## MA679 Midterm Exam

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### Context

A company wants to hire data scientists from pool of people enrolled in the courses conducted by the company. The company wants to know which of these candidates are looking to change their job. Information related to demographics, education, experience are in hands from candidates signup and enrollment. In this exam, your goal is to predict if the candidate is looking for a new job or will work for the current company.

- uid: Unique ID for candidate
- · city: City code
- city\_dev\_index : Development index of the city (scaled)
- gender: Gender of candidate
- relevant\_experience: Relevant experience of candidate
- enrolled university: Type of University course enrolled if any
- education level: Education level of candidate
- major\_discipline :Education major discipline of candidate
- experience years: Candidate total experience in years
- company\_size: No of employees in current employer's company
- company\_type : Type of current employer
- lastnewjob: Difference in years between previous job and current job
- training hours: training hours completed
- change job: 0 Not looking for job change, 1 Looking for a job change

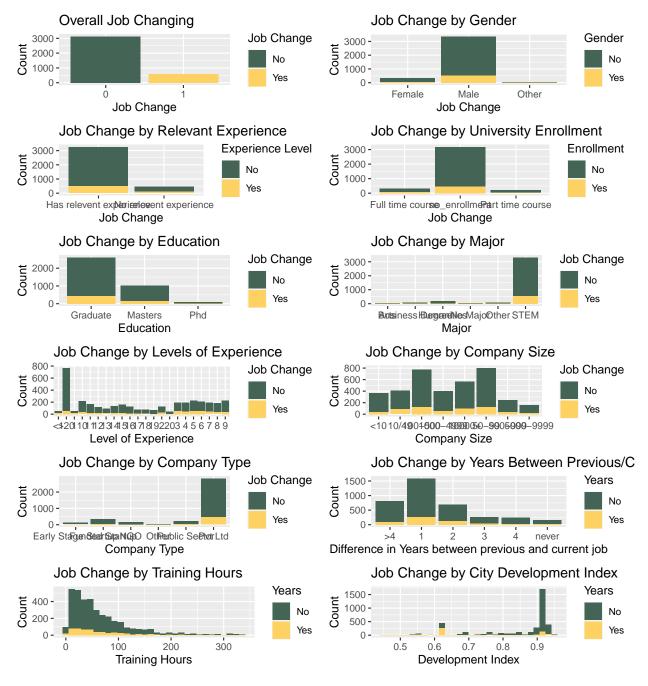
#### Introduction

Understanding that this is a classification problem given a set of 10 factored and one continuous variable - not considering the candidate ID and city code (I figured city code is included within the continuous City Development Index), a classification tree is likely to be the best approach. In using a classification tree, we determine how "well" the model preforms via the error rate and accuracy (James et al, 311). Additionally, there are some advantages to using trees for clarification purposes like being easily interpretable, similar/mirroring human decision making and easily handle qualitative variables (315). There are certain drawbacks to using trees as they can be very non robust and perform poorly when applied to other data sets which can be handled through boosting among other methods.

The approach I've taken is to use a gradient boosting machine (GBM). Boosting works through growing trees sequentially, meaning that each tree is grown from the previous one (321). This helps in the classification by fitting continuously smaller trees to the previously made residuals. The problem and challenge with using a GBM in this approach is the missing values in this data set. Out of the 8000 training values, there are 4,282 incomplete rows ecompassing about 54% of the data!

## Data EDA/Visualization

Unfortunately, it appears that our data is very uneven once we remove NANs rows! Additionally, it appears when grouping by certain variables, we still have a large amount unevenness in job change, with a significantly higher rate of "No" responses. Because of this uneveness-way more people declining jobs compared to accepting them, I would assume more problems in the tree with model specificity when we approach the modeling process.

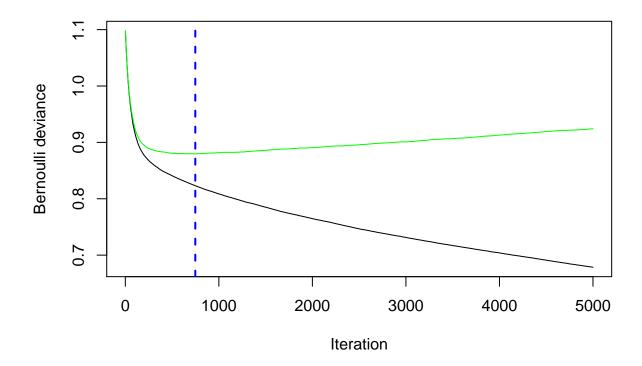


### Dealing with NANs

Because of the unevenness in the data and the large amount of missing values, it is likely beneficial to include them somehow. We have several methods for approaching the NA problem; We could "input" missing values simply as the overall mean of the predictor - this would somewhat balance our data towards the mean, create a dummy variable for "missing" and feed it into the model, or create surrogate variables (Hastie et al, 311). I decided to fabricate the variables from the mean of the overall predictor set as a method to balance the observations and allow parsing to library(gbm). The new generated data EDA shows in Appendix A.

## **Model Selection**

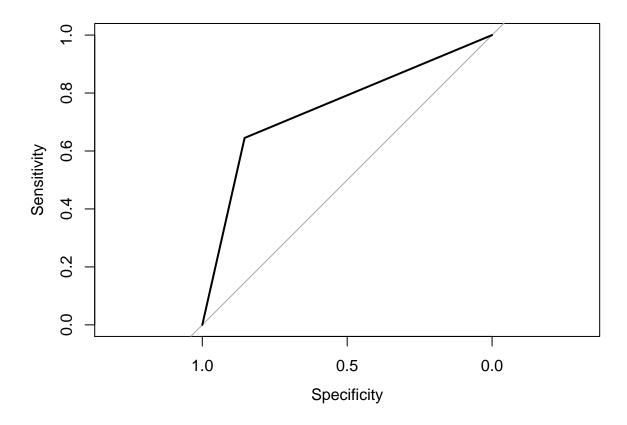
Originally, I was considering filling the NA values with a  $N-(\mu,\sigma^2)$  to add some variation and minimize drawing predictors to their mean; however, I turned to just the mean-fill method to avoid adding further noise. Using the GBM Classification Tree as described above, I fit a model of 5000 trees which, as expected, turned out to be too many (Seen Below)! To deal with this and avoid overfitting to the training set, I used gbm::gbm.pref to return the estimated optimal number of iterations using the cv method (748) - found by determining at what iteration count the trees have the lowest MSE compared to the training error; below we can see the error on test (green line) and the error on train (black line) over increasing trees. The relative influence can be found in Appendix B.



# Model Validation / Evaluation

To validate the model, I split the given train into half and trained the model on the first half, then tested it on the second to generated a confusion matrix (Appendix C). Overall the model preformed well, with an accuracy of 80.25% - 95% confidence intervals being (78.98% and 81.47%) and test MSE of 3.92. Furthermore, as expected, the sensitivity preformed well, and the specificity of the model preformed poorly. For further validation, we can look at the ROC curve (Below) and test the AUC (.7495). As a part of evaluation, I also wanted to test if the model would have preformed better if I had dropped the NA values entirely. The NA dropped model (Appendix D) drew roughly the same variable relative influences, but had worse accuracy,

specificity and MSE when applied to the held test set.



## Discussion

As previously mentioned this model has the shortfall of not being able to intake the 54% incomplete rows in the data and instead intakes the mean value of the variable, which artificially stabilizes the data around the mean. Furthermore, the selected model has a difficult time classifying false positives, as conjecture, this is because of the overall imbalance in the predicted variable. However, the model preformed much better than the NA removed model in overall accuracy and specificity/sensitivity. Special thanks to Justin Singh-M from UC Santa Barbara (https://gist.github.com/program--) for the visualization of tree function!



#### References

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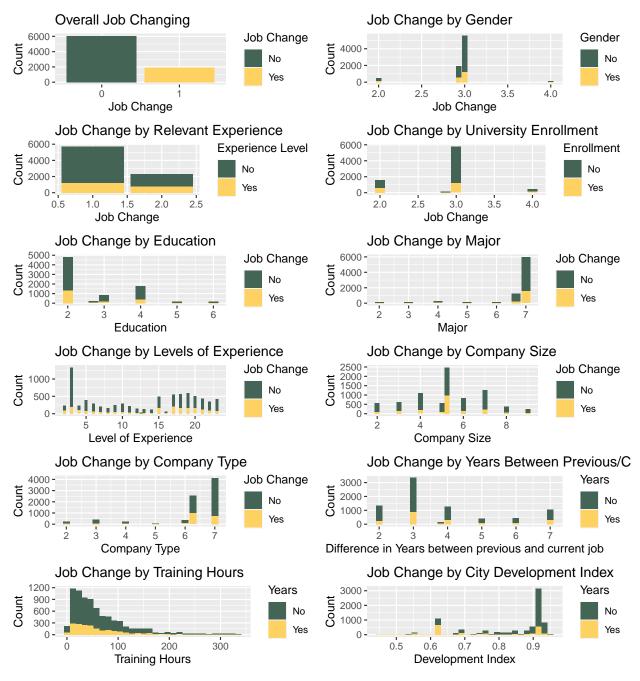
Xavier Robin, Natacha Turck, Alexandre Hainard, Natalia Tiberti, Frédérique Lisacek, Jean-

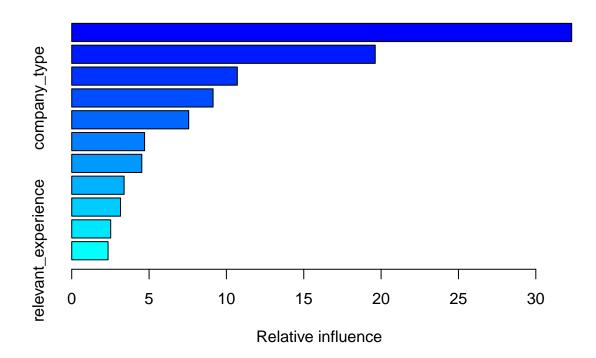
Charles Sanchez and Markus Müller (2011). pROC: an open-source package for R and S+ to analyze and compare ROC curves. BMC Bioinformatics, 12, p. 77. DOI: 10.1186/1471-2105-12-77 http://www.biomedcentral.com/1471-2105/12/77/

Wickham et al., (2019). Welcome to the tidyverse. Journal of Open Source Software, 4(43), 1686, https://doi.org/10.21105/joss.01686

## **Appendix**

#### $\mathbf{A}$





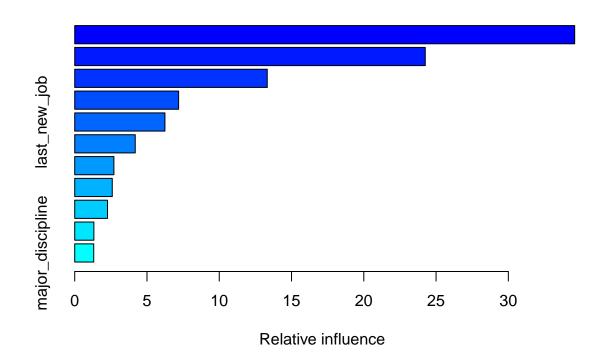
```
##
                                       var
                                             rel.inf
## city_dev_index
                            city_dev_index 32.319667
## training_hours
                            training_hours 19.620727
## experience_years
                          experience_years 10.706254
## company_type
                              company_type 9.137948
## company_size
                              company_size 7.562379
## education_level
                           education_level
                                            4.710258
## last_new_job
                              last_new_job 4.530599
## major_discipline
                          major_discipline
                                            3.386707
## enrolled_university enrolled_university
                                            3.156699
## gender
                                    gender
                                            2.519015
## relevant_experience relevant_experience 2.349747
```

#### $\mathbf{C}$

```
## Confusion Matrix and Statistics
##
##
             Reference
## Prediction
                 0
            0 2574
                    350
##
##
               440
                    636
##
##
                  Accuracy : 0.8025
##
                    95% CI: (0.7898, 0.8147)
##
       No Information Rate: 0.7535
```

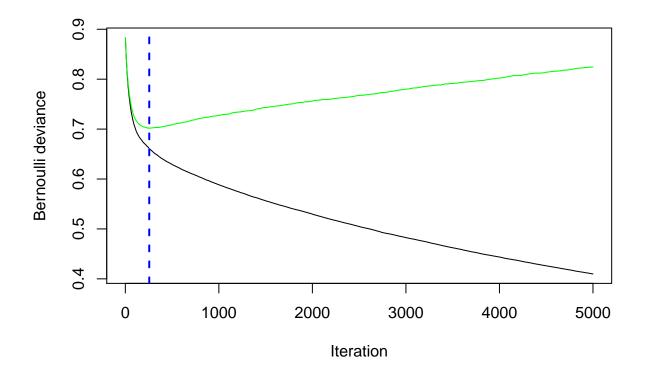
```
##
       P-Value [Acc > NIR] : 1.013e-13
##
                     Kappa : 0.4842
##
##
    Mcnemar's Test P-Value : 0.001543
##
##
               Sensitivity: 0.8540
##
               Specificity: 0.6450
##
##
            Pos Pred Value : 0.8803
##
            Neg Pred Value : 0.5911
##
                Prevalence: 0.7535
            Detection Rate: 0.6435
##
##
      Detection Prevalence : 0.7310
         Balanced Accuracy : 0.7495
##
##
          'Positive' Class : 0
##
##
```

 $\mathbf{D}$ 



	var	rel.inf
city_dev_index	city_dev_index	34.590496
training_hours	training_hours	24.253311
experience_years	experience_years	13.320469
company_size	company_size	7.188066
<pre>last_new_job</pre>	<pre>last_new_job</pre>	6.240360
	<pre>city_dev_index training_hours experience_years company_size last_new_job</pre>	city_dev_index city_dev_index training_hours training_hours experience_years company_size company_size

```
## company_type company_type 4.187622
## education_level education_level 2.710540
## enrolled_university enrolled_university 2.598123
## relevant_experience relevant_experience 2.269638
## gender gender 1.325470
## major_discipline major_discipline 1.315906
## [1] 8.232002
```



```
## Confusion Matrix and Statistics
##
##
             Reference
## Prediction
                 0
                      1
##
            0 1454 131
##
            1 116 158
##
##
                  Accuracy : 0.8671
                    95% CI : (0.8509, 0.8822)
##
##
       No Information Rate: 0.8445
##
       P-Value [Acc > NIR] : 0.003423
##
##
                     Kappa : 0.4831
##
##
    Mcnemar's Test P-Value: 0.373037
##
##
               Sensitivity: 0.9261
               Specificity: 0.5467
##
```

```
Pos Pred Value : 0.9174
##
##
           Neg Pred Value : 0.5766
               Prevalence: 0.8445
##
##
           Detection Rate : 0.7821
     Detection Prevalence : 0.8526
##
##
         Balanced Accuracy: 0.7364
##
##
          'Positive' Class : 0
##
```