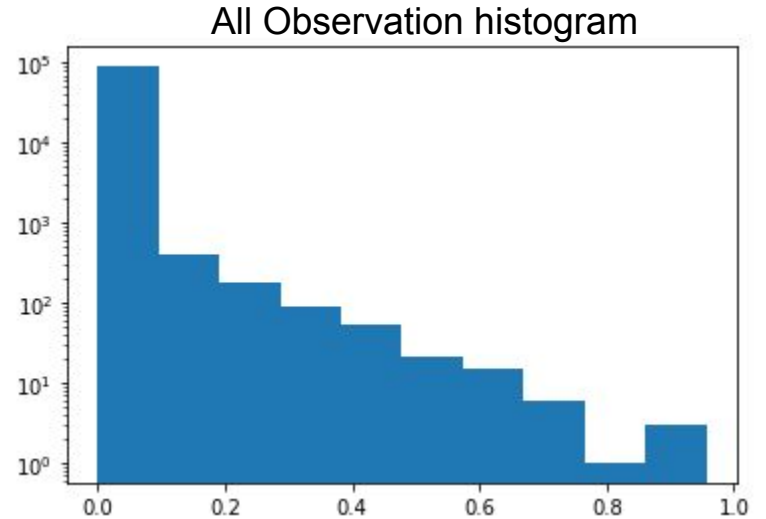


- Matrix Size: 250\*361
- Total Observation: 90250
- Top 0.5% CutOff: 0.16

Very Sparse Matrix, after processing, there are 451 observations of 1, the rest are set to 0.

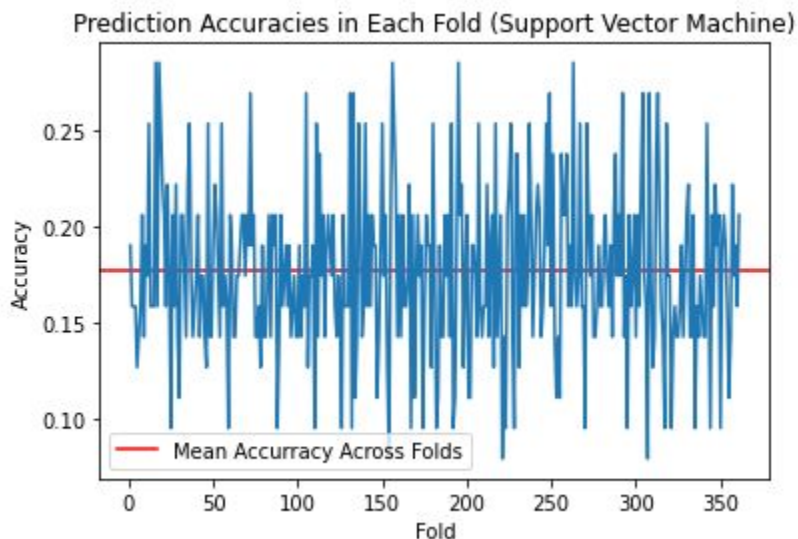


# Plant Species

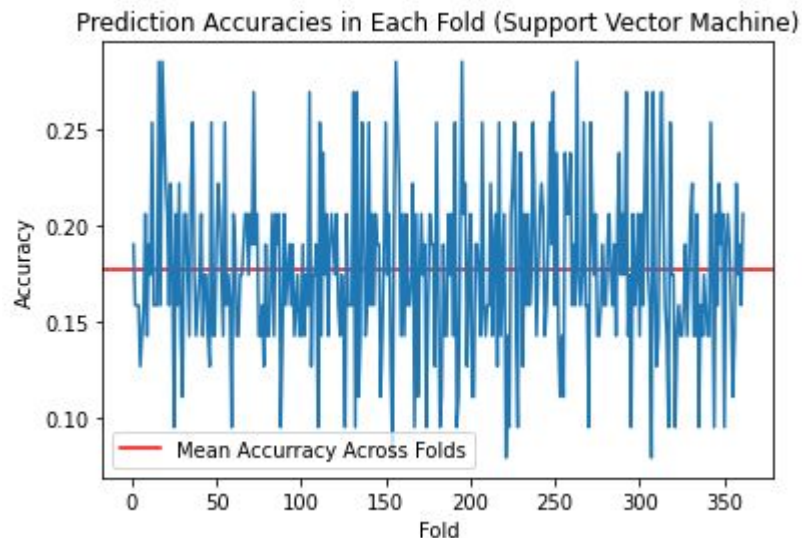
13 Unique Species, 250 samples total, Average 19 samples per label.

{'POPR', 'ELSC', 'POLE', 'TRSP', 'ELEL', 'ELTR', 'POST', 'FETH', 'FEBR', 'FESA', 'POAL', 'ACNE', 'ACLE'}

## Continuous Measurement



## Binary Matrix



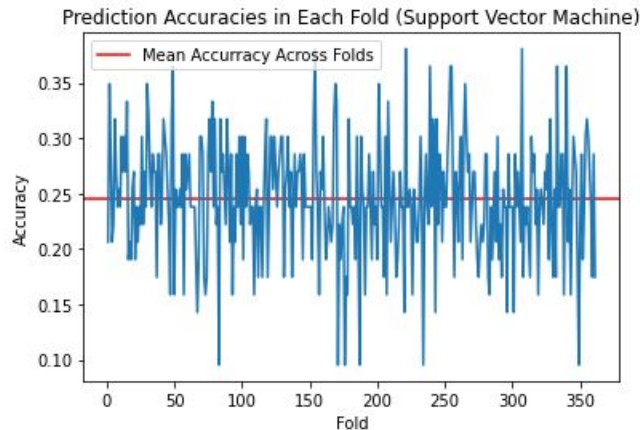
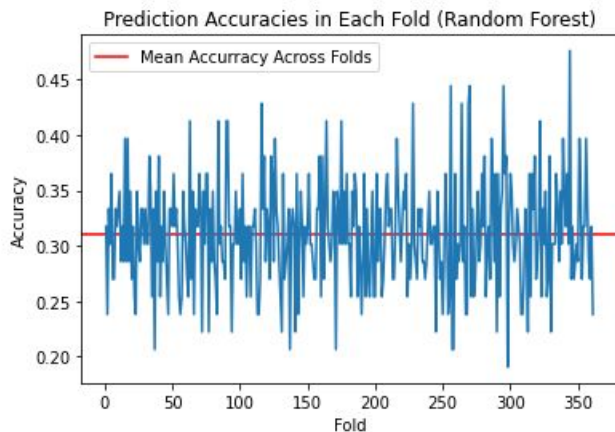
# Plant Species (grouped by genus)

{'PO', 'TR', 'AC', 'FE', 'EL'}

Random Guess Accuracy: 20%

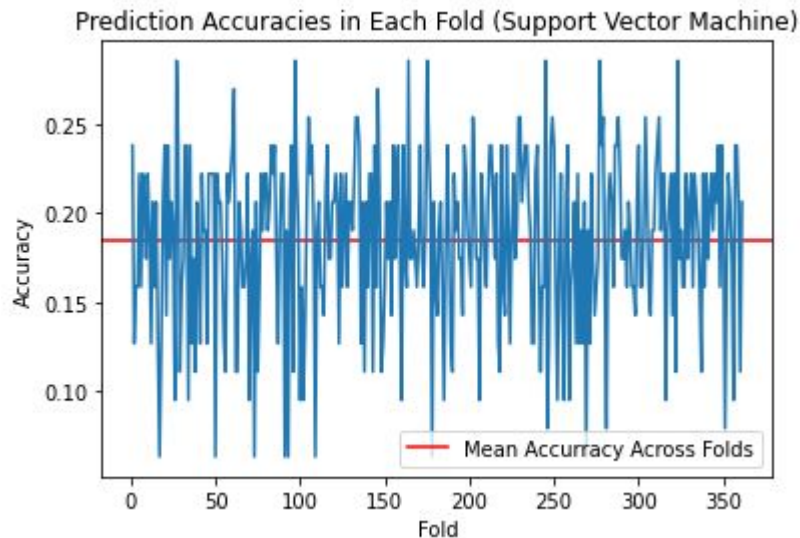
Mean prediction Accuracy using SVM: 24.57%

Mean prediction Accuracy using Random Forest: 31%



# Gradient

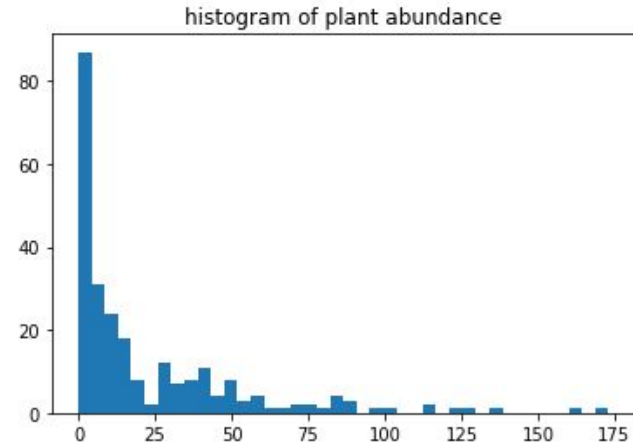
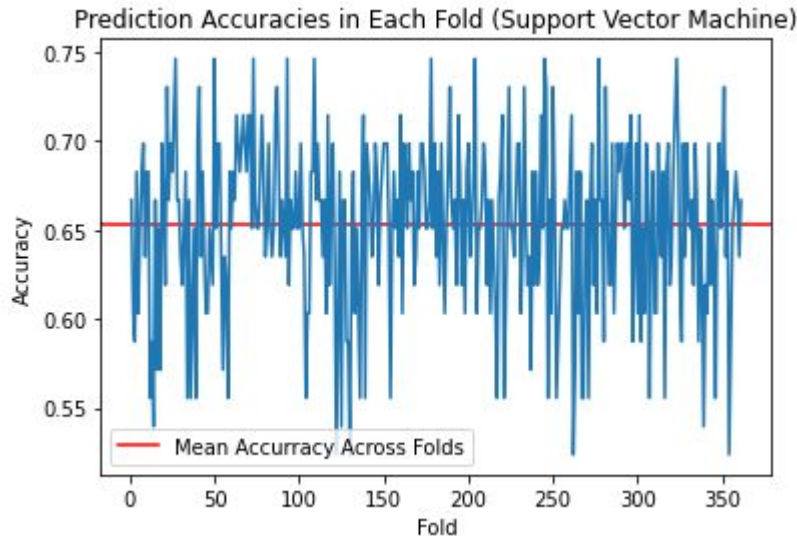
{'Avery', 'Treasury', 'Ruby', 'HunterHill', 'RP', 'Cinnamon', 'KP', 'JC2', 'Teocali', 'JC1'}



# Plant Abundance (Ignoring Species and Genus)

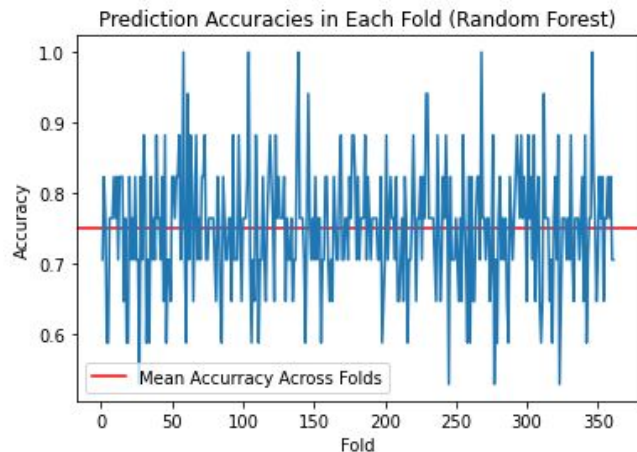
Continuous measurements, generalized into 2 class: {Abundant, Not Abundant}

Currently using 5 as the cutoff.

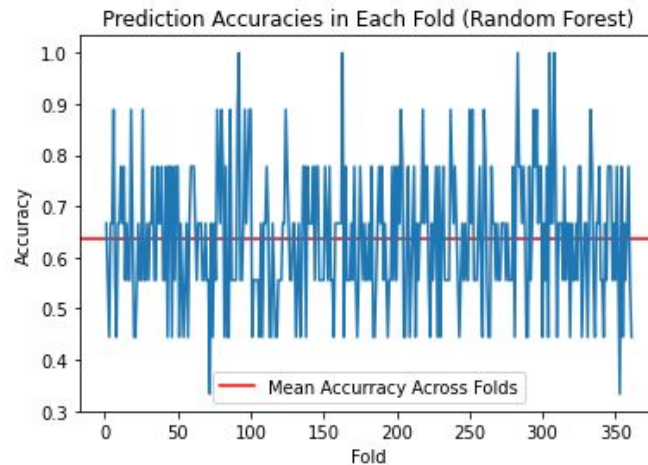


# Plant Abundance by Genus

PO

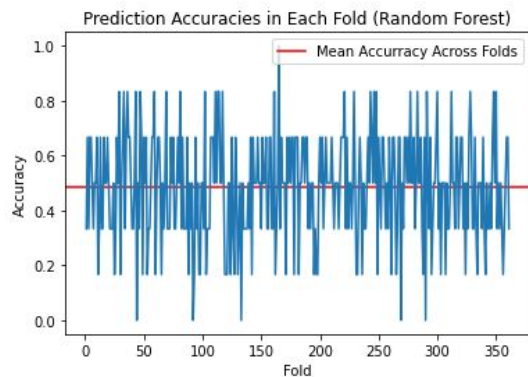


TR

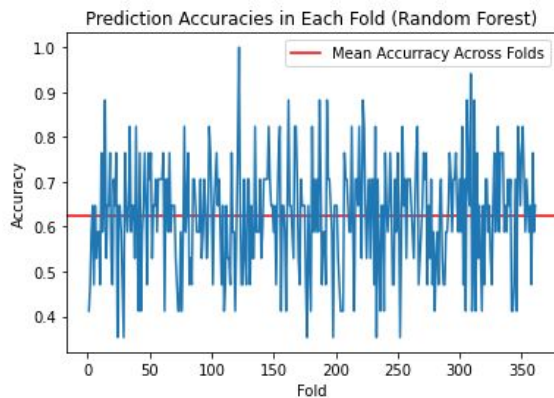


# Plant Abundance by Genus

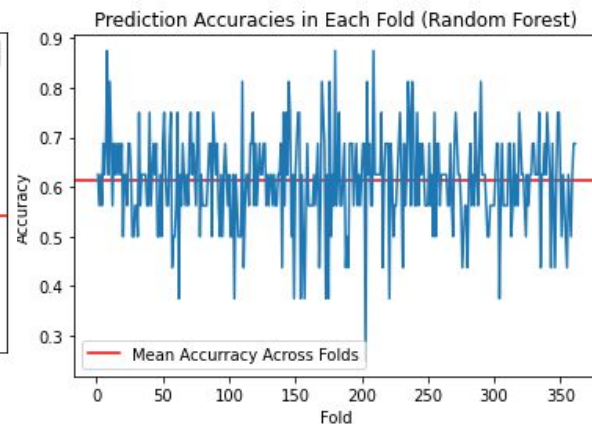
AC



FE



EL



# Plant Abundance by Genus

Mean Accuracy Prediction by Genus:

PO: 0.751

TR: 0.638

FE: 0.623

EL: 0.6125346260387812

AC: 0.483



# POPR

Sample Size is small (20), and extremely unbalanced:

## SVM prediction with all ASVs:

## Real Labels:

['Abundant', 'Abundant', 'Abundant', 'Abundant', 'Abundant', 'Abundant', 'Abundant', 'Abundant', 'Abundant', 'Abundant', 'Abundant',  
'Abundant', 'Abundant', 'Not Abundant', 'Abundant', 'Abundant', 'Abundant', 'Abundant', 'Abundant', 'Abundant', 'Not  
Abundant']

## Predictions:

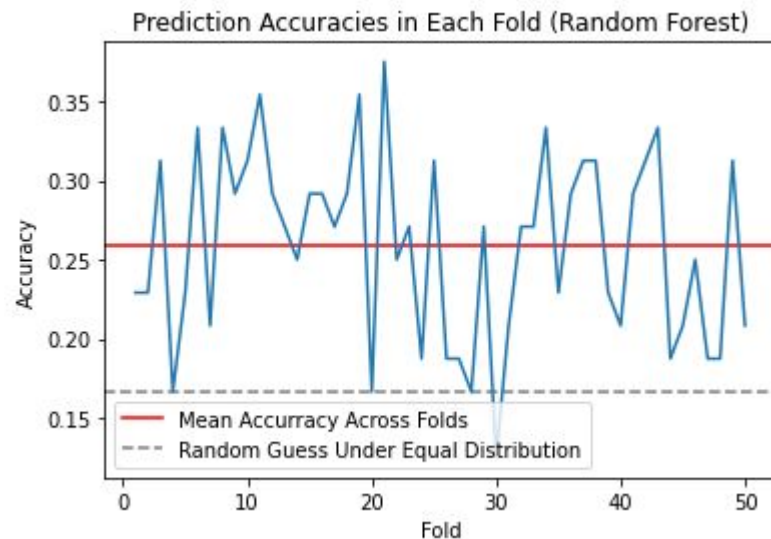
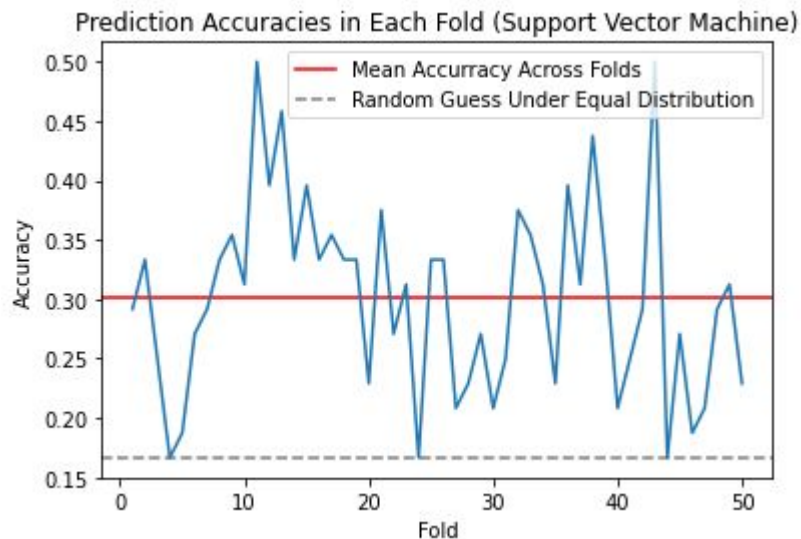
[illegible]

# Species Prediction (One VS Others)

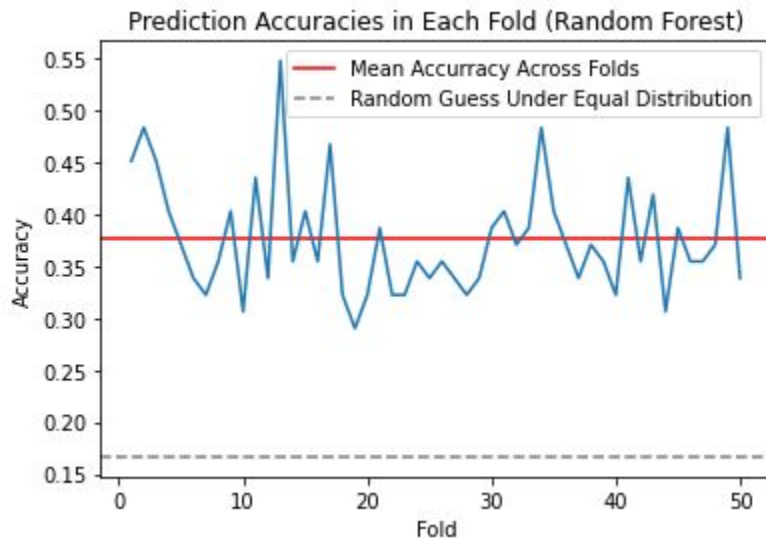
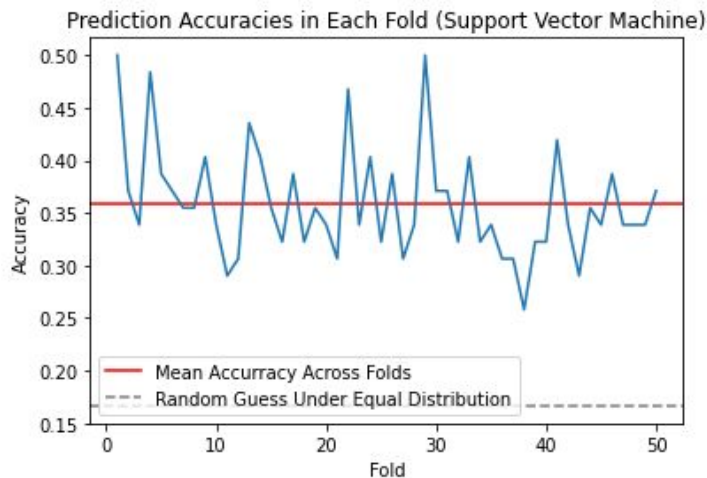
SVM, NB and RF all yield meaningless predictions (no true positives among all species and samples). With one false positive in the prediction of ELTR nad FETH.

Most predictions are completely all negatives.

# Leaves-Gradient

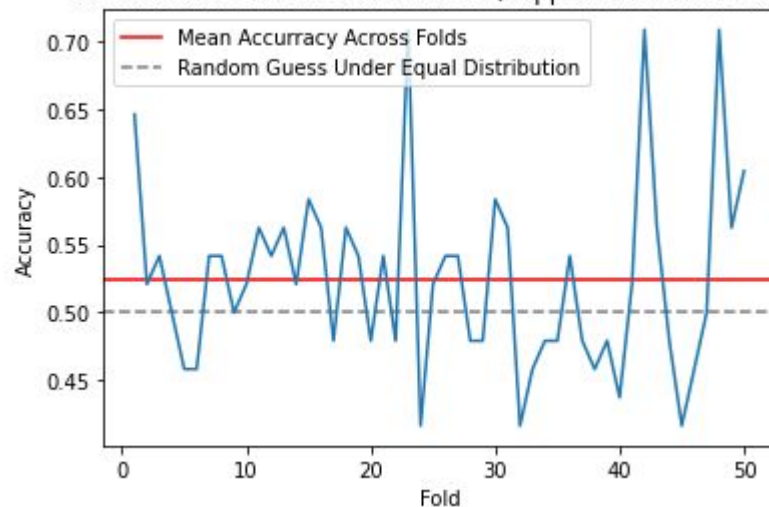


# Root-Gradient

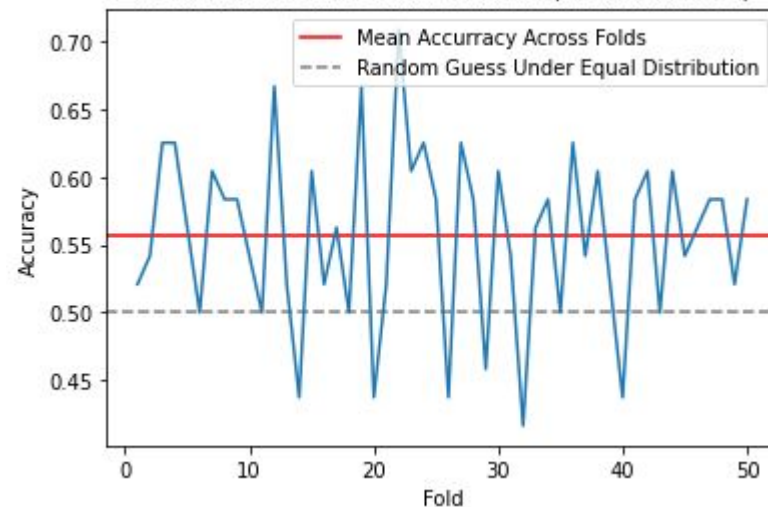


# Leave-Richness

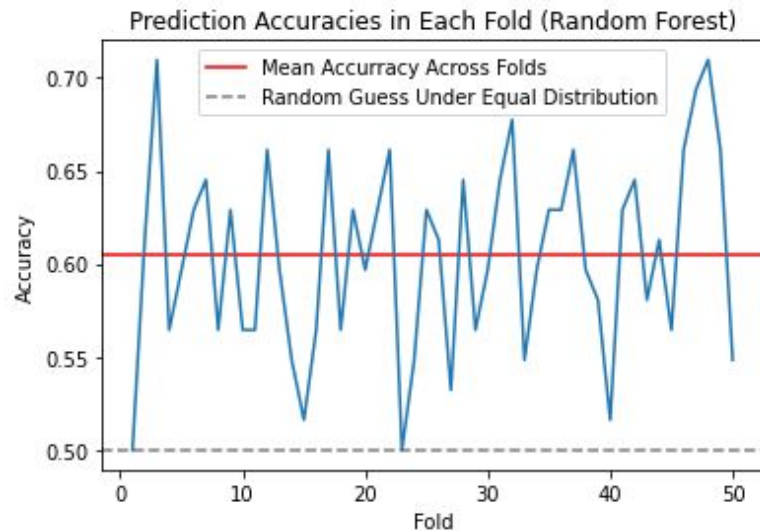
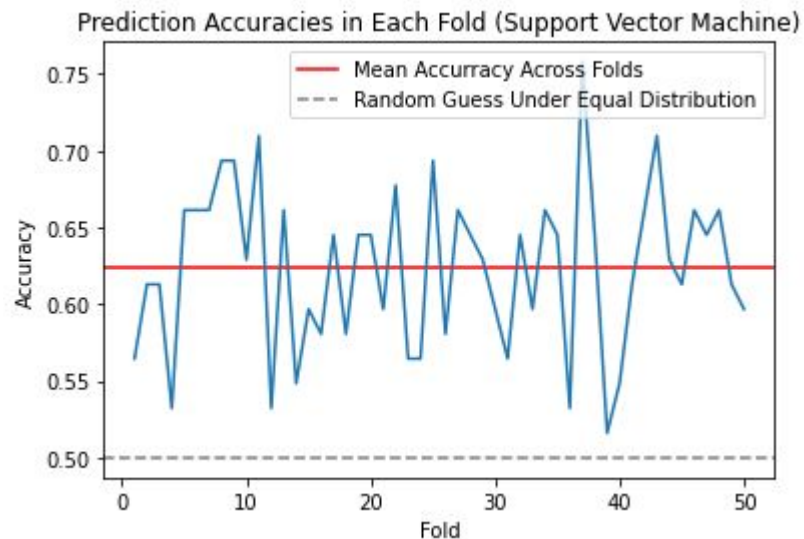
Prediction Accuracies in Each Fold (Support Vector Machine)



Prediction Accuracies in Each Fold (Random Forest)

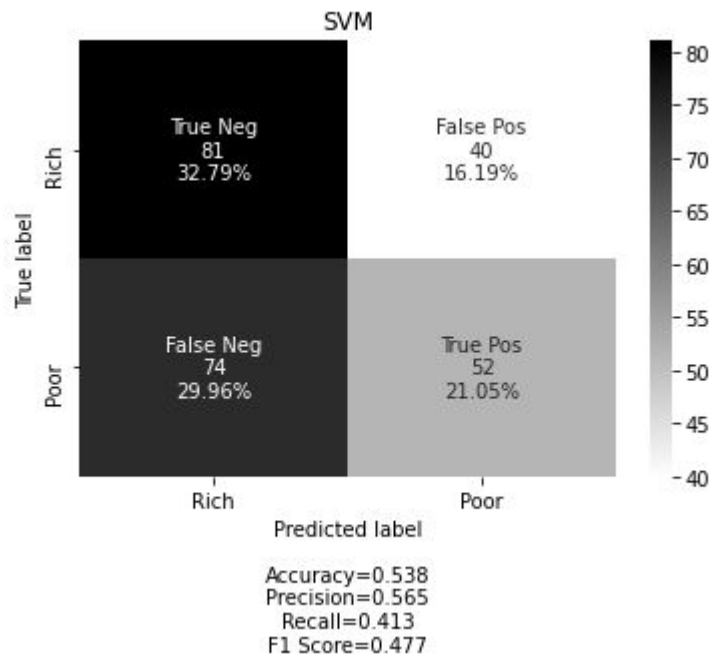
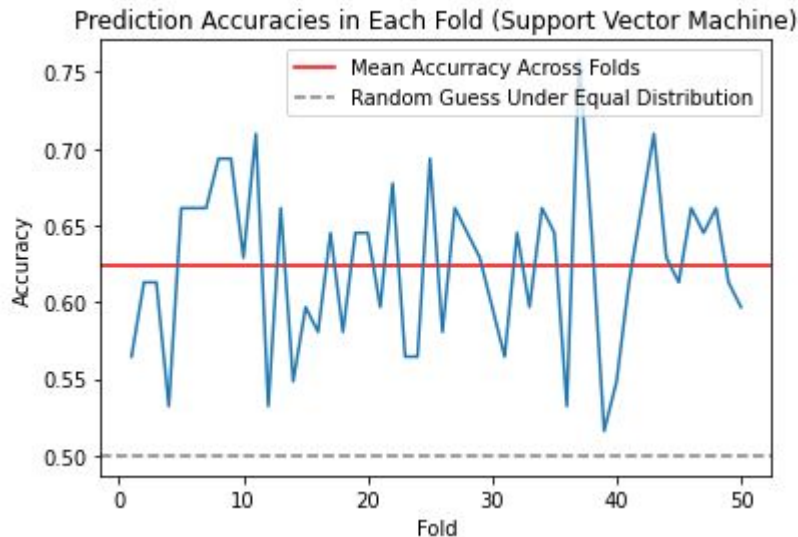


# Root-Richness

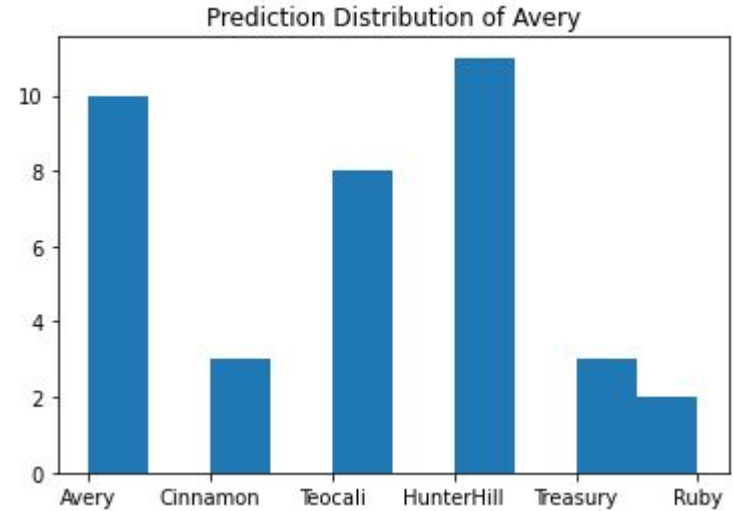
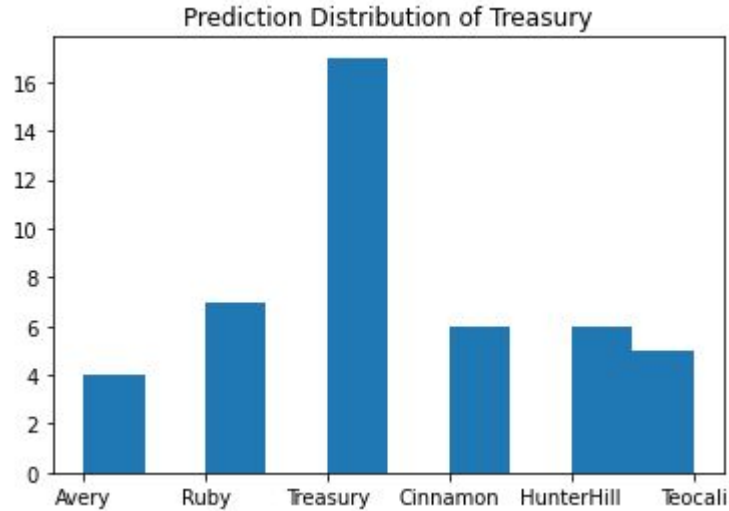


# Richness using total-N (seeing minor signal so far)

Using Top 30 Features



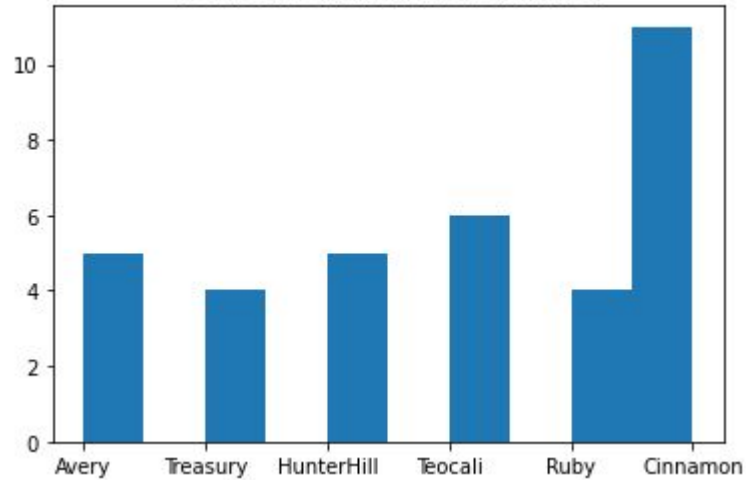
# Gradient Prediction By Groups



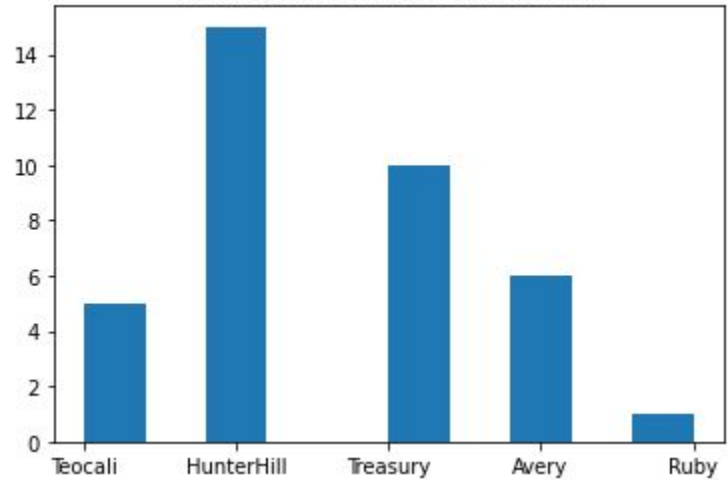


# Gradient Prediction By Species

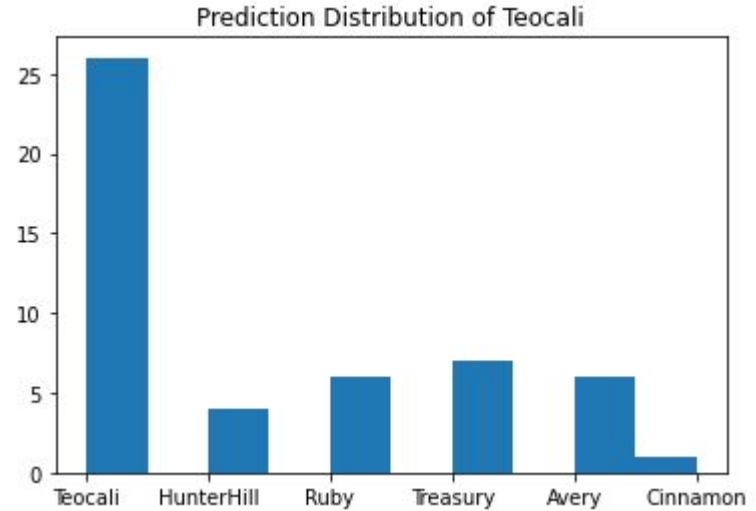
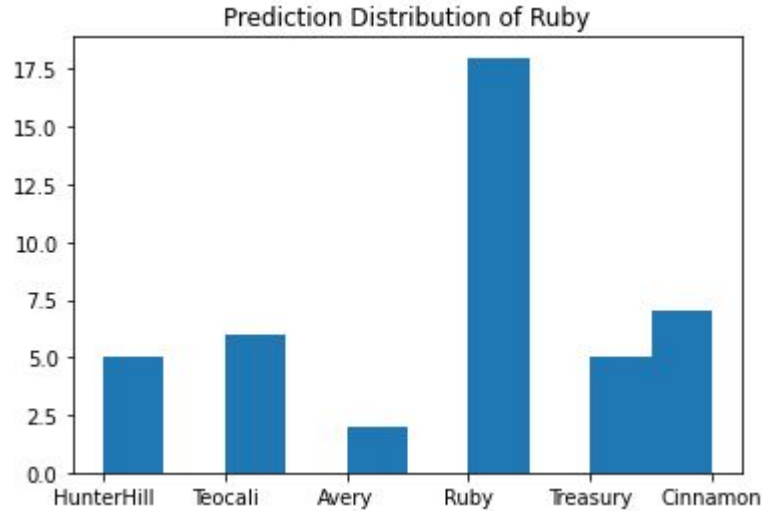
Prediction Distribution of Cinnamon



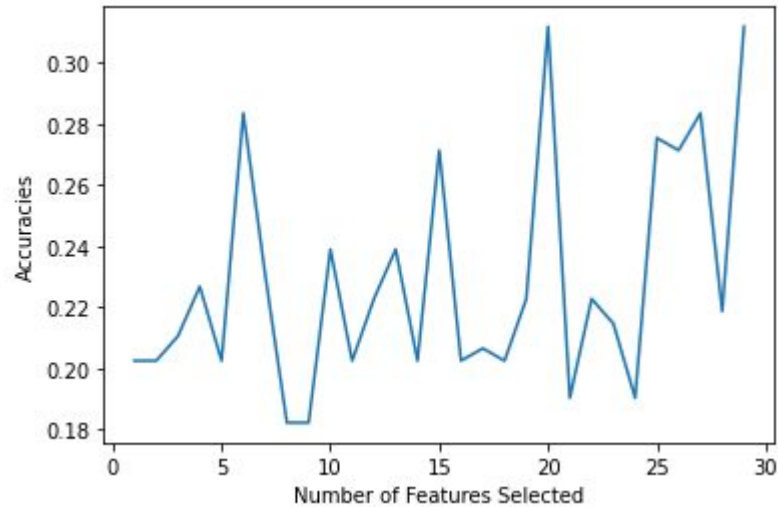
Prediction Distribution of HunterHill



# Gradient Prediction By Species

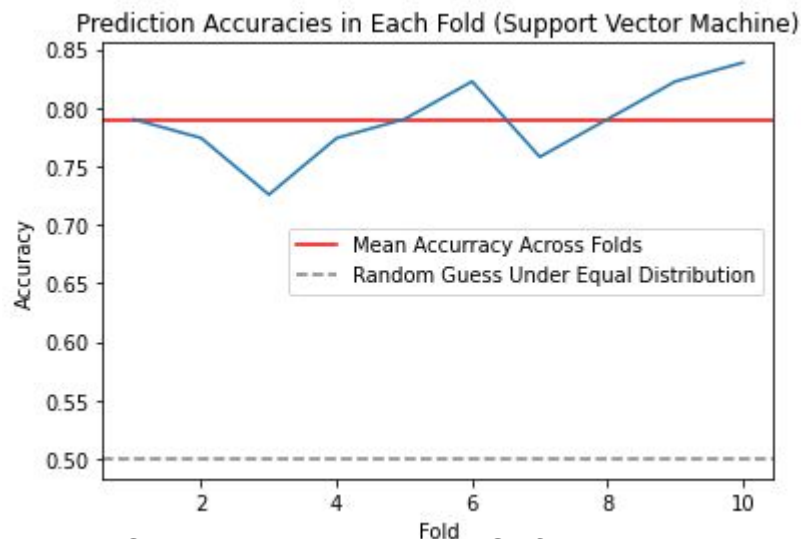


# Feature Selection on Gradient



# Treasury

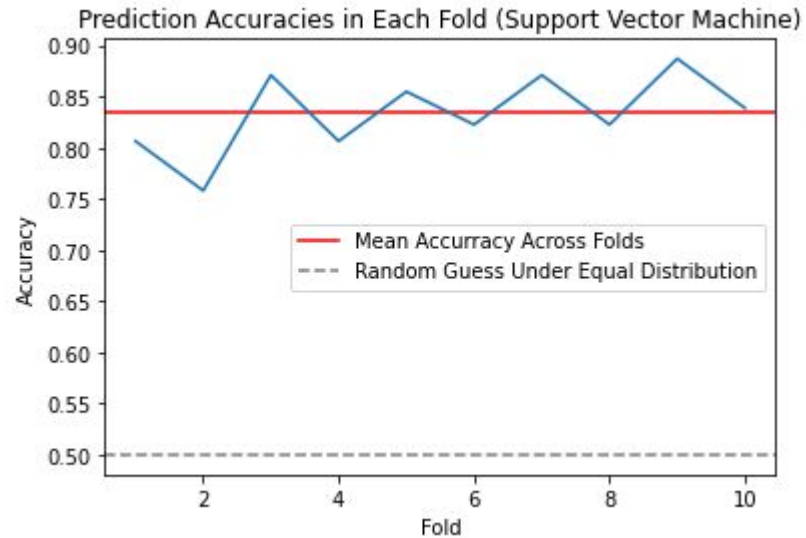
Mean Accuracy Across Folds 0.7887096774193548



One vs Rest without FS, for this plot,  
its binary labels of Treasury vs Not  
Treasury

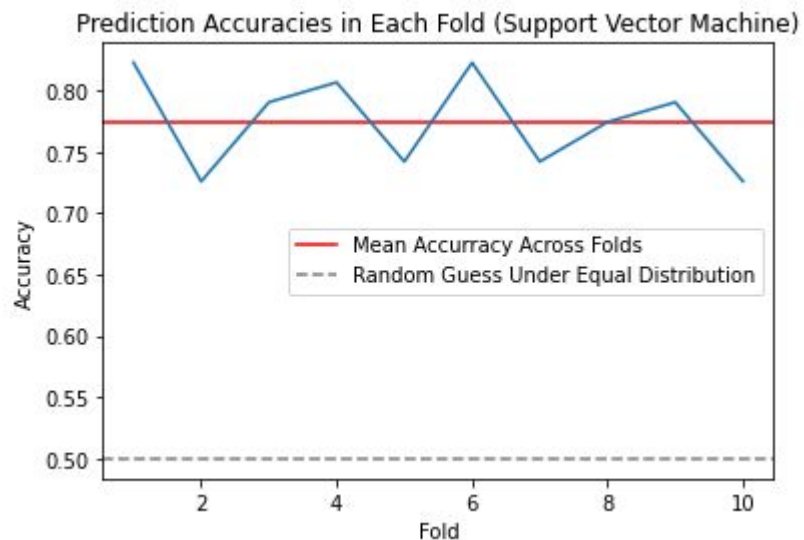
# Cinnamon :

Mean Accuracy Across Folds 0.8338709677419354



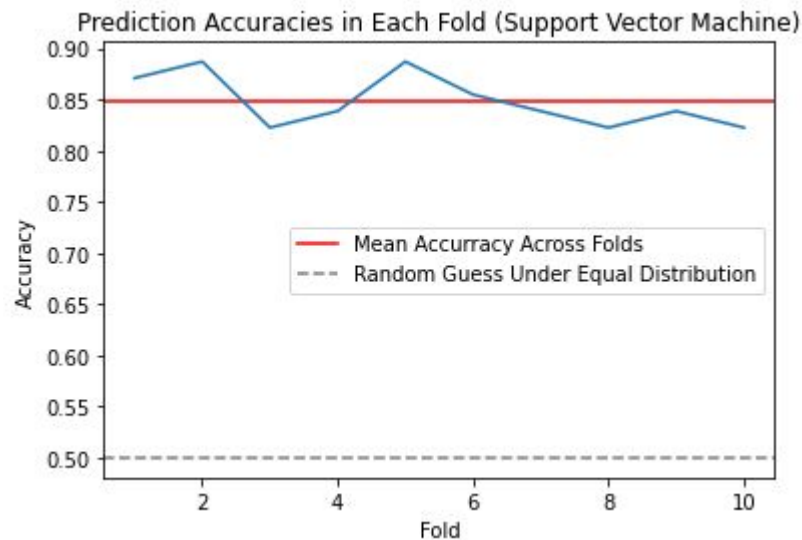
# Avery

Mean Accuracy Across Folds 0.7741935483870968



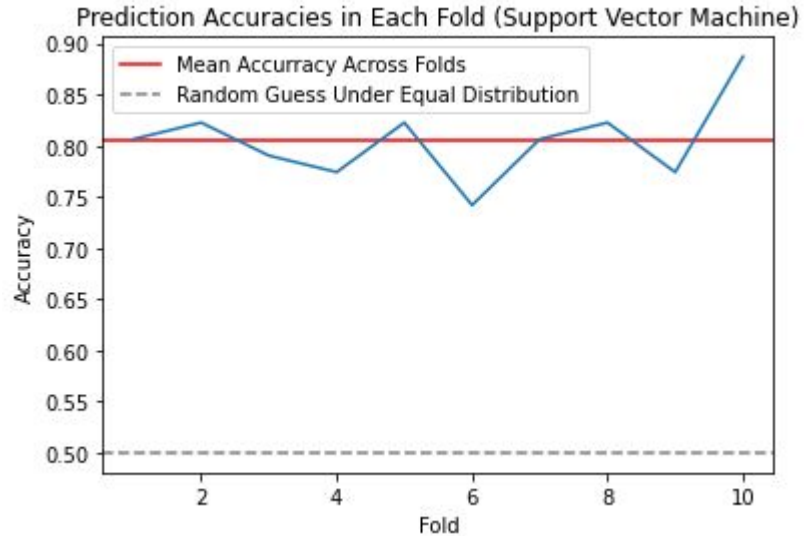
# Hunterhill

Mean Accuracy Across Folds 0.8483870967741935



# Teocali

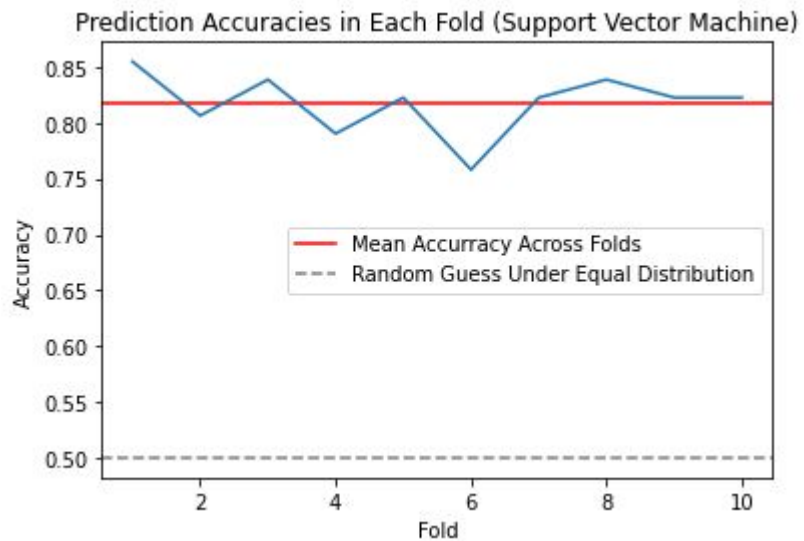
Mean Accuracy Across Folds 0.8048387096774194



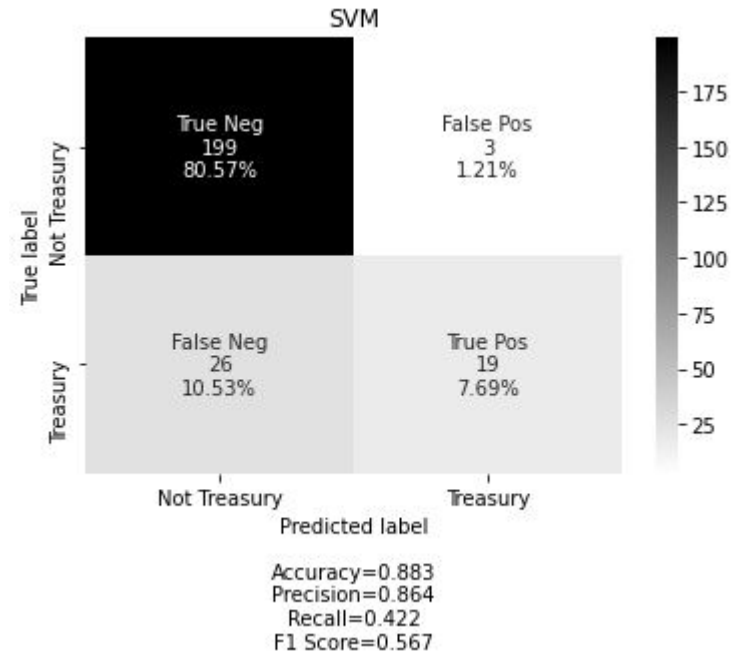
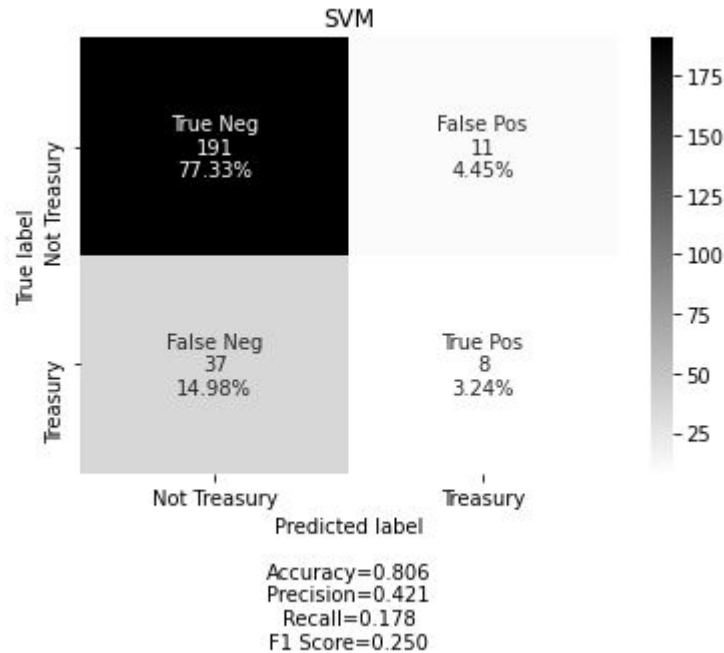


# Ruby

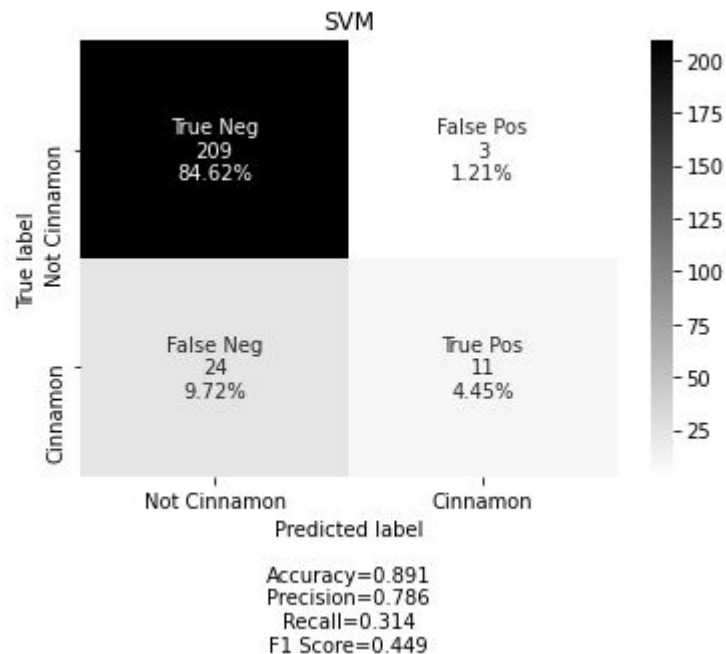
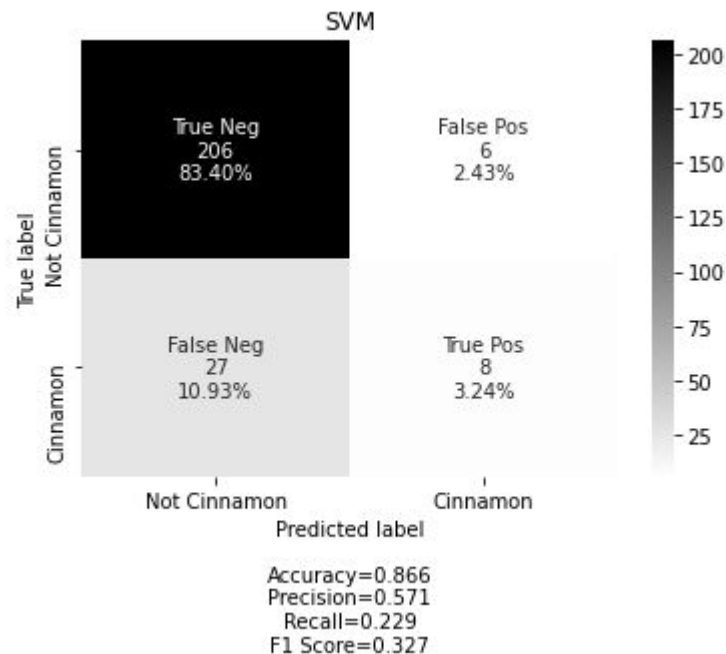
Mean Accuracy Across Folds 0.817741935483871



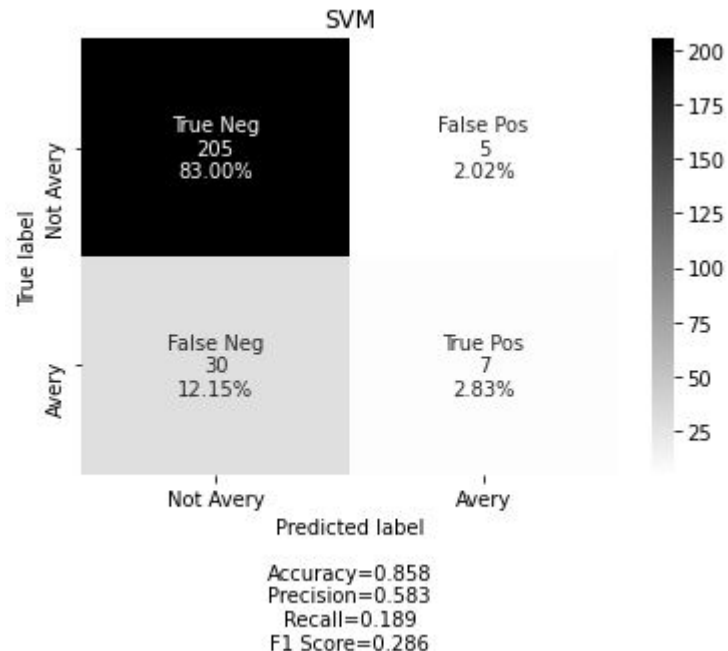
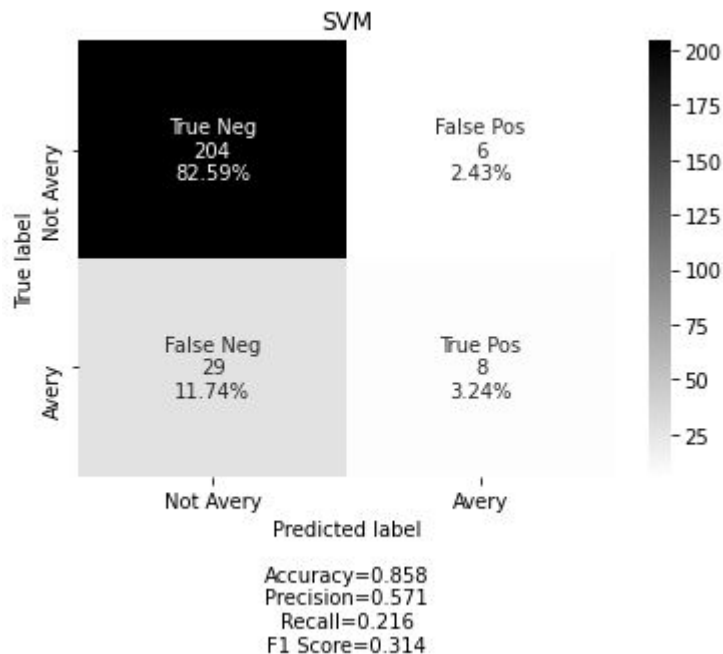
Different Number of features impact different gradient prediction precision  
Top 30 features (left) vs Top 100 features(right) for Treasury, select K best  
with Chi2 scoring function.



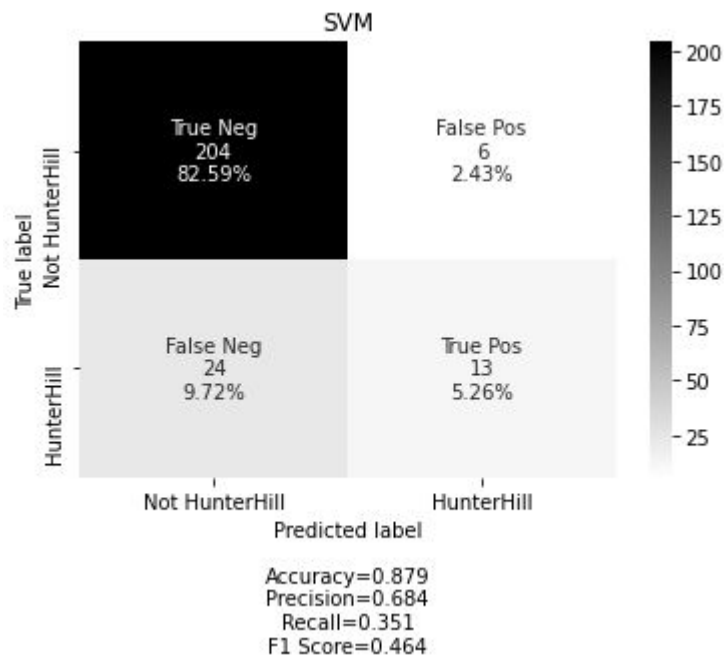
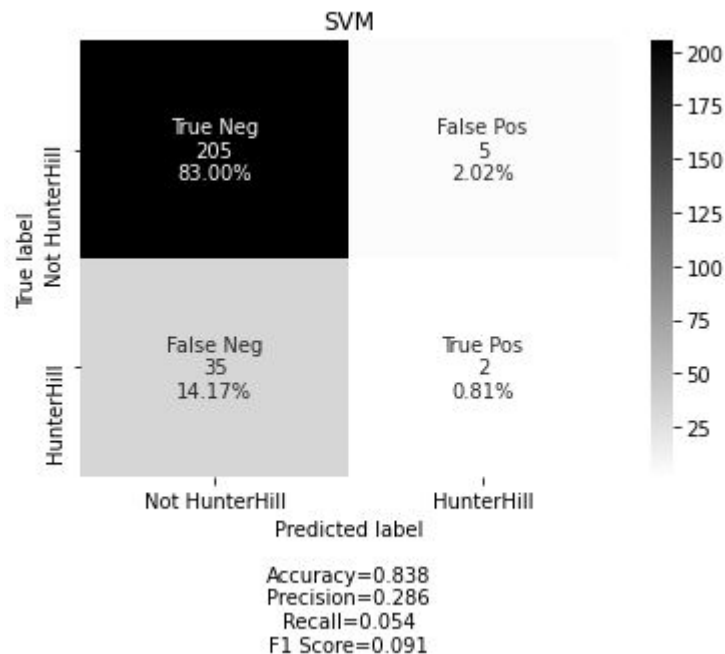
# Cinnamon



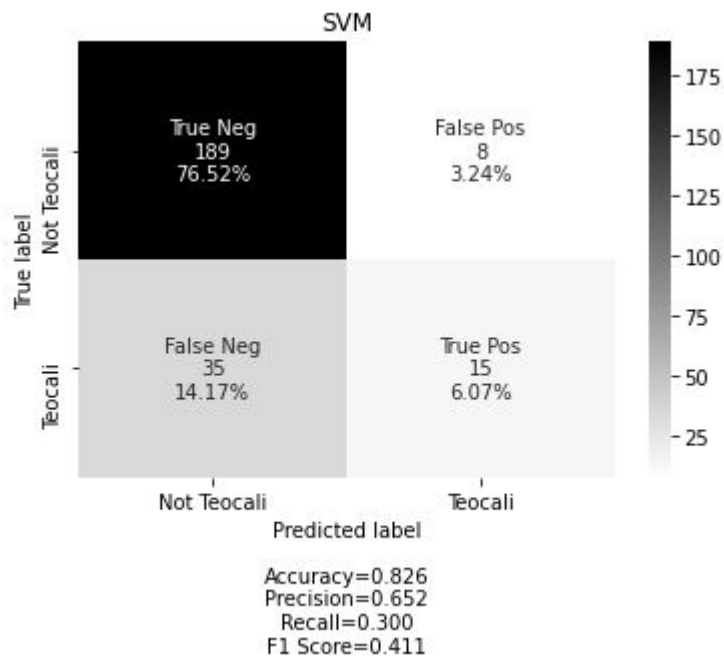
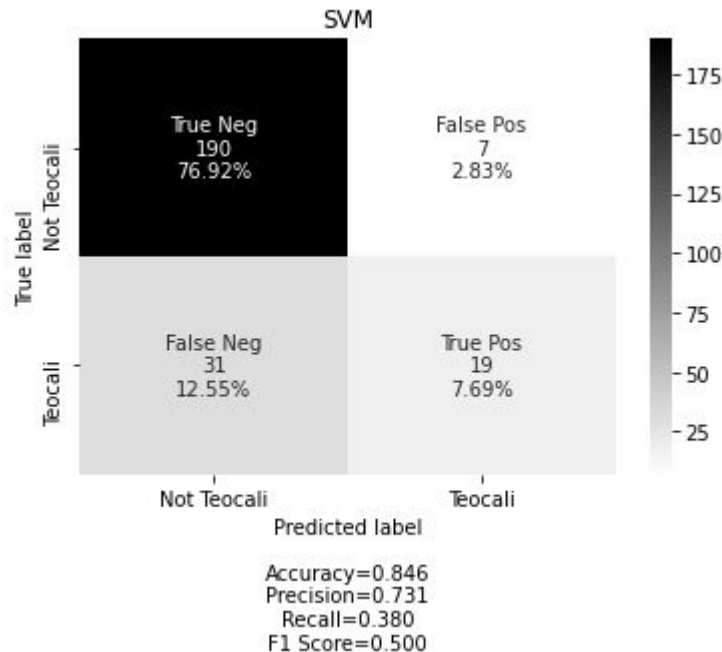
# Avery



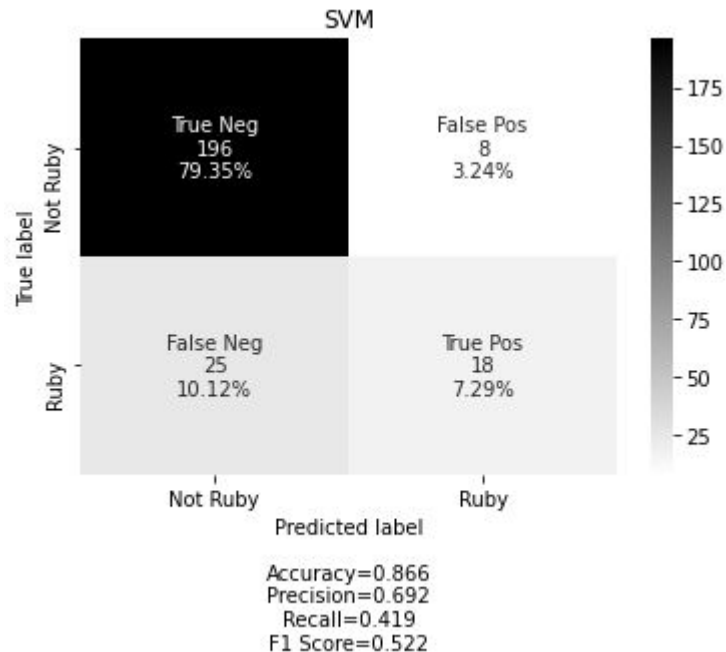
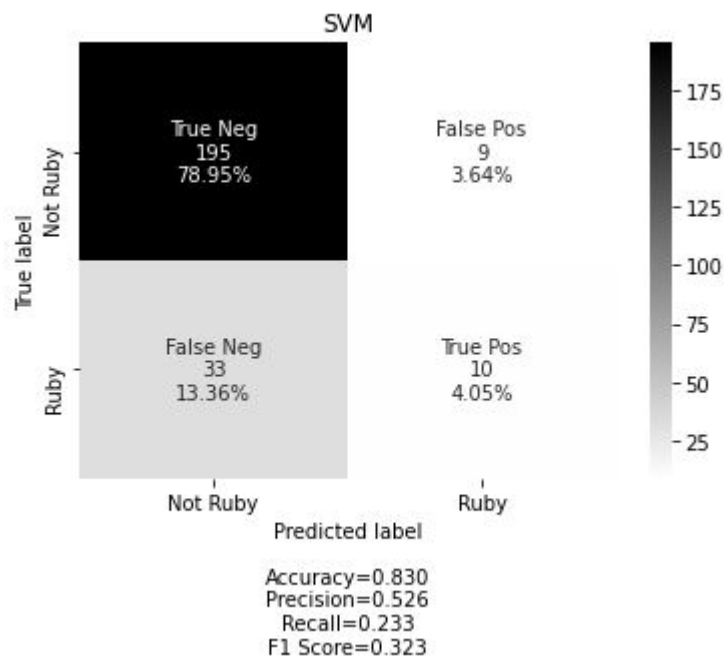
# Hunterhill



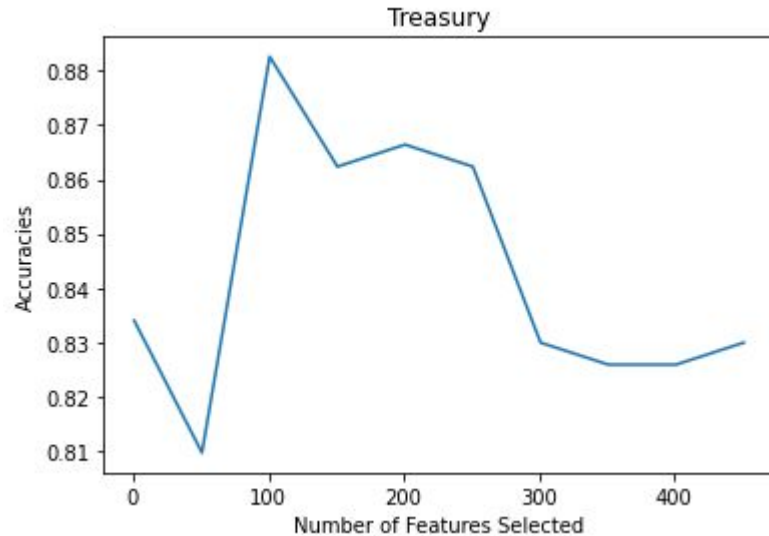
# Teocali



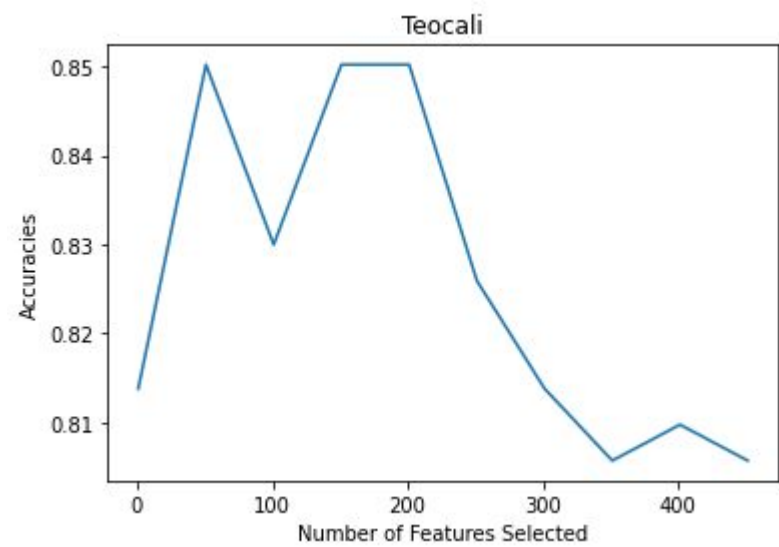
# Ruby

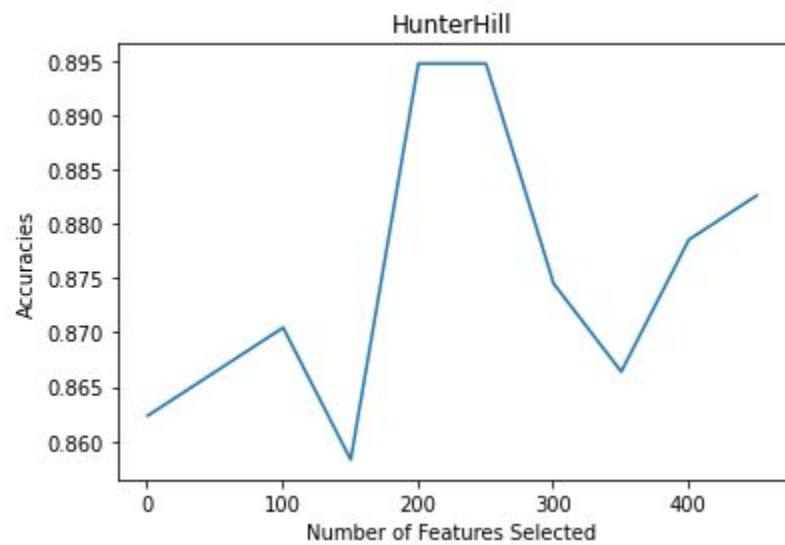


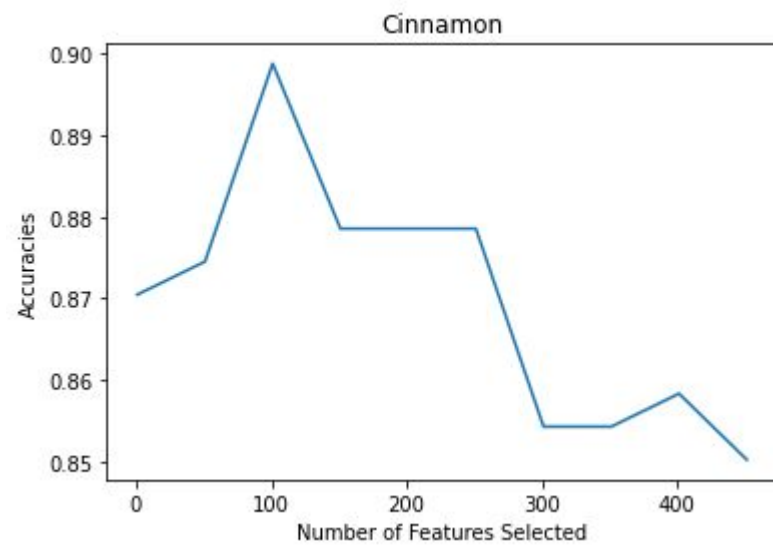
# Number of Top Features Selected (increment of 50)

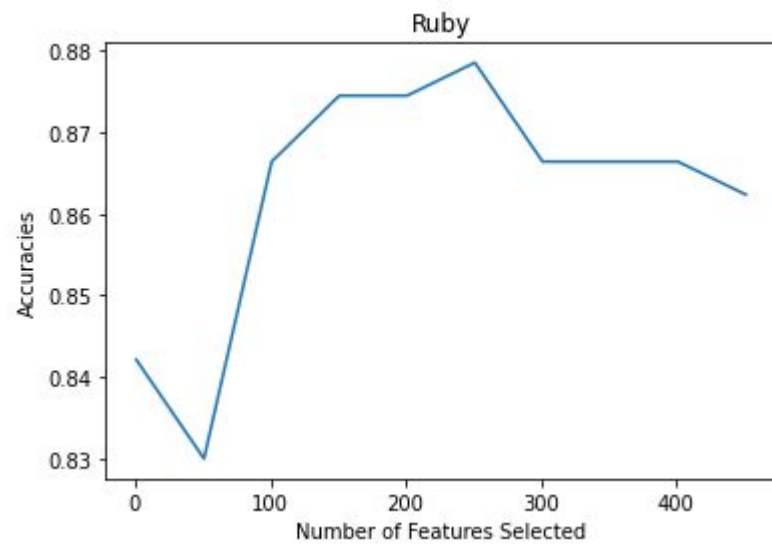


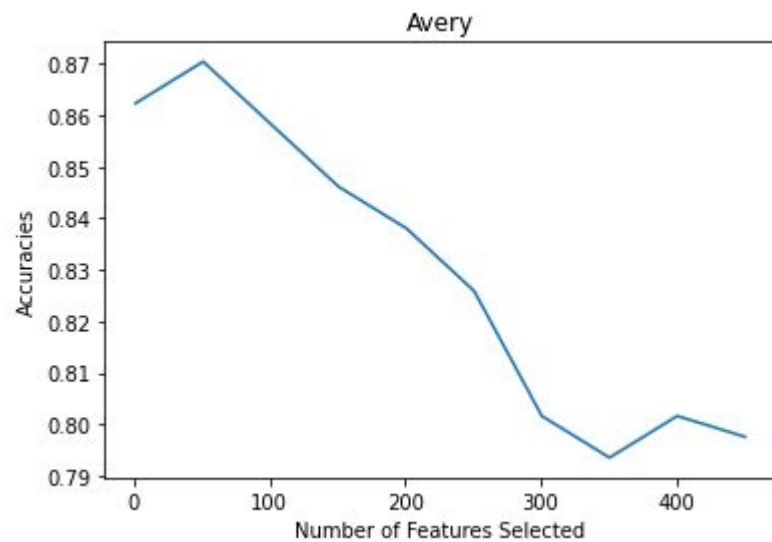




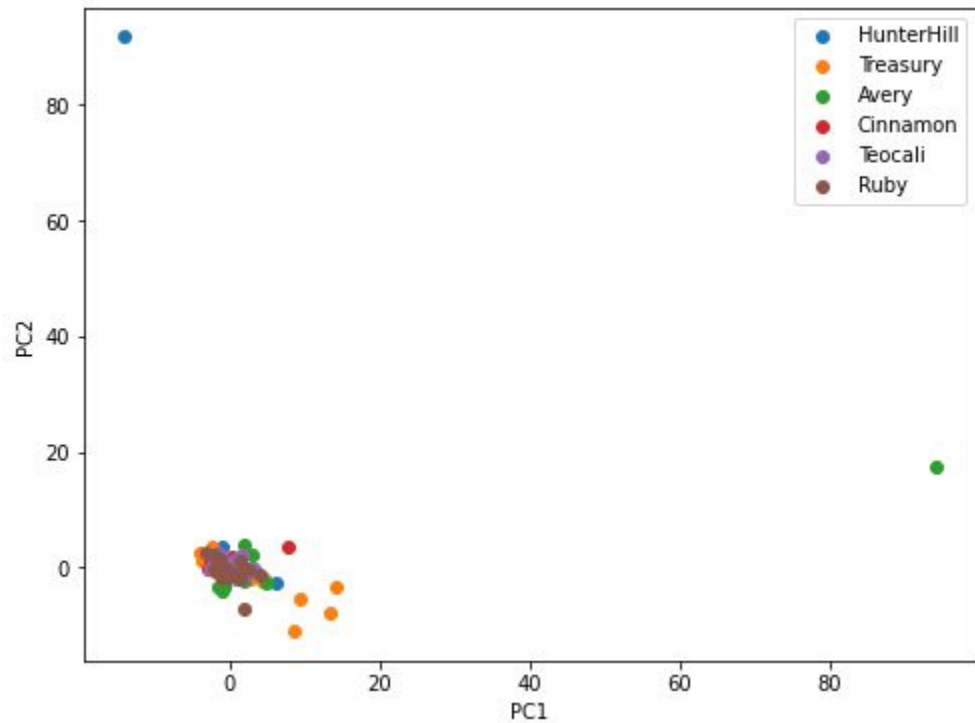




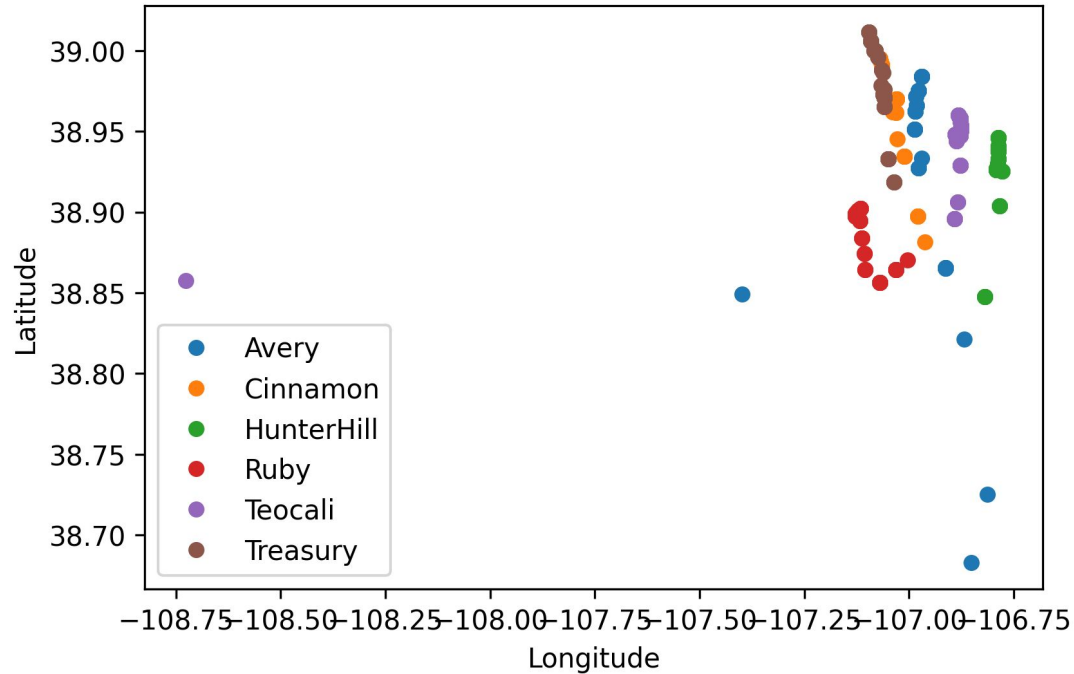




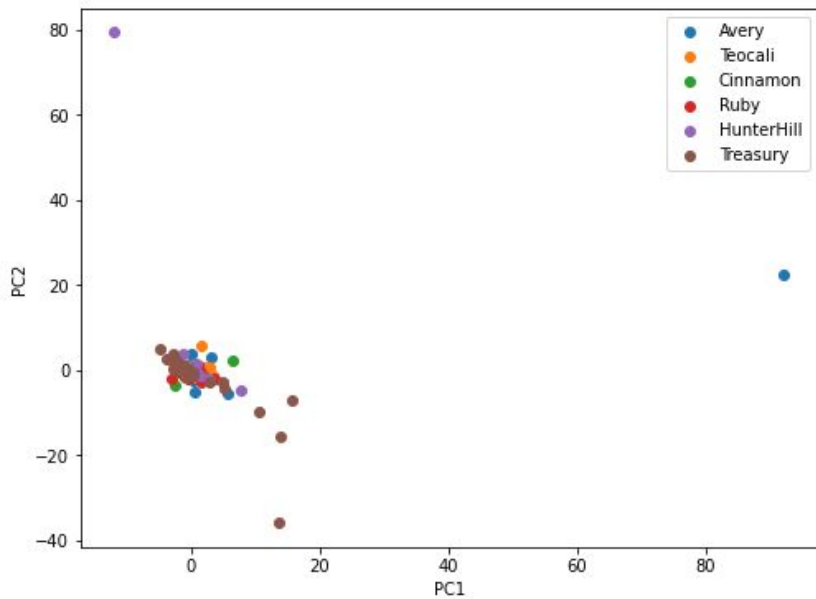
PCA Plot (Not much Variance Explained, as expected, about 1.5 percent variance explained on PC1 and PC2)



## Sample Map (one odd Teocali Sample) (old version with ELEL)

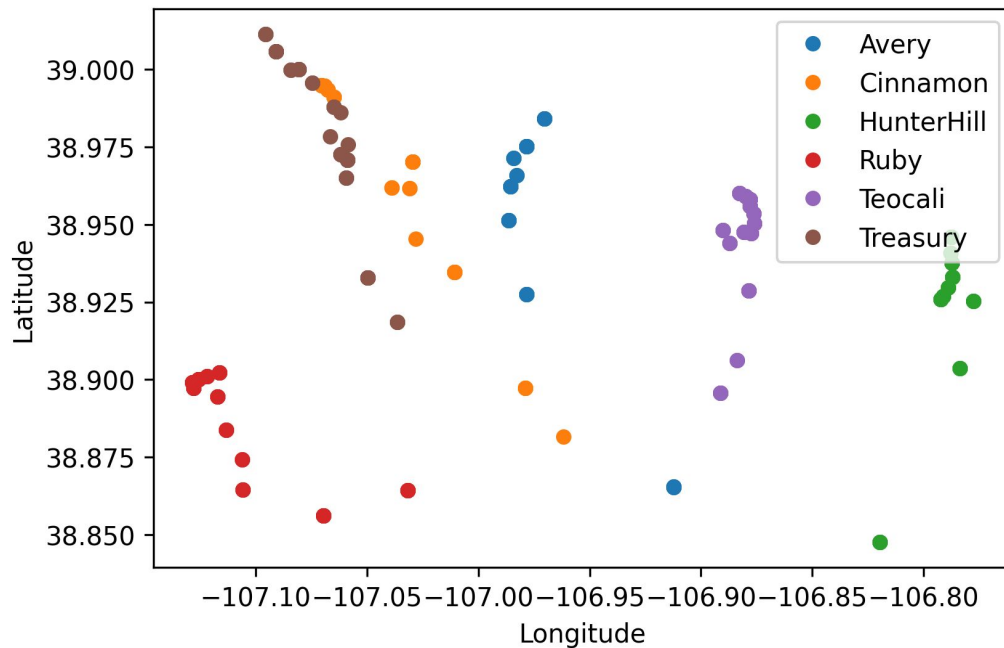


Updated PCA (slightly better, [0.01696325 0.01507139])

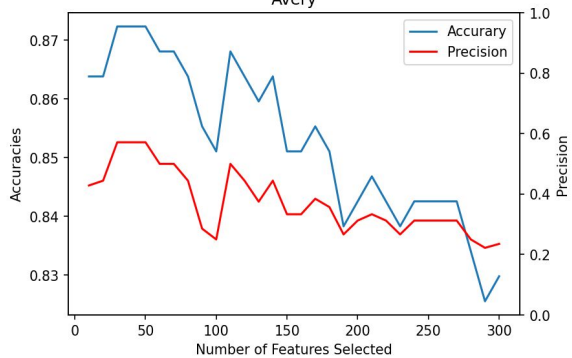




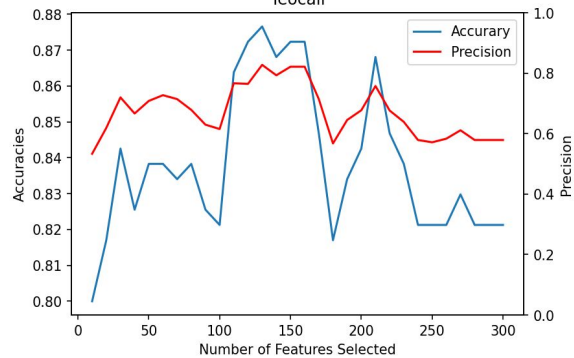
# Updated Map (removed ELEL from data)



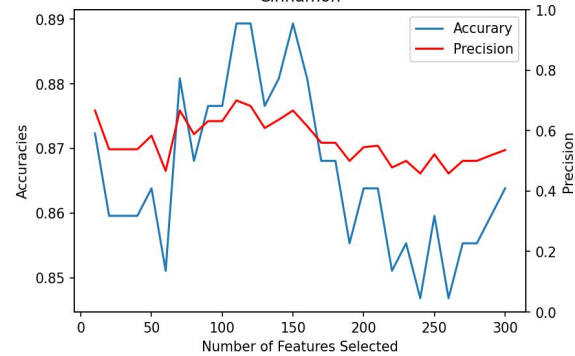
Avery



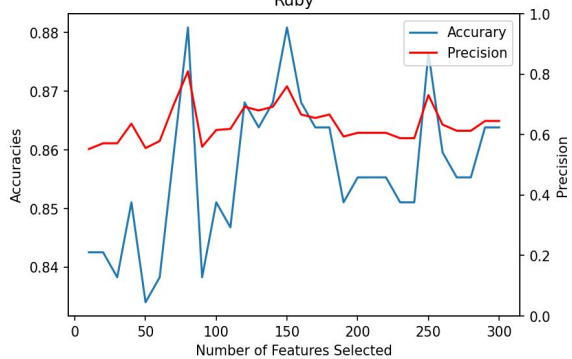
Teocali



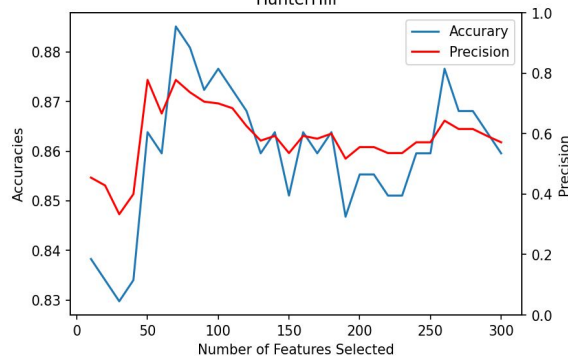
Cinnamon



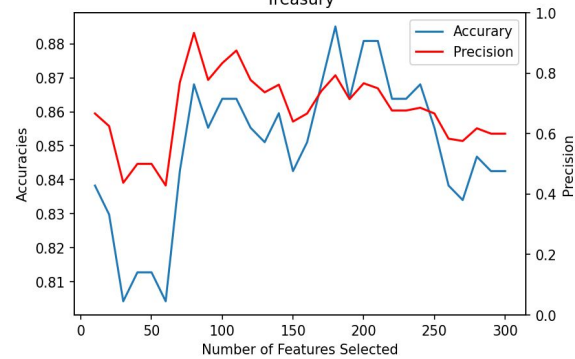
Ruby

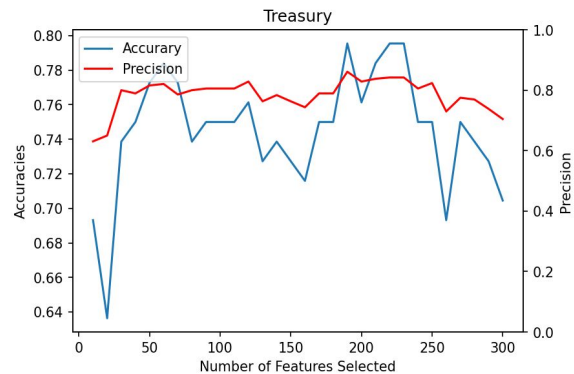
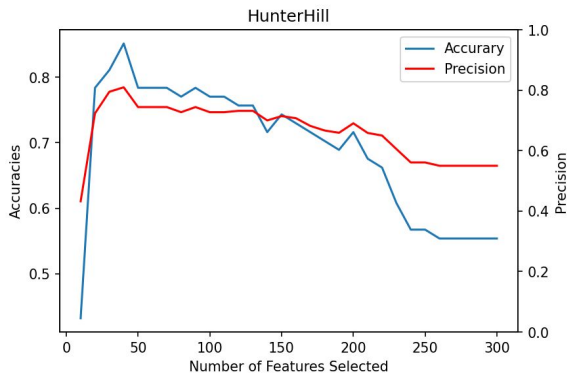
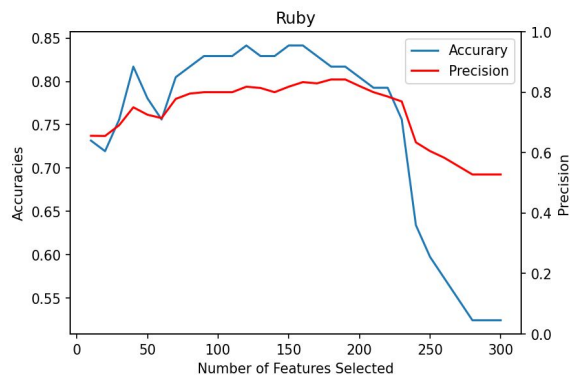
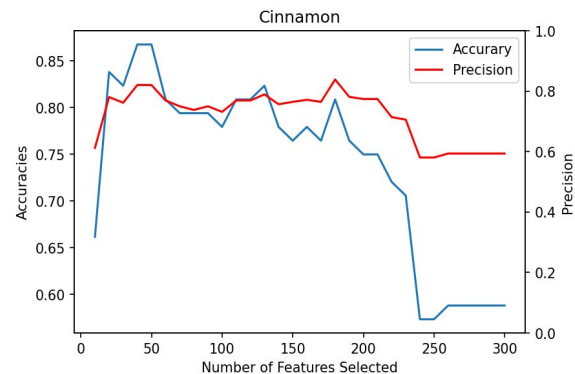
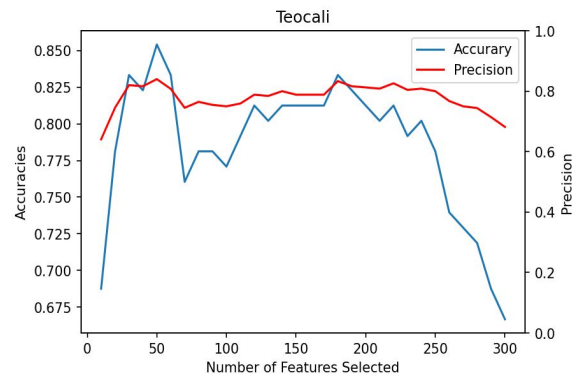
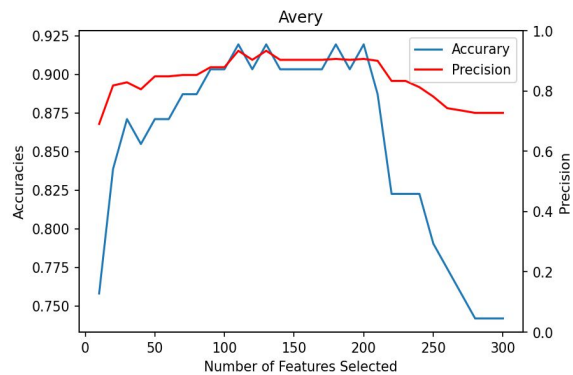


HunterHill

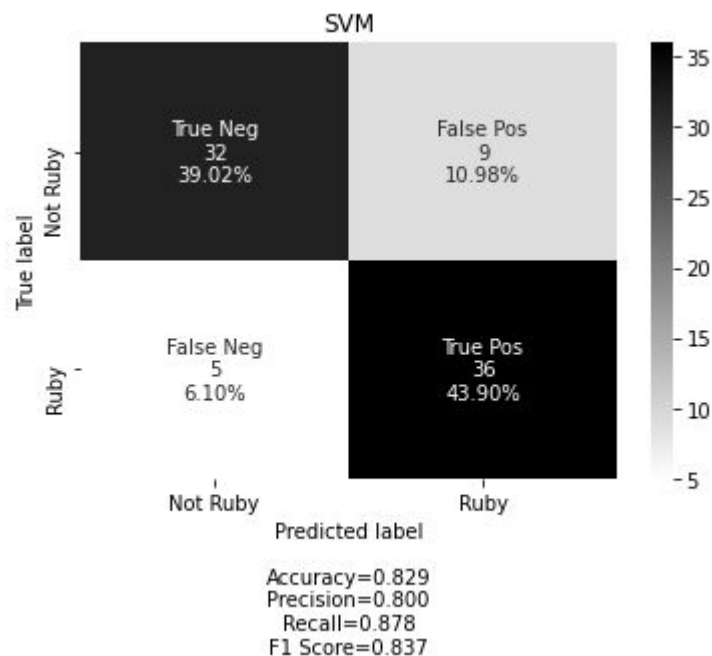


Treasury

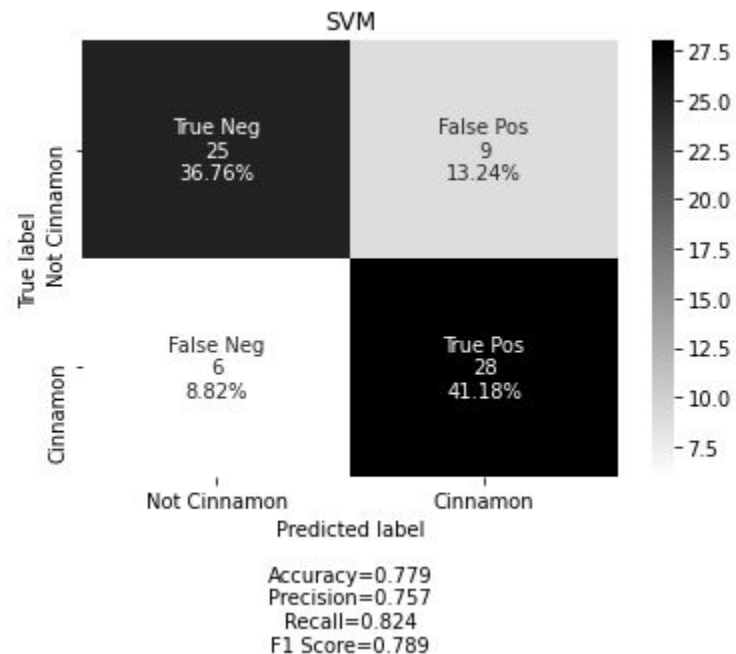




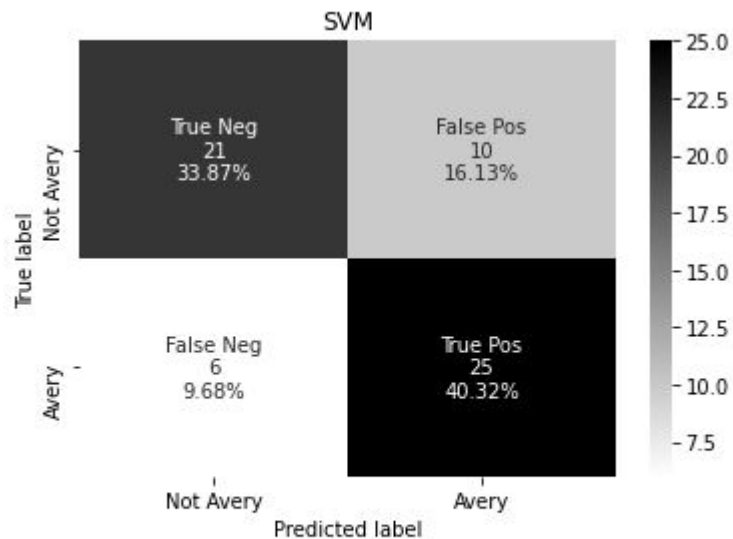
## RUBY



## CINNAMON

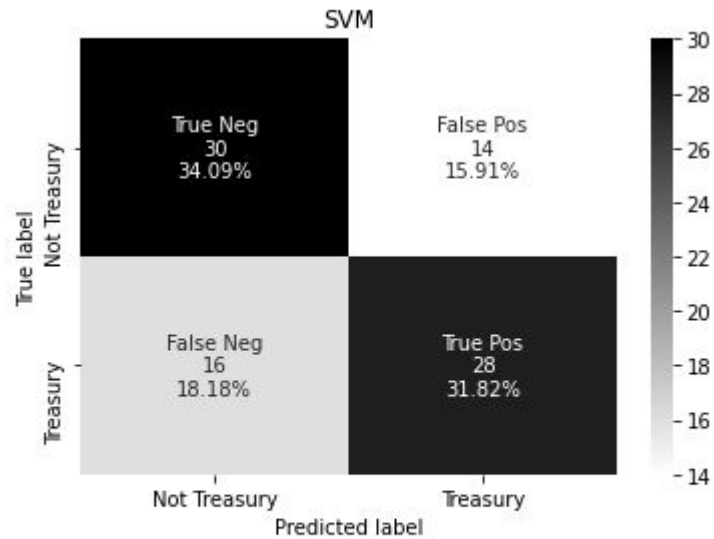


## Avery



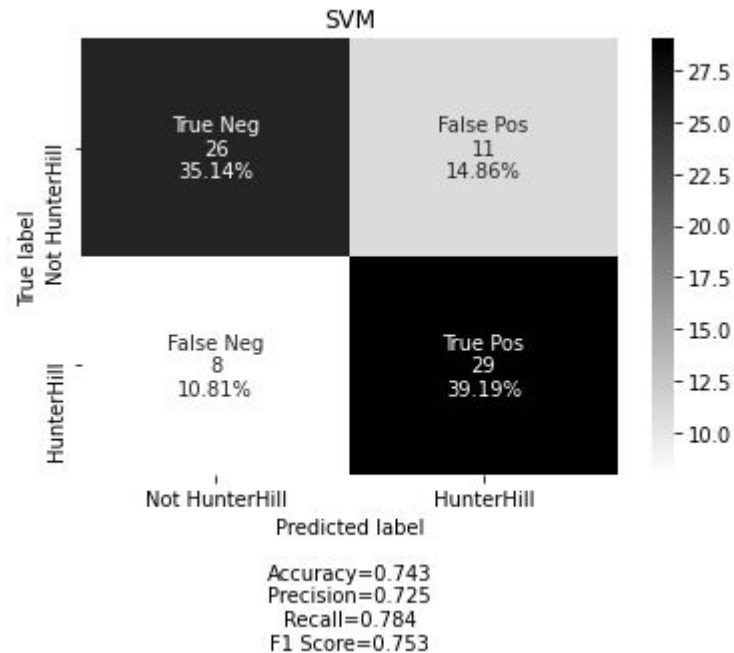
Accuracy=0.742  
Precision=0.714  
Recall=0.806  
F1 Score=0.758

## Treasury

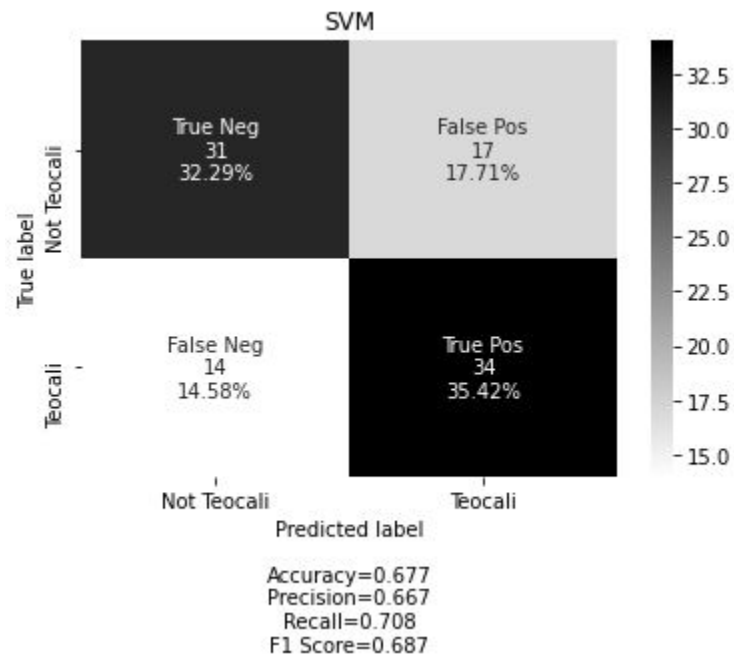


Accuracy=0.659  
Precision=0.667  
Recall=0.636  
F1 Score=0.651

## Hunterhill



## Teocali



# After including 20 percent noise

