## STAT 602 - Final Project

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Libraries used: MASS, foreach, doParallel, knitr, kableExtra, MNLpred, nnet, neuralnet, class, featuretoolsR (some backend python code with reticulate needs to be fixed), magrittr, matrixStats, and e1071

The data being explored in this document is from handwriting of 40 different people (subjects) writing 6 different phrases (conditions) 3 times each and in both cursive and print (group) measured with the MovAlyzeR. This document outlines some exploration to find out if there is reason to believe further work is reasonable or time-worthy (skunk work) and not a full on analysis itself. Accuracies were measured by dividing the number of correct predictions by the total number of predictions. Here, predictions are important and will be used as the metric for model performance.

Initially, the data frame included all of the different "Segments" broken down by the direction of the writing, but the mean of all fo the different trials (each combination of subject + condition + group + replication) was taken as a baseline for predictions. With the new summary data where each row is a unique replication. These values were than checked for values that are only zero's and removed. These variables are RelativeDurationofPrimary, and RelativeSizeofPrimary.

Since from the in class 'Playing with Data' we got some benchmark accuracies from linear discriminate analysis and quadratic discriminate analysis using the principal components of the remaining variables for the different response variables group, subject, condition, and joint which is the combination of all 3. It's important to note that the model for the joint variables doesn't seem to work for the quadratic discriminate analysis

So these are the accuracies to beat through other methods or further exploration. The next thing that I tried to do was remove other variables that didn't appear to show separation in the box plots that were used in the 'Playing with Data' code. This was only for group separation, however. The columns that visually appeared to have no separation were PeakVerticalVelocity, HorizontalSize, AverageNormalizedyJerkPerTrial, AbsoluteJerk, AverageNormalizedJerkPerTrial, NumberOfPeakAccelerationPoints, and AveragePenPressure. This resulted in a higher increase in accuracies for group and condition but not for the subject or joint response variable.

With slight improvements in group and condition but not subject or joint, I thought another place to look would be using the means themselves in the LDA instead of the principal components, as well as scaling the means. Using the mean values from the variables with visible group separation increased the prediction accuracy of group, subject, condition, and joint over the use of principal components for the LDA, but the QDA decreased group and subject accuracies, while slightly boosting condition. Adding back in the variables that appeared to have some separation in the boxplots split by group increased all 4 response variables when using the LDA and subject and condition when using QDA. Quadratic discriminate analysis again had shown a decrease in the prediction accuracy for group when using the means from the data frame used in class like seen with the means of those variables with some visible separation between the 2 group factors - CUR and PRI. Finally, for LDA and QDA I scaled the means of the data used in class which had minor accuracy increases over the raw means for the LDA's group and condition (subject and joint that was the same) where as the QDA didn't change from the unscaled data.

After working with the LDA and QDA above with the column means of the samples, I went with the

suggestion given in lecture about logistic regressions. this didn't work for the joint prediction because there were 'too many (12000) weights'. I also did a single non-cross validated approach just to get some kind of an idea and then followed it with a cross validation for better accuracy judgement. The prediction accuracy of group and subject did increase (the logistic regression without CV was just to see if it was worth looking further into but not itself very trustworthy), and the condition accuracy did show an increase but it wasn't as high as the others (condition accuracy ended up at only 0.425 while group and subject were 0.928 and 0.725, respectively). All values were greater than the baseline set in lecture so the logistic regression shows promise.

Single regular glm was giving errors, I found the multinom logistic regression function which can be used for predicting probabilities of classes for the larger response sizes rather than the response like glm. https://www.r-bloggers.com/how-to-multinomial-regression-models-in-r/

Another avenue that I have used for classifications in the past was using neural networks. The nerualnet package makes this relatively easy without having to code perceptrons from scratch (actually taught how to do that on codecademy under the data science path for python) along with a bunch of customizations. Looking at some of the recommendations for how big to make the hidden layer, a stack overflow poster had said that good guidelines are 2/3 the input + the output, somewhere between the input shape and output shape, or no more than twice the input. I chose to go with 15 as a round number that was close to the 2/3 input + output for group and maintained it for the others. The input data was also normalized (have read some other places this is good practice so it doesn't put inappropriate weights in certain variables and they all start on a "level" playing field). The accuracy of the neural net is surprisingly well for the group and subject (since the condition failed to converge) with 0.931 and 0.621, respectively. This is better than the baselines, however with condition and joint not converging it doesn't warrant further investigation at this time. With different parameteters this may be possible. https://stats.stackexchange.com/questions/181/how-to-choose-the-number-of-hidden-layers-and-nodes-in-a-feedforward-neural-netw

Following the logistic regression, I thought maybe looking at a KNN set of models would provide a good way to classify. For KNN, I used the means data frame that was constructed from the 'Playing with data' along with the scaled verson of the means using scale. The unscaled accuracy with K=1 was lower for all response combinations than was the scaled data. The best joint prediction came from K=1 for the scaled data where the model achieved an accuracy of 0.342. Other combinations included group: subject where the highest accuracy was 0.609 for K=1, group:condition with a best accuracy of 0.465 with K=1, and subject:condition with a best accuracy of 0.346 on the scaled data with K=1. The scaled data also performed better with just the single response variables, and with K=1. Response group was 0.935, subject was 0.624, and condition was 0.481.

Table 1: KNN accuracies of K=1 and K=2 for both scaled and unscaled means data.

	K=1_Unscaled	K=2_Unscaled	K=1_Scaled	K=2_Scaled
knn.g.s.c.acc	0.1104167	0.0895833	0.3423611	0.2916667
knn.g.s.acc	0.3034722	0.2423611	0.6090278	0.5451389
knn.g.c.acc	0.2000000	0.1694444	0.4645833	0.4020833
knn.s.c.acc	0.1159722	0.0979167	0.3458333	0.2888889
knn.g.acc	0.7326389	0.7159722	0.9347222	0.9236111
knn.s.acc	0.3388889	0.3034722	0.6236111	0.5513889
knn.c.acc	0.2458333	0.2284722	0.4805556	0.4006944

When all of the previous methods weren't able to get the condition accuracy up higher than 0.425 (LOOCV logistic regression), I thought it would be good to go back to the beginning and push further with descriptive statistics than what was done in lecture. At first I tried to use a package called featuretoolsR, but there is something that doesn't work right with the backend and reticulate to port into python (https://github.com/magnusfurugard/featuretoolsR). Here, I used the matrixStats package to compute MovAlyzR variable summaries by trial for mean, median, variance, standard deviation, interquartile range,

range, min, and max. After computing that for each of the output variables, the data frame is now 200 predictors, but there are some that have no real use and can be removed by checking if the variance is zero and if so, remove it. The remaining table has 178 predictors (22 columns had a variance of 0 and therefore no difference between all of the samples [rows]) were used for pca and the different components were looped through to determine the best number for the particular response. Using these metrics as predictors for group, subject, condition, and joint proved fairly effective with accuracies of 0.9597, 0.86944, 0.7201, and 0.7076, respectively. So it looks as though an LDA taking more information about the original data is more powerful than just means.

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Figure 1: Principal components of the descriptive statistics values for the 23 columns used in lecture (mean, median, variance, standard deviation, interquartile range, range, min, and max).

On top of just using the values mentioned above, I used the e1071 package to calculate the column skewness, and kurtosis, while adding median absolute deviation from the matrixStats package. After computing these for each column, there were now 34 columns that were removed for having a variance of 0. Adding these extra descriptors increased the accuracy of the three response variables independently, but not the joint response which decreased from 0.7076 to 0.6813, which is higher than the baseline from lecture. With these descriptive data, I attempted XGBoost with high number of K-fold (attempt to get close to LOOCV) but is computationally expensive for only about 30% at best prediction accuracy.

Predictions for the unlabeled data - Using the first descriptive statistics LDA model, the prediction accuracy I would expect to be similar to the test since that was from LOOCV. To be conservative, I would expect predictions for group to be around 90%, subject around 80%, while condition and joint accuracy around 70%. To make the predictions, the testing data will be used in two steps with predict(); first with the prinipal component models, then with the LDA models to arrive at the final predicted classes.

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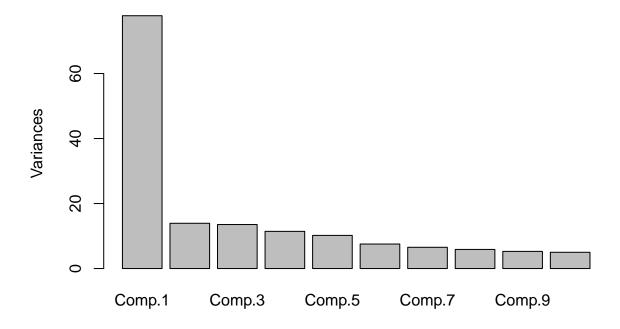


Figure 2: Principal components of the descriptive statistics values for the 23 columns used in lecture (mean, median, variance, standard deviation, interquartile range, range, min, max, skewness, kurtosis and median absolute deviation).

Table 2:

	Group	Subject	Condition	Joint
LDA PCA (cps baseline)	0.8798611	0.3854167	0.3034722	0.2861111
QDA PCA (cps baseline)	0.8256944	0.5444444	0.2638889	NA
LDA PCA vars with separation	0.8951389	0.2895833	0.3263889	0.2347222
QDA PCA vars with separation	0.8687500	0.3986111	0.2743056	NA
LDA on means with separation	0.9145833	0.3979167	0.3722222	0.3756944
QDA on means with separation	0.6979167	0.4736111	0.2847222	NA
LDA on class means	0.9222222	0.7000000	0.4048611	0.5701389
QDA on class means	0.6881944	0.6583333	0.3819444	NA
LDA on scaled class means	0.9229167	0.7000000	0.4055556	0.5701389
QDA on scaled class means	0.6881944	0.6583333	0.3819444	NA
One shot Logistic Regression (no CV) scaled class	0.9326389	0.9430556	0.455556	NA
Logistic Regression LOOCV scaled class	0.9277778	0.7527778	0.4250000	NA
Neural net LOOCV (1/25)	0.9310345	0.6206897	NA	NA
LDA Descriptive Stats - 1	0.9597222	0.8694444	0.7201389	0.7076389
LDA Descriptive Stats - 2	0.9604167	0.8770833	0.7500000	0.6812500