



Final Challenge: Alzheimer Patient Classification

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DATA ANALYSIS

The first step of the challenge is to analyze the datasets. The results are briefly summarized as follows:

- **Dimensionality**: We inspect **p** the number of predictors and **n** the number of samples in each dataset:
 - Task 1: n = 164, $p = 429 \Rightarrow very high dimensionality <math>(p >>> n)$
 - Task 2: n = 172, $p = 63 \Rightarrow Low$ dimensionality (p < n)
 - Task 3: n = 172, p = 593 => very high dimensionality <math>(p >>> n)
- *Balance*: we compare the number of samples in each class:
 - Task 1: 81 AD vs 83 CTL => Balanced dataset.
 - Task 2: 82 AD vs 90 MCI => Balanced dataset.
 - Task 3: 82 CTL vs 90 MCI => Balanced dataset.
- Correlation: we calculate the correlations between each pair of predictors, and we draw the corresponding correlation matrices (Figure 1). We notice the presence of high correlations between some pairs of the variables (dark red) in all tasks.
- *Range*: by printing the summary of each dataset, we notice a difference in the range of variables. E.g., *E2F2* ∈ [8, 13] whereas background ∈ [0.003, 0.007].
- Outliers: the package "rrcovHD" is used to detect outliers based on the result of robust PCA. The results show the existence of 5 outliers in task 1, 6 in task 2 and 4 in task 3. However, it is seen that the outliers in tasks 1 and 3 are farther in distance from the other points compared to the outliers in task 2.

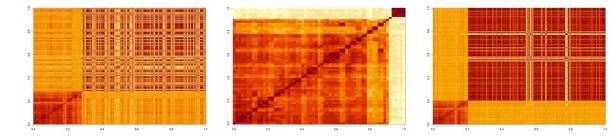


Fig 1: Correlation matrices of task 1, 2 and 3 respectively

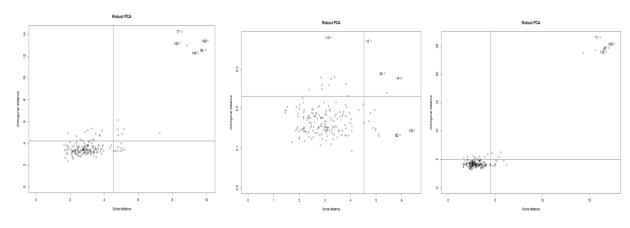
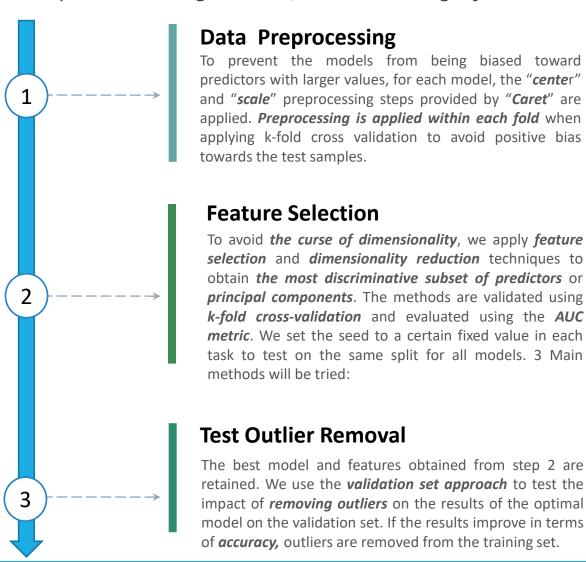


Fig 2: Robus PCA results of task 1, 2 and 3 respectively

METHODOLOGY

In order to solve the classification problem, for each task we will try *different classification algorithms* and use the one that provides the *highest AUC/ROC score using k-fold Cross-Validation*. The following pipeline is applied for each model:



Recursive Feature Elimination

The rfe function of "Caret" is used to perform recursive feature elimination with multiple subset sizes. 5-fold cross validation is used to validate the results. If the model has parameters, grid search is used to test a few values for each subset of features. The subset suggested by RFE is then compared with the other feature selection methods (below) using the same sampling splits (by cross-validation).

Removing Correlated Predictors

A *correlation filter* is used to remove predictors with *correlation higher than a threshold* with any of the other predictors. *Multiple thresholds* (0.6, 0.7, 0.8) are tested for each classifier. If the model has parameters, *grid search* is used to fine-tune the parameters for each correlation filter and the parameter that gives the *highest AUC* is retained.

Principal Component Analysis

PCA is applied with *multiple thresholds* (cutoffs for the cumulative percent of variance to be retained). If the model has parameters, *grid search* is used to fine-tune the parameters and find the *best ones for each PCA threshold*. *K-fold Cross-validation* is used to validate the results using the *AUC metric*.

Example: Task 1 AD vs CTL

To have a closer look at the methodology we will take Task 1 as an example. Similar analysis is performed for the other tasks.

- In task 1, we need to classify patients to 2 classes:
 - AD: Alzheimer Disease
 - o CTL: Control
- To achieve this, we will train the following models:
 - Logistic Regression
 - Linear Discriminant Analysis
 - Quadratic Discriminant Analysis
 - K-Nearest Neighbors
 - Support Vector Machine with Linear kernel
 - Support Vector Machine with Radial Basis Function kernel
 - Random Forest.
- We will take k-NN as an example for demonstration. The same analysis is applied to the other models.
 - First, RFE is applied with subsets of sizes 1, 5, 10, 25, 50, 100 and 250. The results are validated using 5-fold repeated cross validation.
 - The result shows that RFE suggests using **10 predictors**. The results of RFE are summarized in Figure 3. The variable importance is shown in Figure 4.
 - The resulting 10 predictors are saved to be tested later with the same 10-fold cross-validation split with the other k-NN models.

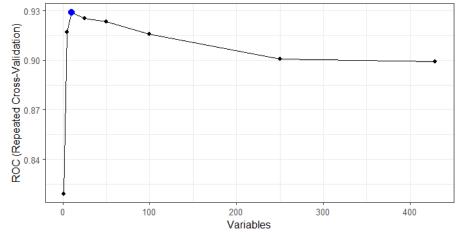


Figure 3: k-NN RFE results

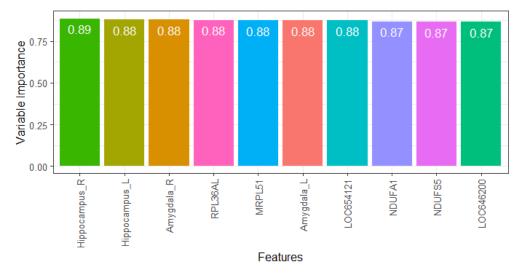


Figure 4: k-NN variable importance

Example: Task 1 (cont'd)

- Next, "recipes" library is used to create a grid of models with different preprocessing steps. We will add the following models to the grid:
 - A baseline with no feature selection (it uses all predictors).
 - A model that uses the predictors suggested by **RFE** (previous slide).
 - Models that use correlation filters with different thresholds (0.6, 0.7, 0.8).
 - Models that apply PCA with different thresholds (0.75, 0.8, 0.85, 0.9, 0.95).
- The models in the grid are then trained using 10-fold repeated cross validation with 5 repeats. A grid search with length 10 is applied to find the optimal parameter k for each model. In all experiments, the data is preprocessed using "scale" and "center" within each fold according to the formula:

$$x'_{ij} = \frac{x_{ij} - \bar{x}_j}{\sigma_j}$$

Where x' is the new value of the data, i is the sample index, j is the predictor index

- The models are compared with respect to the area under the ROC.
- Figure 5 summarizes the results in term of AUC/ROC.
- Table 1 shows the optimal number of neighbors k found by the grid search for each model. As well as the number of predictors/principal components.
- From the results, it is seen that the k-NN model that gives the best median Area Under the ROC is the one that uses the *predictors suggested by RFE* and a k number of neighbors equal to 21. The model achieves AUC/ROC = 0.9531250 and MCC = 0.7745967.
- This model is then saved to be compared with the other classifiers.

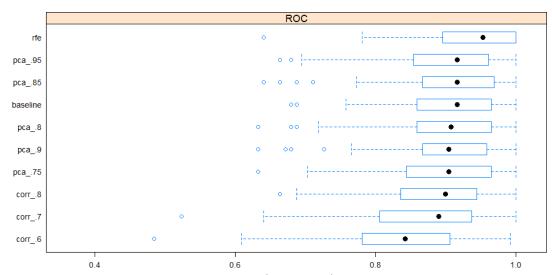


Figure 5: k-NN results

Model	Optimal k	# Variables
Baseline	23	429
RFE	21	10
Correlation Filter 0.6	23	56
Correlation Filter 0.7	13	102
Correlation Filter 0.8	21	188
PCA 0.75	11	9
PCA 0.8	13	15
PCA 0.85	11	25
PCA 0.9	15	41
PCA 0.95	15	70

Table 1: k-NN optimal k parameters and variable numbers

Example: Task 1 (cont'd)

The final analysis results of all models trained on Task 1's dataset are summarized in Table 2.

Model	The best Feature selection method	The number of predictors/principal components used	Optimal model parameters	Area under the ROC	мсс
Logistic Regression	PCA with threshold = 0.8	15	No parameters	0.9531250	0.7638889
Linear Discriminant Analysis	PCA with threshold = 0.9	41	No parameters	0.9557292	0.7692428
Quadratic Discriminant Analysis	PCA with threshold = 0.75	9	No parameters	0.9218750	0.7692428
K-Nearest Neighbors	Recursive Feature Elimination	10	k = 21	0.9531250	0.7745967
SVM with Linear Kernel	No Feature Selection (baseline)	429	C = 1	0.9722222	0.7745967
SVM with Radial Basis Function Kernel	No Feature Selection (baseline)	429	Sigma = 0.001794168 , C = 2	0.9583333	0.7638889
Random Forest	Recursive Feature Elimination	10	mtry = 2	0.9414062	0.7789731

- It is seen from the results that the model that achieved the best Area under the ROC is **Support Vector Machine** with a **linear kernel** that uses **all the predictors** in the dataset.
- We can notice from the results that, unlike SVM, simpler methods such us **k-NN** and **Logistic regression** work better with a **small number of predictors**. In the case of these models, the least complex models gave the best results. It can also be seen that **PCA** and **RFE** perform better than removing correlated predictors for feature selection and dimensionality reduction.
- Finally, We split the dataset to 150 training samples and 14 samples for validation. We test the *accuracy* of the best model (linear SVM) using *all* the samples in the training set from one hand and after removing outliers from another. The results show that in Task 1, removing outliers does not change the results of the model. Therefore, they will not be removed.
- The best model, SVM with Linear kernel is then retrained using the whole training set and used to predict the classes for the test set.

RESULTS FOR TASK 2 AND TASK 3

Task	Model	The best Feature selection method	The number of predictors	Optimal model parameters	AUC/ROC	MCC
-2-	Logistic Regression	PCA with threshold = 0.85	12	None	0.8055556	0.4166667
AD	Linear Discriminant Analysis	PCA with threshold = 0.90	18	None	0.8040123	0.4084912
VS	Quadratic Discriminant Analysis	PCA with threshold = 0.75	6	None	0.7638889	0.4260064
	K-Nearest Neighbors	No feature selection (Baseline)	63	K=15	0.7916667	0.5093840
MCI	SVM with Linear Kernel	PCA with threshold = 0.85	12	C = 1	0.8055556	0.4366100
	SVM with RBF Kernel	PCA with threshold = 0.85	12	Sigma = 0.05678335, C = 0.25	0.8194444	0.4166667
	Random Forest	Correlation Filter with th = 0.80	28	mtry = 19	0.7847222	0.4084912
-3-	Logistic Regression	PCA with threshold = 0.85	16	None	0.888889	0.6017536
MCI	Linear Discriminant Analysis	Recursive Feature Elimination	1	None	0.8750000	0.5493503
	Quadratic Discriminant Analysis	PCA with threshold = 0.85	16	None	0.8472222	0.5277778
VS	K-Nearest Neighbors	Recursive Feature Elimination	5	k = 9	0.8726852	0.5493503
CTL	SVM with Linear Kernel	PCA with threshold = 0.85	16	C = 1	0.8750000	0.6042610
	SVM with RBF Kernel	Recursive Feature Elimination	250	Sigma = 0.005432391 C = 4	0.9027778	06527778
	Random Forest	Recursive Feature Elimination	10	mtry = 5	0.8750000	0.5493503

- We can see from the table that for both tasks **SVM with RBF kernel** gave the best result in terms of AUC/ROC. This model is used to obtain the results for the test set.
- We can see that in Task 2 the MCC is low which reflects the difficulty of separating the two classes AD and MCI and thus deciding whether a patient has Alzheimer's disease, or a mild cognitive impairment based on the predictors provided in the dataset.
- We can also see that in task 2, the curse of dimensionality is less problematic than the other tasks. This is seen in the result of k-NN that usually suffers from this problem, but it performed the best without feature selection in this task.

Thank you!