

On Building A Data Fitting System Using Ad Hoc Models

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ABSTRACT

One class of data is measured or simulated data with error estimation. This data can consist of many continuous dimensions for which values are available only at discrete points. Increasing the number of discrete points at which the data is available can be expensive or even impossible to obtain, but it can still be useful for predicting data trends. Unfortunately, this is difficult when the various dimensions do not follow the same type of fit (linear, logarithmic, polynomial, etc.). Our approach focuses on building decision trees and using them to interpolate new data points that follow existing trends. This is in contrast to previous methods which focused on extrapolating data for specific applications or using purely numerical regression models. By using this approach, sparse data sets or those that exhibit unusual patterns can be analyzed effectively.

Categories and Subject Descriptors

H.2.8 [Database Management]: Database Applications—*data mining*

General Terms

Algorithms

Keywords

data mining, sparse data, interpolation

1. INTRODUCTION

Outline goes here.

- The first item
- The second item
- The third etc ...

1.1 Stuff

This is a subsection.

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1.1.1 More Stuff

This is a subsubsection.

2. RELATED WORK

Another section. I am citing something random [1].

3. METHODOLOGY

3.1 Decision Tree Interpolation

Decision Tree Interpolation follows this process:

1. Build a decision tree from the original data.
2. For each leaf node in the tree:
 - (a) Obtain all attribute values for associated data instances.
 - (b) Define ranges for attribute values.
 - For numeric attributes, define the range using minimum and maximum values.
 - For discrete attributes, define the range as all distinct values.
 - (c) Calculate the distribution of all associated data instances.
 - (d) Create new data points within the ranges that match the statistical distribution.

Note that the number of data points created per leaf node is proportional to the number of data points already classified by that leaf node. This ensures that any interpolated data will follow the overall data distribution, at least relative to the data density per leaf node.

3.2 Interpolated Data Validation

All interpolated data is validated through this process:

1. A new decision tree is built based on the interpolated data. Note that the original data is *not* included here.
2. Both the new and original decision trees are compared for accuracy against the new and original data sets.

Note that any decision tree with an arbitrarily large maximum depth can classify data with perfect accuracy. Defining a low maximum depth means that classification is imperfect, and it is under these conditions that differences in the quality of different decision trees become apparent.

Table 1: Experiment Result Summary

Data Set	Max Tree Depth	OT -> OD	OT -> ND	NT -> OD	NT -> ND
adult_sample	1	0.805527123849	0.780737704918	0.804503582395	0.782786885246
adult_sample	2	0.808597748209	0.809426229508	0.249744114637	0.813524590164
adult_sample	3	0.816786079836	0.850102669405	0.801432958035	0.852156057495
adult_sample	4	0.822927328557	0.794661190965	0.787103377687	0.784394250513
adult_sample	5	0.822927328557	0.84052532833	0.792221084954	0.84052532833
car	1	0.700231481481	0.710648148148	0.700231481481	0.710648148148
car	2	0.777777777778	0.783564814815	0.774305555556	0.789351851852
car	3	0.824074074074	0.809027777778	0.824074074074	0.815972222222
car	4	0.894097222222	0.903935185185	0.889467592593	0.915509259259
car	5	0.96412037037	0.966981132075	0.938078703704	0.982311320755
iris	1	0.666666666667	0.693333333333	0.666666666667	0.693333333333
iris	2	0.96	0.973333333333	0.946666666667	0.986666666667
iris	3	0.973333333333	0.959459459459	0.953333333333	0.972972972973
iris	4	0.98	0.959459459459	0.946666666667	1.0
iris	5	1.0	1.0	0.966666666667	1.0
lung-cancer	1	0.59375	0.6	0.375	0.666666666667
lung-cancer	2	0.625	0.571428571429	0.4375	0.714285714286
lung-cancer	3	0.625	0.642857142857	0.5625	1.0
lung-cancer	4	0.6875	0.428571428571	0.53125	1.0
lung-cancer	5	0.78125	0.538461538462	0.53125	1.0
tic_tac_toe	1	0.699373695198	0.68267223382	0.699373695198	0.68267223382
tic_tac_toe	2	0.705636743215	0.690376569038	0.703549060543	0.696652719665
tic_tac_toe	3	0.769311064718	0.779874213836	0.755741127349	0.758909853249
tic_tac_toe	4	0.831941544885	0.82264957265	0.745302713987	0.856837606838
tic_tac_toe	5	0.918580375783	0.907284768212	0.83611691023	0.933774834437
voting	1	0.95632183908	0.923766816143	0.95632183908	0.923766816143
voting	2	0.95632183908	0.956896551724	0.95632183908	0.956896551724
voting	3	0.963218390805	0.913357400722	0.95632183908	0.927797833935
voting	4	0.963218390805	0.892156862745	0.928735632184	0.90522875817
voting	5	0.972413793103	0.866071428571	0.937931034483	0.895833333333

3.3 Experiment Parameters

We completed 30 experiments based on two variables: data source and maximum depth of the decision tree. The maximum depth ranged from one to five, and there were six data sources pulled from Orange’s documentation data sets.

4. RESULTS

Result data is shown in Table 1. The first two columns list the data source and the maximum depth for generated decision trees. Note that two decision trees are generated per row, one for the original data and one for the new (interpolated) data. The last four columns list the accuracy of both the original and new decision trees against the original and new data sets. For example, the column labeled “OT -> ND” shows the accuracy of the original decision tree (OT) when used to classify instances in the new data set (ND). Results and outliers for each of the data sets will be explored further here.

4.1 adult_sample

The only significant outlier in this group was for a max tree depth of two. The “NT -> OD” column shows an accuracy of 0.2497 while the other accuracy values ranged from 0.8085 to 0.8135. Ignoring the outlier, accuracy ranged from 0.7807 to 0.8522 for this data source.

4.2 car

Results in this section largely followed expected trends.

Accuracy ranged from 0.7002 to 0.9823 for this data source, and accuracy increased in each column as the max tree depth increased.

4.3 iris

Results in this section followed expectations, but classification started out relatively inaccurate. At a max tree depth of 1, accuracy ranged from 0.6667 to 0.6933. At a max tree depth of 2, accuracy increased rapidly to range from 0.9467 to 0.9867. Accuracy remained above 0.9467 for all other entries.

4.4 lung-cancer

Accuracy had enormous variance here, ranging from 0.375 to 1.0 across all categories. While both decision trees increased in accuracy as max depth increased for their own data sets, they did not consistently increase in accuracy for the other data set.

4.5 tic_tac_toe

Results in this section were mostly consistent, and accuracy ranged from 0.6827 to 0.9338. Accuracy tended to increase as max tree depth increased, with only slight exceptions.

4.6 voting

While there was no clear trend in the results for this data source, accuracy was consistently greater than or equal to 0.8661 for all entries. All but three values were above 0.9,

so accuracy was generally high across all instances.

5. DISCUSSION

Another section.

6. CONCLUSION AND FUTURE WORK

Last section.

7. REFERENCES

- [1] M. Bowman, S. K. Debray, and L. L. Peterson.
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