Anomaly Detection and JMP[®] Pro

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Outline

- Project goal
- Background
- Imbalanced Data script: dialog, models and sampling techniques, evaluation report
- Real data examples
- Simulations
- Conclusions
- Future work
- References



Project Goals

- To highlight aspects of the imbalanced data problem in the context of classification into a minority and a majority class, where the minority class is under-represented relative to the majority class.
- To provide users with a tool that allows them to explore predictive models that are available in JMP Pro, in conjunction with sampling techniques that are useful in modeling imbalanced data.
- To show examples of the value of the Precision-Recall curve in imbalanced situations.
- To share conclusions about the relative performance of the prediction models and sampling techniques that we studied.
- To provide suggestions about when class imbalance may become an issue for typical modeling techniques.



Background What is the Imbalanced Data Problem?

- Binary response variable
 - # observations at one response level >> # observations at other response level
 - Call the response levels "majority" and "minority"
- Minority level is generally the level of interest
 - Examples include: detection of fraud, disease, credit risk
- Want to predict class membership based on regression variables.
- Some traditional measures of classification accuracy are not appropriate for imbalanced data.



Background Obtaining a Classification Model

- A predictive model that assigns probabilities of membership into the minority class is developed.
- Classification using the predictive model requires selection of a *threshold* value.
- An observation whose predicted probability of membership (or "score") exceeds the threshold value is classified into the minority class.
- Thus, the threshold value defines the classification scheme.
- One tries to choose a threshold value to optimize various criteria, such as the misclassification rate, the true positive rate, the false positive rate, precision, recall, etc.



Background Threshold for Prediction



- A data set consists of 1,452 observations, with only 78 in the minority class.
- The plot shows predictive probabilities of membership in the minority class (thresholds) based on a given model.
- Two thresholds are shown: 0.90 and 0.75.
- Each defines a classification rule.
- As the threshold decreases, more minority instances are identified. But the false positive rate also increases. **Sas**

- For a binary response, one measure of accuracy is the *confusion matrix*.
- It is based on selection of a given threshold.
- The threshold in JMP is 0.5 by default, or you can set a threshold using the Profit Matrix column property.

Confusion Matrix	Predicted Yes	Predicted No		
Actual Yes	True Positive	False Negative		
Actual No	False Positive	True Negative		

- A related summary measure: Accuracy = (TP + TN) / (TP + FP + TN + FN)
- JMP reports: **Misclassification Rate** = 1 Accuracy







- Misclassification rate breaks down with severe imbalance
- Consider the case of a 2% minority class:
 - You can achieve 98% accuracy simply by predicting all majority cases!
 - This would be a bad classifier, however.
- Each threshold value defines a classification scheme and confusion matrix
- Consider curves that plot classification behavior across all thresholds:
 - Precision-Recall Curves
 - Receiver Operating Characteristic (ROC) Curves
 - Gains Curves



• For a given threshold:

Predicted Class								
Actual Class	Count	Minority	Majority	Row Total				
	Minority	ТР	FN	TP + FN = P				
	Majority	FP	TN	FP + TN = N				
	Col Total	TP + FP	TN + FN					

- Sensitivity = True Positive Rate = TP / P
- Specificity = True Negative Rate = TN / N
- 1 Specificity = False Positive Rate = FP / N
- Precision = Positive Predictive Value = TP / (TP + FP)
- Recall = Sensitivity



Background Comparison of Curves

• The PR, ROC, and Cumulative Gains curves are related:

Plot		Y Axis	X Axis		
PR Curve	Precision	True Positives/ (True + False Positives)	Recall	True Positive Rate	
ROC Curve	Sensitivity	True Positive Rate	1 - Specificity	False Positive Rate	
Cumulative Gains Curve	Cumulative Gains	True Positive Rate	Portion	Proportion of Top- Ranked Observations	

• The ideal curve has the Y axis quantity equal to 100%.



Background Precision-Recall Curve



- Precision-Recall (PR) Curve
 - Plots precision versus recall
 - Precision = TP / (TP + FP)
 - Recall = TP / P
- Precision is the Positive Predictive Value
- Recall is the True Positive Rate (Sensitivity)
- The PR curve is preferred for imbalanced data.



Background ROC Curve



• ROC Curve

- Plots sensitivity vs. 1 specificity
- Sensitivity = TP / P
- 1 Specificity = FP / N
- Sensitivity is the True Positive Rate (Recall)
- 1 Specificity is the False Positive Rate



Background Cumulative Gains Curve



- Cumulative Gains Curve
 - Plots cumulative gains vs. portion of the data
 - Cumulative Gains = TP / P (Sensitivity)
 - Portion = proportion of the observations ranked by their probability of membership in the minority class



Background Solutions for Imbalanced Data Problems

- Sampling methods
 - Make modifications to impose a more balanced distribution
- Cost-sensitive methods
 - Use cost to differentiate misclassification consequences or to combine models in an ensemble
 - Incorporate cost information into the classification scheme
- Kernel-based methods
 - Support vector machines (SVMs); can also be integrated with sampling methods



Background Sampling Methods Approaches

- Sampling methods involve modifications to impose a more balanced distribution
 - Random oversampling and undersampling
 - Informed undersampling (EasyEnsemble, BalanceCascade)
 - Synthetic sampling with data generation (SMOTE)
 - Adaptive synthetic sampling (ADA-SYN)
 - Sampling with data cleaning (Tomek links)
 - Cluster-based sampling method
 - Integration of sampling and boosting



Imbalanced Data in JMP

- We want to address imbalanced data sets using JMP Pro.
- How can we implement sampling techniques and combine them with JMP Pro platforms to perform data analysis?
- Chose appropriate JMP Pro platforms.
- Chose a variety of sampling techniques.
- We created a script that enables users to fit and compare models for imbalanced data with a binary response.



Imbalanced Data in JMP

JMP Pro Platforms

- Naïve Bayes
- Neural Networks
 - NTanH(3) Model
- Bootstrap Forest
 - Default options
- Boosted Tree
 - Default options
- Logistic Regression
- Generalized Regression
 - Adaptive Lasso
 - All two-way interactions

Sampling Techniques

- No Weighting
- Weighting
- Random Undersampling
- Random Oversampling
- SMOTE*
- Tomek Links*
- * These techniques are implemented using R.



• No Weighting

- Original data
- Baseline comparison
- Weighting
 - Upweight each observation of the minority class by the same ratio
 - Define the ratio as # majority observations / # minority observations



- Random Undersampling
 - Randomly select a set of observations from the **majority** class
 - Remove this set from the data to decrease the total number of observations
- Random Oversampling
 - Randomly select (with replacement) a set of observations from the **minority** class
 - Add this set to the data to increase the total number of observations

For both methods, the sets are selected such that the sizes of the minority and majority classes are equal.



- Synthetic minority oversampling technique (SMOTE)
 - A more sophisticated form of oversampling adding more minority cases
 - Generates new data observations that are similar to the existing minority class observations, rather than simply replicating them
 - Perform K Nearest Neighbors on the minority class

•
$$x_{new} = x_i + (\hat{x}_i - x_i) + \delta$$

- x_i minority class observation
- \hat{x}_i one of the nearest neighbors for x_i
- δ random number in [0,1]

Figures from He and Garcia (2009; section 3.1)





• Tomek Links

- A more sophisticated form of undersampling removing majority cases
- Removes observations from the majority class that are "close" to minority class observations to better define cluster borders
- Find pairs of nearest neighbors, (x_i, x_j) , that fall into different classes to reduce overlapping of majority and minority instances.
 - x_i in minority class
 - x_j in majority class
 - Remove x_j from data



Imbalanced Data in JMP Dialog Window

Imbalanced Data - IMP Pro

- Choose model and sampling technique combinations
 - For use with SMOTE and Tomek, data are standardized
- Validation options
 - A validation set is used for all fitting options
- Random Seed
 - Sets seed for sampling schemes as well as random validation within platforms
 - Results not identical between JMP 14 and JMP 15 due to changes in random seeds
 - JMP 15 used in this presentation

		5 A
nning on Data Table: Mammogra	phy	
elect Columns	Cast Selected Columns into Roles	Action —
Class attr1 attr2 attr3 attr4 attr5 attr6	Binary Class Variable X, Predictors attr1 attr2 attr3 attr4 attr5 attr6 optional 	OK Cancel Remove Recall
Models Naive Bayes Neural Network Bootstrap Forest Boosted Tree Logistic Regression Generalized Regression Select All Models	Validation Options Training Proportion 0.55 Validation Proportion 0.15 Test Proportion 0.3 Model Options	
ampling Techniques	Set Random Seed 123456	

Note: SMOTE and Tomek are disabled if there is no R connection, if there is only one predictor, or if some predictors are categorical. If there is no R connection, the AUC values for the PR curves are unavailable in the report.



No Weighting

Under Sampling

Select All Techniques

Over Sampling
 SMOTE

Weighting

Tomek

Imbalanced Data in JMP Dialog Window

🙀 Imbalanced Data - JMP Pro		-		×
Running on Data Table: seismic				
Select Columns	Cast Selected Columns into Roles		Action	
seismic seismoacoustic shift genergy gpuls gdenergy gdpuls gdpuls ghazard nbumps	Binary Class Variable th class X, Predictors th shift genergy gpuls gdenergy optional		OI Can Rem	cel ove all
Models Neural Network Bootstrap Forest Boosted Tree Logistic Regression Generalized Regression Select All Models Sampling Techniques No Weighting Weighting	Validation Options Training Proportion 0.55 Validation Proportion 0.15 Test Proportion 0.3 Model Options Set Random Seed .			
Vider Sampling Over Sampling SMOTE Tomek Select All Techniques	Note: SMOTE and Tomek are disabled if there is no R connecti there is only one predictor, or if some predictors are categoric there is no R connection, the AUC values for the PR curves are unavailable in the report.	ion, if al. If		



Imbalanced Data in JMP Dialog Window

🙀 Imbalanced Data - JMP Pro	_	
Running on Data Table: seismic		
Select Columns seismic seismoacoustic shift genergy gpuls gdenergy gdenergy gdpuls gdpuls gdpuls	Cast Selected Columns into Roles Binary Class Variable L class X, Predictors gdpuls optional Image: Class Variable	Action OK Cancel Remove Recall
A nbumps nbumps2 Models Naive Bayes	Validation Options	
Bootstrap Forest Bootstrap Forest Logistic Regression Generalized Regression Select All Models	Training Proportion 0.55 Validation Proportion 0.15 Test Proportion 0.3	
Sampling Techniques	Set Random Seed	
Under Sampling Over Sampling SMOTE Tomek Select All Techniques	Note: SMOTE and Tomek are disabled if there is no R connection, if there is only one predictor, or if some predictors are categorical. If there is no R connection, the AUC values for the PR curves are unavailable in the report.	



Data Sets Studied

- Considered nine data sets.
- Minority class representation runs from 35.90% to 0.17%

	Data Set	N Predictors	N Continuous	N Nominal	N	N Minority Class	Minority %
1	lonosphere	34	32	2	351	126	35.90%
2	Pima Indians	8	8	0	768	268	34.90%
3	Diabetes Modified	10	9	1	442	121	27.38%
4	Ecoli	7	7	0	336	77	22.92%
5	New Thyroid	5	5	0	215	35	16.28%
6	Seismic	18	14	4	2584	170	6.58%
7	Wilt Data	5	5	0	4839	261	5.39%
8	Mammography	6	6	0	11183	260	2.32%
9	Credit Card Fraud	30	30	0	284807	492	0.17%



Data Sets Studied

- The three data sets with the highest minority class percentage showed less difference in terms of classification models and sampling methods than did the other data sets.
- However, even for Diabetes Modified.jmp, with a 27.4% minority proportion, the PR curves differentiate between models, while the ROC curves are similar.





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- The data table Mammography.jmp is based on a set of digitized film mammograms, used in a study of microcalcifications in mammographic images.
- There are six continuous predictors and 11,183 observations.
- Each record is classified as "1", representing calcification, or "0", representing no calcification.
- How might one use the Imbalanced Data script, and the Evaluation Report, to select a model?
- Details are given in the following slides, which are for your reference.





- Run the Imbalanced Data script with your data table of interest as the active data table.
- The script opens the dialog to the right.
- Make appropriate selections and run the script.

💱 Imbalanced Data - JMP Pro	-	-		×
Running on Data Table: Mammog	raphy			
Select Columns	Cast Selected Columns into Roles	_	Actio	n —
Class attr1 attr2 attr3 attr4 attr5 attr6	Binary Class Variable X, Predictors A attr1 attr2 attr3 attr4 attr5 attr6 optional		O Can Rem Rec	K cel ove
Models				
 ✓ Naive Bayes ✓ Neural Network ✓ Bootstrap Forest ✓ Boosted Tree ✓ Logistic Regression ✓ Generalized Regression ✓ Salert All Models 	Validation OptionsTraining Proportion0.55Validation Proportion0.15Test Proportion0.3			
Sampling Techniques	Model OptionsSet Random Seed123456			
 ✓ No Weighting ✓ Weighting ✓ Under Sampling ✓ Over Sampling ✓ SMOTE ✓ Immek 	Note: SMOTE and Tomek are disabled if there is no R connection, if there is only one predictor, or if some predictors are categorical. If there is no R connection, the ALIC values for the PR curves are unavailable in the			



Select All Techniques

report.

Data Sets Studied

- When you run the Imbalanced Data script, the following are provided:
 - The Evaluation Report, called "Imbalanced Data for <current data table>"
 - The Techniques and Thresholds data table, which contains scripts for the Evaluation Report and the Summary Table.
 - The Summary Table
 - The Training Set these are the observations used to fit the models, and they include the validation set selected using the specifications in the dialog
 - The Test Set this is the independent set of observations used to produce the Techniques and Thresholds data table and the Evaluation Report.
- The Techniques and Thresholds table contains the detailed data used to produce the Evaluation Report.
- The Summary Table links to the Techniques and Thresholds table, and thus to the Evaluation Report.



▼ Techniques and Threshol ▷	۹ 🔪 💽							
Source	F	Model Type	Sampling Method	Model and Sampling Method	Probs	Class	Precision	Recall
Summary Table	1	Naive Bayes	No Weighting	Naive No Weighting	1.0000000	•	0.68571	0.000000
Evaluation Report	2	Naive Bayes	No Weighting	Naive No Weighting	1.0000000	1	0.68571	0.307692
Columna (12/0)	3	Naive Bayes	No Weighting	Naive No Weighting	1.0000000	1	0.68571	0.307692
Model Type	4	Naive Bayes	No Weighting	Naive No Weighting	1.0000000	1	0.68571	0.307692
Sampling Method	5	Naive Bayes	No Weighting	Naive No Weighting	1.0000000	1	0.68571	0.307692
Model and Spling Method	6	Naive Bayes	No Weighting	Naive No Weighting	1.0000000	1	0.68571	0.307692
A Probs	7	Naive Bayes	No Weighting	Naive No Weighting	1.0000000	1	0.68571	0.307692
L Class	8	Naive Bayes	No Weighting	Naive No Weighting	1.0000000	1	0.68571	0.307692
Precision Pecell	9	Naive Bayes	No Weighting	Naive No Weighting	1.0000000	0	0.68571	0.307692
A Sensitivity	10	Naive Bayes	No Weighting	Naive No Weighting	1.0000000	1	0.68571	0.307692
1 - Specificity	11	Naive Bayes	No Weighting	Naive No Weighting	1.0000000	1	0.68571	0.307692
Cumulative Gains	12	Naive Bayes	No Weighting	Naive No Weighting	1.0000000	1	0.68571	0.307692
Portion	13	Naive Bayes	No Weighting	Naive No Weighting	1.0000000	1	0.68571	0.307692
ROC Curve AUC	14	Naive Bayes	No Weighting	Naive No Weighting	1.0000000	0	0.68571	0.307692
PR Curve AUC	15	Naive Bayes	No Weighting	Naive No Weighting	1.0000000	1	0.68571	0.307692
	16	Naive Bayes	No Weighting	Naive No Weighting	1.0000000	1	0.68571	0.307692
Kows 120 816	17	Naive Bayes	No Weighting	Naive No Weighting	1.0000000	1	0.68571	0.307692
Selected 0	18	Naive Bayes	No Weighting	Naive No Weighting	1.0000000	0	0.68571	0.307692
Excluded 0	19	Naive Bayes	No Weighting	Naive No Weighting	1.0000000	1	0.68571	0.307692
Hidden 0	20	Naive Bayes	No Weighting	Naive No Weighting	1.0000000	1	0.68571	0.307692
Labelled 0	21	Naive Bayes	No Weighting	Naive No Weighting	1.0000000	0	0.68571	0.307692

 The Techniques and Thresholds data table shows each selected modeling and sampling technique, its probability thresholds, and the computed values that are plotted on the curves.

 Note the Summary Table and Evaluation Report table scripts.



- The Summary table gives AUC values for each selected method.
- It also provides an easy way to select curves for methods in the Evaluation report, or rows in the Techniques and Thresholds table.

💌 Summary Table for Mammography 👂	۹ 🖉 💽				N Non-Missing		
Source		Model Type	Sampling Method	Model and Sampling Method	Probabilities	PR Curve AUC	ROC Curve AUC
Bar Graph Comparison	1	Boosted Tree	No Weighting	BoostedTree No Weighting	3356	0.6002	0.9341
	2	Boosted Tree	Over Sampling	BoostedTree Over Sampling	3356	0.5163	0.9491
	3	Boosted Tree	SMOTE	BoostedTree SMOTE	3356	0.5699	0.9437
Columns (6/0)	4	Boosted Tree	Tomek	BoostedTree Tomek	3356	0.5889	0.9344
🔥 Model Type 🚇	5	Boosted Tree	Under Sampling	BoostedTree Under Sampling	3356	0.5086	0.9462
🔥 Sampling Method	6	Boosted Tree	Weighting	BoostedTree Weighting	3356	0.5221	0.9499
📕 Model and Sampling Method 🚇	7	Bootstrap Forest	No Weighting	BootstrapForest No Weighting	3356	0.5739	0.9461
A DR Currie ALIC @	8	Bootstrap Forest	Over Sampling	BootstrapForest Over Sampling	3356	0.5402	0.9438
	9	Bootstrap Forest	SMOTE	BootstrapForest SMOTE	3356	0.4791	0.9432
	10	Bootstrap Forest	Tomek	BootstrapForest Tomek	3356	0.5360	0.9421
	11	Bootstrap Forest	Under Sampling	BootstrapForest Under Sampl	3356	0.1348	0.8373
Rows	12	Bootstrap Forest	Weighting	BootstrapForest Weighting	3356	0.4908	0.9435
All rows 36	13	Generalized Regr	No Weighting	GenReg No Weighting	3356	0.6022	0.9351
Selected 0	14	Generalized Regr	Over Sampling	GenReg Over Sampling	3356	0.6304	0.9460
Excluded 0	15	Generalized Regr	SMOTE	GenReg SMOTE	3356	0.6397	0.9455
Hidden 0	16	Generalized Regr	Tomek	GenReg Tomek	3356	0.6027	0.9350
	17	Generalized Regr	Under Sampling	GenReg Under Sampling	3356	0.4339	0.9507



Summary			
Report Details			
Model Specifications			
Binary Class Variable	Predictors		
Class	attr 1 attr2 attr3 attr4 attr5 attr6		
Models Naive Bayes Neural Network Bootstrap Forest Boosted Tree Logistic Regression Generalized Regression	Sampling Techniques No Weighting Weighting Under Sampling Over Sampling SMOTE Tomek	Validation Options Training Proportion: 0.55 Validation Proportion: 0.15 Test Proportion: 0.3	Reproducibilit 123456

i rainind Binar Class Variable Number Percent 97.7% 7646 2.3% Total 7828 100.0% Test Binary **Class Variable Number Percent** 0 3277 97.7% 2.3% 78 Total 3355 100.0%

- Run the Evaluation Report script in the Techniques and Thresholds data table to obtain the Evaluation report.
- The Summary outline provides details about the report and information about the analysis that generated the report.
- This outline is followed by the Precision-Reliability Curves, ROC Curves, and Cumulative Gains Curves outlines.





- For the methods and sampling techniques considered, the ROC curves are similar and have high AUC values.
- It is tempting to select Neural No Weighting, or perhaps Neural SMOTE, as the best techniques, as these have the highest AUC values.



• But the ROC curves for Neural No Weighting and Neural SMOTE are very similar. How do you choose between them?







 The PR curves differ substantially for the models considered.





- In particular, the PR curves for Neural No Weighting and Neural SMOTE differ.
- Neural No Weighting has the higher AUC value.





- Suppose you are considering a threshold that gives sensitivity (or recall) around 0.90.
- The Neural No Weighting method gives greater precision than the Neural SMOTE method.





 To see this difference on the ROC curve, you would have to expand the horizontal scale.



- From the Techniques and Thresholds table, we see that Neural No Weighting is more precise at sensitivity 0.897 than Neural SMOTE.
- For Neural No Weighting, of the 8.5% of cases tested, 24.5% are positive.
- For Neural SMOTE, of the 9.9% of cases tested, 21.0% are positive.
- Neural No Weighting gives higher precision with fewer tests than does Neural SMOTE. It follows that Neural No Weighting has a lower false positive rate (1 – Specificity).

Model and Sampling Method	Probs	Class	Precision	Recall	Sensitivity	1 - Specificity	Cumulative Gains	Portion
Neural No Weighting	0.0227600	0	0.24476	0.897436	0.897436	0.065914	0.89744	0.085246
Neural SMOTE	0.0363622	0	0.21021	0.897436	0.897436	0.080256	0.89744	0.099255









• The probabilities of class membership, which define the thresholds, have quite different distributions for the two techniques.

- However, this is not of interest.
- Only the ranking of the thresholds is relevant.



Data Sets Studied Wilt

- Wilt.jmp contains data from a remote sensing study.
- The study involved detecting diseased trees using Quickbird satellite imagery.
- The data set consists of five continuous variable measuring various aspects of image segments.
- The binary response categorizes each image as containing diseased trees or not.
- There are 4,839 images.



Data Sets Studied Wilt



 The model accounts for differences in ROC curves and AUC values, with Naïve Bayes and Bootstrap Forest not performing as well as other models.

- Neural models appear to perform the best.
- Sampling technique has little effect, except for Bootstrap Forest.



Data Sets Studied Wilt



- AUC Values by Model and Sampling Method BoostedTree No Weighting 0.919 BoostedTree Over Sampling 0.935 0.923 BoostedTree SMOTE BoostedTree Tomek 0.93 0.907 BoostedTree Under Sampling BoostedTree Weighting 0.926 BootstrapForest No Weighting 0.758 BootstrapForest Over Sampling 0.464 BootstrapForest SMOTE 0.651 BootstrapForest Tomek 0.817 0.624 BootstrapForest Weighting 0.214 GenReg No Weighting 0.885 0.852 GenReg Over Sampling GenReg SMOTE 0.88 GenReg Tomek 0.885 GenReg Under Sampling 0.784 GenReg Weighting 0.858 Logistic No Weighting 0.744 Logistic Over Sampling 0.734 Logistic SMOTE 0.738 Logistic Tomek 0.743 Logistic Under Sampling 0.637 Logistic Weighting 0.736 Naive No Weighting 0.487 Naive Over Sampling 0.486 0.476 Naive SMOTE Naive Tomek 0.483 0.483 Naive Under Sampling Naive Weighting 0.489 Neural No Weighting 0.967 0.953 Neural Over Sampling Neural SMOTE 0.966 Neural Tomek 0.963 Neural Under Sampling 0.912 Neural Weighting 0.959 0.0 0.2 0.4 0.6 0.8 1.0 1.2 1.4 PR Curve AUC
- Differences are more apparent for PR curves and their AUC values.
- Although model seems to have the largest impact, sampling technique has an effect as well.



Data Sets Studied Credit Card Fraud

0.983

0.981

0.982

0.979

0.983

0.982

0.974

0.98

0.928

0.949

0.982

0.98

0.978

0.98

0.979

0.983

0.984

0.979

0.982

0.983

0.979

0.942

0.944

0.942

0.942

0.943

0.944

0.979

0.971

0.981

0.982

0.975

0.973

0.2 0.4 0.6 0.8 1.0 1.2 1.4

ROC Curve AUC



- The ROC curves and their AUC values show little difference among models, other than for Naïve Bayes.
- The curves and AUC values show virtually no differences among sampling technique, other than for Weighting.



0.0

Data Sets Studied **Credit Card Fraud**

0.825

0.782

0.742

0.724 0.734

0.831

0.84

0.834

0.84

0.843

0.845

0.744

0.8

0.791

0.799

0.771

0.731

0.771

0.752

0.745 0.0 0.2 0.4 0.6 0.8 1.0 1.2 1.4

0.676

PR Curve AUC

0.689

0.599

0.765 0.667

0.714



- The PR curves and their AUC values show major differences both among models and sampling techniques.
- Some models and sampling techniques identify the topscored 85% or so of minority observations with much higher precision than **Sas** others.

Data Sets Studied Assessment of Differences

- As expected, differences between PR and ROC curves are most evident for data sets with a small minority representation.
- For such data sets, PR curves are more informative than ROC curves.

	Data Set	N Predictors	N Continuous	N Nominal	N	Minority %	Informal Assessment of ROC vs PR Curve Differences
1	lonosphere	34	32	2	351	35.90%	Minor
2	Pima Indians	8	8	0	768	34.90%	Some
3	Diabetes Modified	10	9	1	442	27.38%	Some
4	Ecoli	7	7	0	336	22.92%	Major
5	New Thyroid	5	5	0	215	16.28%	Minor
6	Seismic	18	14	4	2584	6.58%	Some, but no models perform well
7	Wilt Data	5	5	0	4839	5.39%	Major
8	Mammography	6	6	0	11183	2.32%	Major
9	Credit Card Fraud	30	30	0	284807	0.17%	Major



Data Sets Studied Minority Proportion < ~0.05 Mammography (2.32%)

BoostedTree No Weighting 0.825 BoostedTree Over Sampling 0.714 BoostedTree SMOTE 0.782 BoostedTree Tomek 0.831 BoostedTree Under Sampling 0.742 BoostedTree Weighting 0.724 BootstrapForest No Weighting 0.734 BootstrapForest Over Sampling 0.84 BootstrapForest SMOTE 0.765 BootstrapForest Tomek 0.667 BootstrapForest Under Sampling 0.198 BootstrapForest Weighting 0.834 GenReg No Weighting 0.84 GenReg Over Sampling 0.744 GenReg SMOTE 0.843 GenReg Tomek 0.845 0.401 GenReg Under Sampling GenReg Weighting 0.00172 Logistic No Weighting 0.8 0.689 Logistic Over Sampling Logistic SMOTE 0.791 Logistic Tomek 0.799 Logistic Under Sampling 0.599 Logistic Weighting 0.00172 Naive No Weighting 0.0766 Naive Over Sampling 0.0687 Naive SMOTE 0.078 Naive Tomek 0.0765 Naive Under Sampling 0.0966 Naive Weighting 0.0686 Neural No Weighting 0.771 Neural Over Sampling 0.731 Neural SMOTE 0.771 Neural Tomek 0.752 Neural Under Sampling 0.676 Neural Weighting 0.745 ____ 0.0 0.2 0.4 0.6 0.8 1.0 1.2 1.4 PR Curve AUC

Model and Sampling Method

Credit Card Fraud (0.17%)



Model and Sampling Method

Wilt (5.39%)

BoostedTree No Weighting	0.919				
BoostedTree Over Sampling	0.935				
BoostedTree SMOTE	0.923				
BoostedTree Tomek	0.93				
BoostedTree Under Sampling	0.907				
BoostedTree Weighting	0.926				
BootstrapForest No Weighting	0.758				
BootstrapForest Over Sampling	0.464				
BootstrapForest SMOTE	0.651				
BootstrapForest Tomek	0.817				
BootstrapForest Under Sampling	0.624				
BootstrapForest Weighting	0.214				
GenReg No Weighting	0.885				
GenReg Over Sampling	0.852				
GenReg SMOTE	0.88				
GenReg Tomek	0.885				
GenReg Under Sampling	0.784				
GenReg Weighting	0.858				
Logistic No Weighting	0,744				
Logistic Over Sampling	0.734				
Logistic SMOTE	0.738				
Logistic Tomek	0.743				
Logistic Under Sampling	0.637				
Logistic Weighting	0.736				
Naive No Weighting	0.487				
Naive Over Sampling	0.486				
Naive SMOTE	0.476				
Naive Tomek	0.483				
Naive Under Sampling	0.483				
Naive Weighting	0.489				
Neural No Weighting	0.967				
Neural Over Sampling	0.953				
Neural SMOTE	0.966				
Neural Tomek	0.963				
Neural Under Sampling	0.912				
Neural Weighting	0.959				
0	0 0.2 0.4 0.6 0.8 1.0 1.2 1.4				
	PR Curve AUC				

Data Driven Simulations

Structure

- Simulations based on two of the studied data sets
 - Mammography and Wilt
- Use the sample size of the data set
 - N = 11,183 in Mammography
 - N = 4,839 in Wilt
- Use the covariance structure of the data set
- Vary the mean vector of the minority class
 - The original mean vector from the data
 - Mean vector that is half the original distance from the majority mean vector
 - Mean vector that is twice the original distance from the majority mean vector
- Vary the proportion of minority class observations
 - Proportion vector (.002, .005, .01, .02, .04, .06, .1, .15, .25, .5)
- Evaluation based on AUC from ROC and PR curves
- 250 iterations for each combination



Simulations Based on Mammography Data



2% minority proportion and original mean vector

Ssas

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Simulations Based on Mammography Data Original mean vector





Simulations Based on Mammography Data Conclusions

- The Boosted Tree, Neural Network, and Naïve Bayes models perform well.
- Undersampling performs poorly for almost all models up to about 10% minority proportion.
- Sometimes no weighting performs better than some of the simpler sampling techniques (weighting, oversampling, and undersampling).
- SMOTE and Tomek consistently perform as well as or better than no weighting.
- There is variation in sampling technique performance for all models except Naïve Bayes.



Simulations Based on Wilt Data



6% minority proportion and original mean vector



Simulations Based on Wilt Data Original mean vector





Simulations Based on Wilt Data Conclusions

- Insights obtained from exploring the data indicate that the minority/majority class overlap in the Wilt data is greater than in the Mammography data.
- The Boosted Tree and Neural Network models perform best.
- There is not much variation in the sampling techniques, except when the distance between means is doubled.



Simulation Study Conclusions

- Undersampling performs poorly compared to other sampling techniques.
 - In simulations based on the Mammography data, it performs poorly for almost all models up to about 10% minority proportion.
 - In simulations based on the Wilt data, it performs poorly for almost all models when the distance between the means is doubled.
- The Boosted Tree and Neural Network models perform the best.
 - Naïve Bayes performs better in simulations based on the Mammography data.
 - Generalized regression performs better in simulations based on the Wilt data.
- There appears to be an interaction between model type and distance between means in their impact on performance.
 - When classes are well separated, logistic and generalized regression perform well, but perform very poorly for classes that overlap.
- Bootstrap Forest has the most variability.



Conclusions

- PR curves highlight differences in sampling methodologies whereas ROC curves tend to mask these differences.
- For highly imbalanced data, PR curves give insight on how to choose a "better" modeling technique – one that gives greater precision for a given true positive rate, thus resulting in fewer false positives.
- The separation between means and the minority proportion have an impact on which models and sampling techniques perform well.
 - We suggest using the Imbalanced Data script whenever the minority proportion is less than 10%.
- The Imbalanced Data script is useful in evaluating and selecting models, whether or not the binary class is imbalanced.



Future Work

- Extend the Imbalanced Data script:
 - Add new models: SVM
 - Add new sampling methods: combined SMOTE/Tomek
 - Allow categorical predictors for SMOTE, Tomek, and SMOTE/Tomek sampling methods.
 - Add model specification options
 - Generalized Regression: validation and estimation methods
 - Tree models: tree and resampling specification options
 - Neural nets: multiple hidden layers, boosting
- Study cases where there are more predictors than observations (n < p)



Possible Simulation Study Extensions

- Use different covariance structures.
- Standardize the distances between means.
- Explore the impact of dimensionality.
- Explore model specifications and model options for a specific class of models, perhaps Gen Reg.

Be able to better answer the question: "At what point are my data so imbalanced that I need to worry about the imbalance?"



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