,'Years of Education')



## Data Spliting and Scaling

```
from sklearn.model selection import train test split
from sklearn.preprocessing import StandardScaler
\# X, y
x cols = ['Gender code', 'Visit', 'Age', 'EDUC', 'SES', 'MMSE', 'CDR code', 'eTIV', 'nWBV', 'MR Delay']
X = df rmna[x cols]
y = df rmna['Group code'].astype('category')
# Training & Test
X_train, X_test, y_train, y_test = train test split(X, y, test size = 0.33, random state = 42)
num cols = ['Visit', 'Age', 'EDUC', 'SES', 'MMSE', 'eTIV', 'nWBV', 'MR Delay']
# normalization
# training set
X train norm = StandardScaler().fit transform(X train[num cols])
training norm col = pd.DataFrame(X train norm, index = X train[num cols].index, columns = X train[num cols].columns)
X train.update(training norm col)
# test set
X test norm = StandardScaler().fit transform(X test[num cols])
test norm col = pd.DataFrame(X test norm, index = X test[num cols].index, columns = X test[num cols].columns)
X test.update(test norm col)
```

## Modeling (cont)

# Random Forest (bagging)

```
from sklearn.ensemble import RandomForestClassifier
rf= RandomForestClassifier(random state = 123)
rf.fit(X train, y train)
y predict = rf.predict(X test)
mean squared error(y test, y predict)
se rf = np.round(mean squared error(y test, y predict),2)
acc rf = np.round(accuracy score(y test, y predict),2)
print('MSE:', se rf)
print('Accuracy:', acc rf)
fea ip = rf.feature importances
sns.barplot(x = fea ip, y = X.columns)
MSE: 0.24
Accuracy: 0.76
 from sklearn.neighbors import KNeighborsClassifier
 y predict = KNeighborsClassifier(n neighbors=5).fit(X train, y train).predict(X test)
 se knn = np.round(mean squared error(y test, y predict),2)
 acc knn = np.round(accuracy score(y test, y predict),2)
 print('MSE:', se knn)
 print('Accuracy:', acc knn)
 MSE: 0.48
 Accuracy: 0.52
 # Naive Bayes
 from sklearn.naive bayes import GaussianNB
 v predict = GaussianNB().fit(X train, v train).predict(X test)
 se nb = np.round(mean squared error(y test, y predict),2)
 acc nb = np.round(accuracy score(y test, y predict),2)
 print('MSE:', se nb)
 print('Accuracy:', acc nb)
 MSE: 0.2
 Accuracy: 0.8
```

```
# Gradient Boosting (boosting)
# sequential improvement of models by training on their errors
# improves errors, one tree each step
from sklearn.ensemble import GradientBoostingClassifier

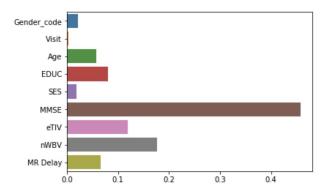
gb = GradientBoostingClassifier(random_state = 0)
gb.fit(X_train, y_train)
y_predict = gb.predict(X_test)
mean_squared_error(y_test, y_predict)

se_gb = np.round(mean_squared_error(y_test, y_predict),2)
acc_gb = np.round(accuracy_score(y_test, y_predict),2)
print('MSE:', se_gb)
print('Accuracy:', acc_gb)

fea_ip = gb.feature_importances_
sns.barplot(x = fea_ip, y = X.columns)
```

MSE: 0.18 Accuracy: 0.82

<matplotlib.axes.\_subplots.AxesSubplot at 0x1a2f3e64a8>



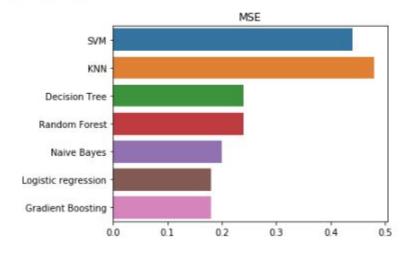
## Model comparison (MSE, Test Accuracy)

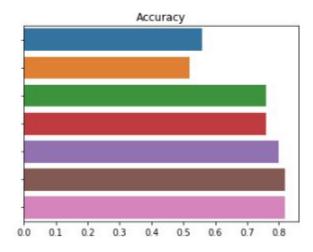
```
# visualize all methods
methods = ['SVM', 'KNN', 'Decision Tree', 'Random Forest', 'Naive Bayes', 'Logistic regression', 'Gradient Boosting']

se = [se_clf, se_knn, se_dtr, se_rf, se_nb, se_lr, se_gb]
acc = [acc_clf, acc_knn, acc_dtr, acc_rf, acc_nb, acc_lr, acc_gb]

fig, (ax1, ax2) = plt.subplots(ncols=2, figsize=(12, 4), sharey=True)
sns.barplot(x = se, y = methods, ax = ax1).set_title('MSE')
sns.barplot(x = acc, y = methods, ax = ax2).set_title('Accuracy')
```

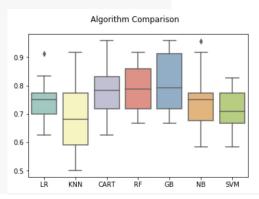
Text(0.5,1,'Accuracy')





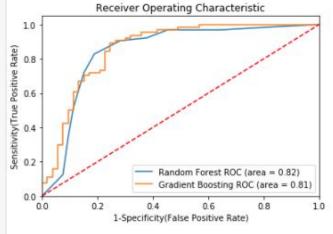
### Model comparison (CV scores)

```
# prepare models
models = []
models.append(('LR', LogisticRegression()))
models.append(('KNN', KNeighborsClassifier()))
models.append(('CART', DecisionTreeClassifier()))
models.append(('RF', RandomForestClassifier()))
models.append(('GB', GradientBoostingClassifier()))
models.append(('NB', GaussianNB()))
models.append(('SVM', SVC()))
# evaluate each model in turn
results = []
names = []
scoring = 'accuracy'
for name, model in models:
    kfold = KFold(n splits=10, random state=123)
    cv results = cross val score(model, X train, y train, cv=kfold, scoring=scoring)
    results.append(cv results)
    names.append(name)
    msg = "%s: %f (%f)" % (name, cv results.mean(), cv results.std())
    print(msg)
# boxplot algorithm comparison
fig = plt.figure()
fig.suptitle('Algorithm Comparison')
ax = fig.add subplot(111)
sns.boxplot(names, results, palette="Set3")
ax.set xticklabels(names)
```



## Performance (ROC curves)

```
# ROC curve RandomForest and GradientBoosting
import matplotlib.pyplot as plt
from sklearn.metrics import roc curve, auc, roc auc score
plt.figure()
# Add the models to the list that you want to view on the ROC plot
models = [
    'label': 'Random Forest',
    'model': RandomForestClassifier().
    'label': 'Gradient Boosting',
    'model': GradientBoostingClassifier(),
# Below for loop iterates through your models list
for m in models:
    model = m['model'] # select the model
   model.fit(X train, y train) # train the model
   y pred=model.predict(X test) # predict the test data
# Compute False postive rate, and True positive rate
    fpr, tpr, thresholds = roc curve(y test, model.predict proba(X test)[:,1])
# Calculate Area under the curve to display on the plot
    auc = roc auc score(y test, model.predict(X test))
# Now, plot the computed values
    plt.plot(fpr, tpr, label='%s ROC (area = %0.2f)' % (m['label'], auc))
# Custom settings for the plot
plt.plot([0, 1], [0, 1], 'r--')
plt.xlim([0.0, 1.0])
plt.ylim([0.0, 1.05])
plt.xlabel('1-Specificity(False Positive Rate)')
plt.ylabel('Sensitivity(True Positive Rate)')
plt.title('Receiver Operating Characteristic')
plt.legend(loc="lower right")
plt.show() # Display
```

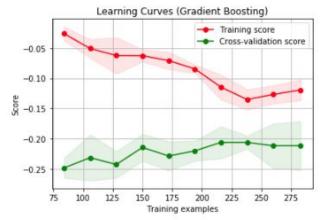


### Hyperparameter Optimization

Accuracy: 0.84

```
from sklearn.model selection import GridSearchCV
  gb = GradientBoostingClassifier(criterion='friedman mse', init=None,
                 learning rate=0.05, loss='deviance', max depth=3,
                 max features='sgrt', max leaf nodes=None,
                 min impurity decrease=0.0, min impurity split=None,
                 min samples leaf=1, min samples split=20,
                 min_weight fraction leaf=0.0, n estimators=60,
                 presort='auto', random state=10, subsample=0.8, verbose=0,
                 warm start=False)
  gb.fit(X train, y train)
  y predict = gb.predict(X test)
  mean squared error(y test, y predict)
  se gb = np.round(mean squared error(y test, y predict),2)
  acc gb = np.round(accuracy score(y test, y predict),2)
  print('MSE:', se gb)
  print('Accuracy:', acc gb)
  MSE: 0.16
```

### Not too bad



Need more data...

# THANK YOU!