Disputed Workers' Compensation Claims in New York State

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Background:

- Workers' compensation claims sent to the New York State Workers' Compensation Board (WCB)
- About 5% of claims have are disputed (controverted) by insurance companies
 - Unbalanced data set
- Some information we have:
 - Part of body injured, type of injury, cause of injury
 - Geographical region
 - Weekly salary of employee
 - Dates various forms are recieved by WCB

Objective:

- Predict which Workers' Compensation Board claims were controverted by the insurance companies.
- Ultimately could help determine which claims should have more resources allocated to them upfront.

Data Sources and References:

NYS Workers' Compensation Board Data - Over 1 million data points:

https://www.kaggle.com/new-york-state/nys-assembled-workers'-compensation-claims/

https://www.kaggle.com/new-york-state/nys-assembled-workers'-compensation-claims/

https://data.ny.gov/Government-Finance/Assembled-Workers-Compensation-Claims-Beginning-20/jshw-gkgu

https

OIICS Codes (appears version 1.01 used): https://wwwn.cdc.gov/wisards/oiics/Trees/MultiTree.aspx?Year=2007)

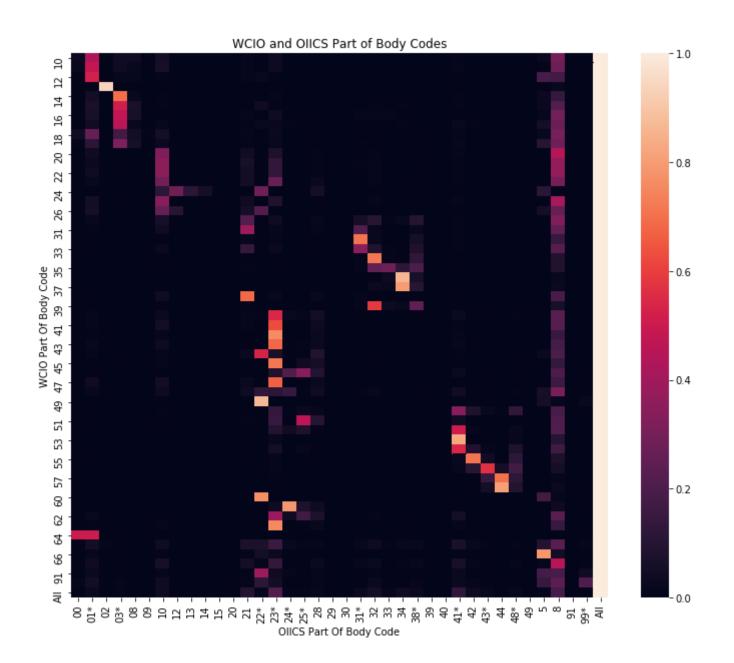
Claims Process:

http://www.wcb.ny.gov/content/main/onthejob/HowSystemWorks.jsp (http://www.wcb.ny.gov/content/main/onthejob/HowSystemWorks.jsp)

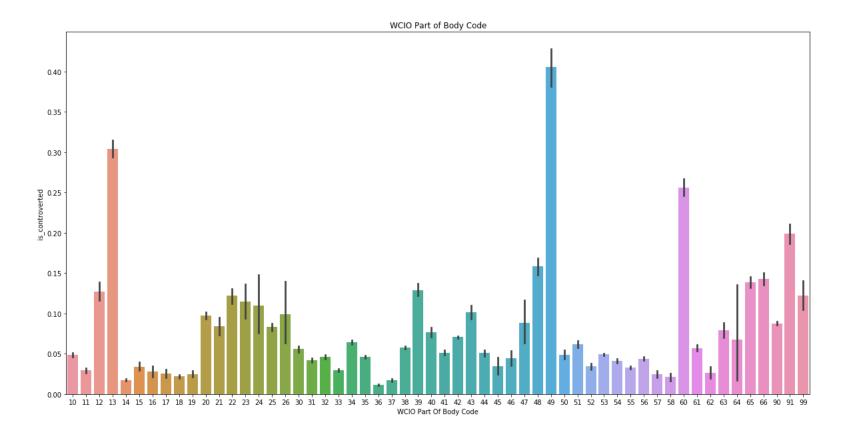
New York State Regions - Combine counties into regions: https://en.wikipedia.org/wiki/Category:Regions of New York (state))

Exploratory Data Analysis

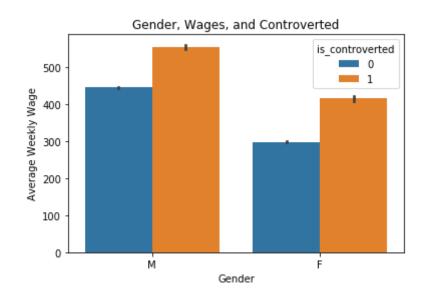
- Two different sets of injury codes WCIO and OIICS
- Cause of injury correlated with type of injury? Redundant features?



In [13]: plt.figure(figsize=(20, 10))
 sns.barplot(x='WCIO Part Of Body Code', y='is_controverted', data=df)
 plt.title('WCIO Part of Body Code')
 plt.show()



```
In [18]: sns.barplot(x='Gender', y='Average Weekly Wage', hue='is_controverted', data=df)
    plt.title('Gender, Wages, and Controverted')
    plt.show()
```



Modeling - Selected Features

- WCIO codes: more fields filled out
- Omitted age: too many NaNs
- Used accident and assembly dates:
 - Not time to ANCR (difference between those dates)
 - Time of year more important than difference between dates
- Gender
- Region
- Type of insurance
 - Not insurance carrier too many unique carriers

Model 1: Random Forest

- Fits a number of decision tree classifiers on sub-samples of the dataset.
- Less prone to overfitting than a decision tree
- Label encoded data set
- Look at feature importances
- For handling the unbalanced dataset, try regular, under sampled, and SMOTE
 - Regular sampling good accuracy, poor recall
 - Under sampled and SMOTE, overfit to data, high recall, low precision

Results

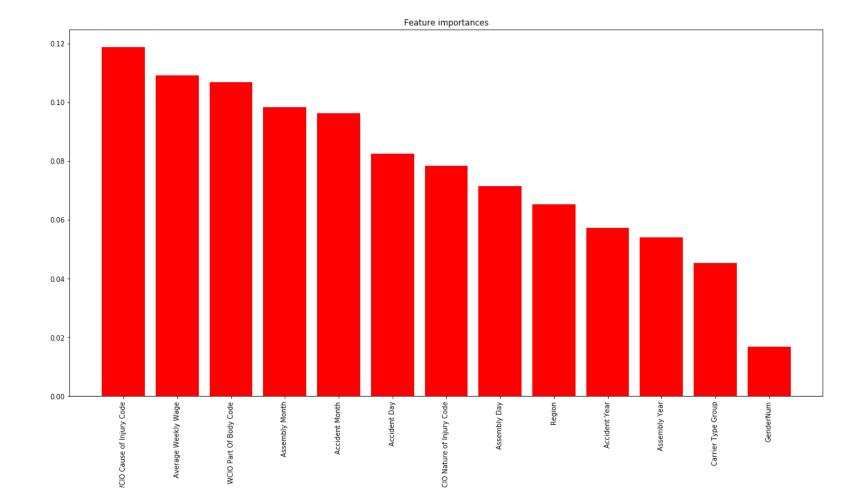
• Score: 0.743

• Precision: 0.141

• Recall: 0.696

```
In [52]: importances = rfc.feature_importances_
    indices = np.argsort(importances)[::-1]

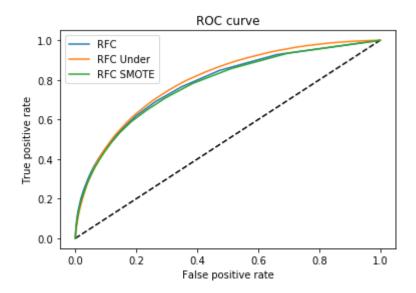
# Plot the feature importances of the forest
    plt.figure(figsize=(20,10))
    plt.title("Feature importances")
    plt.bar(range(len(indices)), importances[indices], color="r", align="center")
    plt.xticks(range(len(indices)), X.columns[indices], rotation=90)
    plt.xlim([-1, len(indices)])
    plt.show()
```



```
In [55]: # Look at the ROC curves for each sampling
    plt.figure(1)
    plt.plot([0, 1], [0, 1], 'k--')
    fpr_rfc, tpr_rfc, _ = roc_curve(y_test, y_pred_prob_rfc)
    fpr_rfc_under, tpr_rfc_under, _ = roc_curve(y_test, y_pred_prob_rfc_under)
    fpr_rfc_SMOTE, tpr_rfc_SMOTE, _ = roc_curve(y_test, y_pred_prob_rfc_SMOTE)

plt.plot(fpr_rfc, tpr_rfc, label='RFC')
    plt.plot(fpr_rfc_under, tpr_rfc_under, label='RFC Under')
    plt.plot(fpr_rfc_SMOTE, tpr_rfc_SMOTE, label='RFC SMOTE')

plt.xlabel('False positive rate')
    plt.ylabel('True positive rate')
    plt.title('ROC curve')
    plt.legend(loc='best')
    plt.show()
```



Model 2: Gradient Boost Classifier

- Tried regular, under, and SMOTE sampling of data
- Under and regular performed about the same
 - Would use under sampled faster
- Slower than random forest

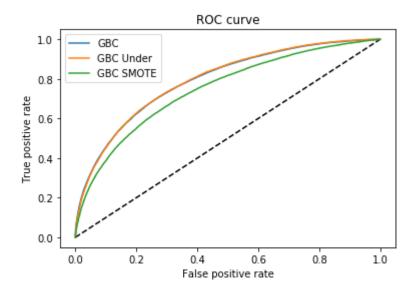
Results:

- Accuracy Score: 0.746
- Recall Score: 0.684
- Precision Score: 0.141

```
In [59]: # Look at the ROC curves for each sampling
    plt.figure(1)
    plt.plot([0, 1], [0, 1], 'k--')
    fpr_gbc, tpr_gbc, _ = roc_curve(y_test, y_pred_prob_gbc)
    fpr_gbc_under, tpr_gbc_under, _ = roc_curve(y_test, y_pred_prob_gbc_under)
    fpr_gbc_SMOTE, tpr_gbc_SMOTE, _ = roc_curve(y_test, y_pred_prob_gbc_SMOTE)

plt.plot(fpr_gbc, tpr_gbc, label='GBC')
    plt.plot(fpr_gbc_under, tpr_gbc_under, label='GBC Under')
    plt.plot(fpr_gbc_SMOTE, tpr_gbc_SMOTE, label='GBC SMOTE')

plt.xlabel('False positive rate')
    plt.ylabel('True positive rate')
    plt.title('ROC curve')
    plt.legend(loc='best')
    plt.show()
```



Light Gradient Boosting

- Grows models leaf-wise instead of depth
- Computationally less expensive than Gradient Boosting
- Can handle categorical variables label encoded
- Used regular dataset, undersampled, and SMOTE
- All three performed about the same, use the under sampled data because it is less computationally expensive

Scores

Accuracy Score LightGBM: 0.755

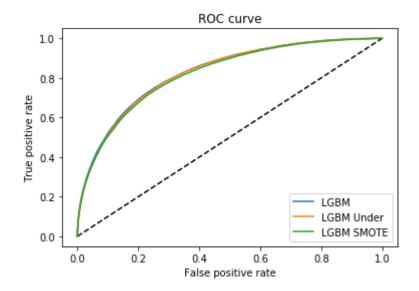
• Recall Score: 0.733

• Precision Score: 0.155

```
In [64]: plt.figure(1)
    plt.plot([0, 1], [0, 1], 'k--')
    fpr_lgbm, tpr_lgbm, _ = roc_curve(y_test, y_pred_proba_lgbm)
    fpr_lgbm_under, tpr_lgbm_under, _ = roc_curve(y_test, y_pred_proba_lgbm_under)
    fpr_lgbm_SMOTE, tpr_lgbm_SMOTE, _ = roc_curve(y_test, y_pred_proba_lgbm_SMOTE)

plt.plot(fpr_lgbm, tpr_lgbm, label='LGBM')
    plt.plot(fpr_lgbm_under, tpr_lgbm_under, label='LGBM Under')
    plt.plot(fpr_lgbm_SMOTE, tpr_lgbm_SMOTE, label='LGBM SMOTE')

plt.xlabel('False positive rate')
    plt.ylabel('True positive rate')
    plt.title('ROC curve')
    plt.legend(loc='best')
    plt.show()
```



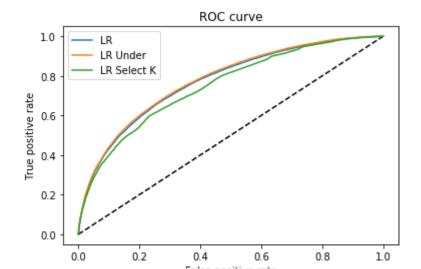
Logistic Regression

- One hot encode categorical features
- Lasso regression: number of features is > 200 after one hot encoding
- SMOTE does not work well on this number of features
 - Slow, many NAN with one hot categorical
 - also tried using SMOTENC too many categorical features
- Try undersampling majority class and using all data points
- Select K Best to reduce features ahead of times

Scores

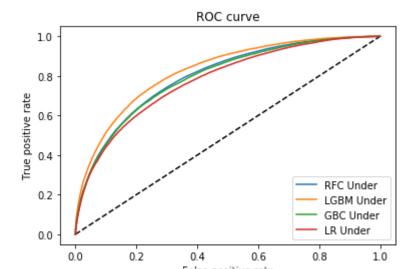
- Logistic Regression Under Sampled Score: 0.721
- Logistic Regression Under Sampled precision score 0.129
- Logistic Regression Under Sampled recall score 0.681

```
In [81]:
         # Look at the ROC curves for each of the three forest models to see which of the s
         ampling methods
         # made the biggest difference or if they are the same
         plt.figure(1)
         plt.plot([0, 1], [0, 1], 'k--')
         fpr lr, tpr lr, = roc curve(y test one, y pred prob lr)
         fpr lr under, tpr lr under, = roc curve(y test one, y pred prob lr under)
         fpr lr k, tpr lr k, = roc curve(y test2, y pred prob lr k)
         #fpr lr SMOTE, tpr lr SMOTE, = roc curve(y test, y pred prob lr SMOTE)
         plt.plot(fpr lr, tpr lr, label='LR')
         plt.plot(fpr lr under, tpr lr under, label='LR Under')
         plt.plot(fpr lr k, tpr lr k, label='LR Select K')
         #plt.plot(fpr lr SMOTE, tpr lr SMOTE, label='LR SMOTE')
         plt.xlabel('False positive rate')
         plt.ylabel('True positive rate')
         plt.title('ROC curve')
         plt.legend(loc='best')
         plt.show()
```





```
In [82]:
         plt.figure(1)
         plt.plot([0, 1], [0, 1], 'k--')
         # Use the unique names for y test here to make sure they don't plot incorrectly
         fpr rfc under, tpr rfc under, = roc curve(y test label, y pred prob rfc under)
         fpr lr under, tpr_lr_under, _ = roc_curve(y_test_one, y_pred_prob_lr_under)
         fpr lgbm under, tpr lgbm under, = roc curve(y test label, y pred proba lgbm unde
         r)
         fpr gbc under, tpr gbc under, = roc curve(y test label, y pred prob gbc under)
         plt.plot(fpr rfc under, tpr rfc under, label='RFC Under')
         plt.plot(fpr lgbm under, tpr lgbm under, label='LGBM Under')
         plt.plot(fpr gbc under, tpr gbc under, label='GBC Under')
         plt.plot(fpr lr under, tpr lr under, label='LR Under')
         plt.xlabel('False positive rate')
         plt.ylabel('True positive rate')
         plt.title('ROC curve')
         plt.legend(loc='best')
         plt.show()
```

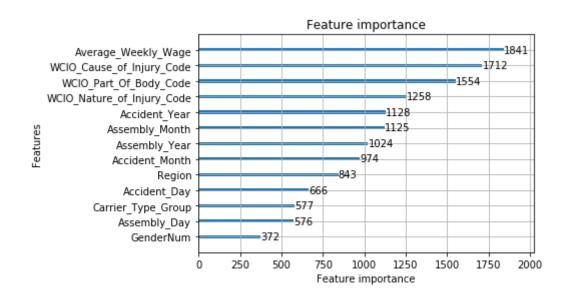


Cross Validation LGBM

- Check what the log-loss is when training set split into 5 folds
- Looked at STD to see if much deviation between the log-loss
 - Found a low STD (.006), the difference between the log-loss in folds was low
- Use to determine over-fit to part of the training set
- Early stopping

```
In [93]: # Plot the feature importances
    print('Feature Importances')
    ax = lgbm.plot_importance(clf_under, max_num_features=20)
    plt.show()
```

Feature Importances



Optimize Parameters for LightGBM

- Set up a grid search
- Did larger ranges for parameters at different times for computational reasons
- Narrowed down to different ranges
- Focused on sub_feature, min_data_leaf, bagging_fraction, bagging_freq, learning_rate, num_leaves

Model Shortcomings:

- Expert in the field could determine other usable features
- Recall comes at high precision cost
- Only optimized the best model
- Dates are not treated cyclically
 - January 1, December 12
- SMOTE implementation
 - Better for continuous variables

Conclusion:

- Best Model: Light Gradient Boost Model
 - Best area under receiver operating characteristic
 - Fast
- Best Sampling Method:
 - Under Sampling
- Potential Future Projects:
 - Predict which cases go to Supreme Court
 - Look into regions of New York with high rates of claims, why is that?
 - Cluster analysis into claims
- If you don't want your insurance claim controverted:
 - Puncture your great toe in May in the Finger Lakes region of New York
 State