MLAssignment1

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1 Introduction

In this study, we implemented the XGBoostTree class in Java for the Weka framework. This class was then utilized within the provided XGBoost class, aiming to facilitate to construct a comprehensive XGBoost learner in Weka.

The core principle of XGBoost is using the Taylor expansion of the loss function, specifically first and second derivative, to refine model predictions in each iteration, seeking to minimize the overall objective function. XGBoost trees incorporate regularization to reduce the risk of overfitting. The tree structures are optimized by gradient boosting framework and evaluate splits using the 'Gain'.

In this study the primary output is the algorithm module designed for integration with Weka and Java.

To evaluate the implemented XGBoost tree learning algorithm, we will compare it with the original XGBoost by performing experiments using Weka experimenter and then analyse the capabilities and potential refinements.

2 Algorithm and Implementation

In the XGBoost class, it is already implemented that choosing between Logloss and SquaredError based on the nominal/numeric class. First and second derivatives of the loss function are computed. Additionally, for the logistic type, a transformation using the *Sigmoid* function is performed:

$$\sigma(x) = \frac{1}{1 + e^{-x}}$$

In the XGBoostTree class, the first and second-order derivatives, represented by g and h respectively, are received and then integrated into the tree's decision calculations.

The primary goal of boosting ensemble algorithms is to minimize a given objective function. This can be represented as:

$$Obi(\Theta) = L(\Theta) + \Omega(\Theta)$$

XGBoost uses a Taylor expansion up to the second order to handle this loss function, approximating it as:

$$\sum_{i} \left[L(y_i, y_{iK-1}) + g_i f_K(x_i) + \frac{1}{2} h_i f_K^2(x_i) \right]$$

In the tree, there is the following mapping relationship:

$$f_K(x) = w_{q(x)}$$

where q(x) is the index of the leaf node (numbered from left to right). w represents the value of the leaf node.

The objective function can ultimately be transformed to:

Obj =
$$\sum_{j=1}^{T} \left[G_j w_j + \frac{1}{2} (H_j + \lambda) w_j^2 \right] + \gamma T$$

To find the optimal w_j , we can differentiate the objective equation with respect to w_j and set it to zero:

$$\frac{\partial \text{Obj}}{\partial w_j} = G_j + (H_j + \lambda)w_j = 0$$

From which:

$$w_j^* = -\frac{G_j}{H_j + \lambda}$$

This gives us the optimal weight for a leaf.

For the Gain, it is used to measure the reduction in loss:

Gain =
$$\frac{1}{2} \left[\frac{G_L^2}{H_L + \lambda} + \frac{G_R^2}{H_R + \lambda} - \frac{(G_L + G_R)^2}{(H_L + H_R) + \lambda} \right] - \gamma$$

In the XGBoostTree class specifically, the workflow can be found in the appendix.

In XGBoost, we also introduced a random seed and used it in XGBoostTree class, referred to as seed, to ensure consistent randomness across multiple runs.

3 Experimental Results and Discussion

3.1 Experimental Procedure and Results

In the Weka experimenter, we conducted comparative experiments with the official XGBoost using 10-fold cross-validation run 10 times across all classification and regression datasets. We then examined the classification accuracy at a significance level of 0.05 and compared the root relative squared error in both classification and regression. The results are shown in the table below:

Percent_correct Analysis for classification

Tester: weka.experiment.PairedCorrectedTTester -G 4,5,6 -D 1 -R 2 -S 0.05 -result-matrix "weka.experiment.ResultMatrixPlainText -mean-prec 2 -stddev-prec 2 -col-name-width 0 -row-name-width 25 -mean-width 0 -stddev-width 0 -sig-width 0 -count-width 5 -print-col-names -print-row-names -enum-col-names"

Analysing: Percent_correct

Datasets: 14 Resultsets: 2

Confidence: 0.05 (two tailed)

Sorted by: -

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Dataset	(1) sklearn.S	(2) meta.X
balance-scale-weka.filter	95.57	73.55 *
wisconsin-breast-cancer-w	96.61	95.59
pima_diabetes-weka.filter	73.11	65.93 *
ecoli-weka.filters.unsupe	95.86	93.45 *
Glass-weka.filters.unsupe	83.02	69.61 *
ionosphere-weka.filters.u	92.99	88.46 *
iris-weka.filters.unsuper	100.00	100.00
optdigits-weka.filters.un	99.21	97.90 *
pendigits-weka.filters.un	99.68	98.60 *
segment-weka.filters.unsu	99.69	98.78 *
sonar-weka.filters.unsupe	83.85	76.24 *
vehicle-weka.filters.unsu	98.38	96.31 *
vowel-weka.filters.unsup	99.25	93.41 *
waveform-weka.filters.uns	88.63	77.83 *

Key:

Percent_correct Analysis for classification(only recorded differences)

 $Analysing: Root_relative_squared_error$

Datasets: 14 Resultsets: 2

 $^{(1) \} sklearn. Scikit Learn Classifier \ \verb|'-batch 100 -learner XGBC | lassifier -parameters "gamma=1, subsample=0.5, colsample_bynode=0.5, tree_method="exact" | lassifier -parameters | lassifier$

[&]quot; -py-command python -py-path default -server-name none' -6212485658537766441

 $^{(2)\, {\}tt meta.XGBoost}$ '-S 1 -I 100 -W trees.XGBoostTree -- -S 1 -colsample_bynode 0.5 -eta 0.3 -gamma 1.0 -lambda 1.0 -max_depth 6 -min_child_weight 1.0 -subsample 0.5' -862299050221542512

Dataset	(1) sklearn.S	(2) meta.X
2dplanes	(100) 23.24	95.64 v
bank8FM	$(100) \ 31.10$	99.99 v
cpu_act	(100) 13.74	15.71 v
cpu_small	(100) 15.81	17.48 v
$delta_ailerons$	(100) 100.00	100.00
$delta_elevators$	$(100)\ 100.00$	100.00
$diabetes_numeric$	(100) 90.46	95.94
kin8nm	(100) 59.83	82.81 v
$machine_cpu$	(100) 34.79	53.23 v
puma8NH	(100) 63.11	97.14 v
pyrim	(100) 99.81	102.26
stock	(100) 14.14	26.18 v
triazines	(100) 100.21	100.56
wisconsin	(100) 113.15	109.67
	(v/ /*)	(8/6/0)

3.2 Experimental Observations and Reflections

- 1. The self-implemented XGBoost is significantly slower, especially on the 2dplane dataset. The Elapsed-Time training is 0.76 to 21.92. A probable reason could be that the my XGBoost has not optimized sorting. For every iteration, it uses Utils.sortWithNoMissingValues to sort all training instances anew. This hypothesis needs further validation. Given time, it might be better to employ sorting methods learned in the course to reduce the time complexity.
- 2. Regarding classification accuracy and root relative squared error metrics, the performance on some datasets is noticeably inferior compared to the official XGBoost, while on other datasets, it's nearly equivalent. Overall, there is a performance gap between the self-implemented XGBoost and the official one both in classification accuracy and root relative squared error.

4 Appendix A: XGBoostTree Class Flowchart

