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Data Preparation for Gradient Boosting with XGBoost in Python

by Jason Brownlee on August 22, 2016 in XGBoost



XGBoost is a popular implementation of Gradient Boosting because of its speed and performance.

Internally, XGBoost models represent all problems as a regression predictive modeling problem that only takes numerical values as input. If your data is in a different form, it must be prepared into the expected format.

In this post, you will discover how to prepare your data for using with gradient boosting with the XGBoost library in Python.

After reading this post you will know:

- How to encode string output variables for classification.
- How to prepare categorical input variables using one hot encoding.
- How to automatically handle missing da

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Let's get started.

- **Update Sept/2016**: I updated a few small typos in the impute example.
- Update Jan/2017: Updated to reflect changes in scikit-learn API version 0.18.1.
- Update Jan/2017: Updated breast cancer example to converted input data to strings.



Data Preparation for Gradient Boosting with XGBoost in Python Photo by Ed Dunens, some rights reserved.

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Label Encode String Class Values

The iris flowers classification problem is an example of a problem that has a string class value.

This is a prediction problem where given measurements of iris flowers in centimeters, the task is to predict to which species a given flower belongs.

Below is a sample of the raw dataset. You can learn more about this dataset and download the raw data in CSV format from the UCI Machine Learning Repository.

```
1 5.1,3.5,1.4,0.2,Iris-setosa
2 4.9,3.0,1.4,0.2,Iris-setosa
3 4.7,3.2,1.3,0.2,Iris-setosa
4 4.6,3.1,1.5,0.2,Iris-setosa
5 5.0,3.6,1.4,0.2,Iris-setosa
```

XGBoost cannot model this problem as-is as it requires that the output variables be numeric.

We can easily convert the string values to integer values using the LabelEncoder. The three class values (Iris-setosa, Iris-versicolor, Iris-virginica) are mapped to the integer values (0, 1, 2).

```
1 # encode string class values as integers
2 label_encoder = LabelEncoder()
3 label_encoder = label_encoder.fit(Y)
4 label_encoded_y = label_encoder.transform(Y)
```

We save the label encoder as a separate object so that we can transform both the training and later the test and validation datasets using the same encoding scheme.

Below is a complete example demonstrating how to load the iris dataset. Notice that Pandas is used to load the data in order to handle the string class values.

```
1 # multiclass classification
2 import pandas
3 import xgboost
4 from sklearn import model_selection
5 from sklearn.metrics import accuracy_score
6 from sklearn.preprocessing import LabelEncoder
7 # load data
8 data = pandas.read_csv('iris.csv', header=None)
9 dataset = data.values
10 # split data into X and y
11 X = dataset[:,0:4]
12 Y = dataset[:,4]
13 # encode string class values as integers
14 label_encoder = LabelEncoder()
15 label_encoder = label_encoder.fit(Y)
16 label_encoded_y = label_encoder.transform(Y)
17 \text{ seed} = 7
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18 test_size = 0.33
```

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```
19 X_train, X_test, y_train, y_test = cross_validation.train_test_split(X, label_encoded]
20 # fit model no training data
21 model = xgboost.XGBClassifier()
22 model.fit(X_train, y_train)
23 print(model)
24 # make predictions for test data
25 y_pred = model.predict(X_test)
26 predictions = [round(value) for value in y_pred]
27 # evaluate predictions
28 accuracy = accuracy_score(y_test, predictions)
29 print("Accuracy: %.2f%" % (accuracy * 100.0))
```

Running the example produces the following output:

Notice how the XGBoost model is configured to automatically model the multiclass classification problem using the **multi:softprob** objective, a variation on the softmax loss function to model class probabilities. This suggests that internally, that the output class is converted into a one hot type encoding automatically.

One Hot Encode Categorical Data

Some datasets only contain categorical data, for example the breast cancer dataset.

This dataset describes the technical details of breast cancer biopsies and the prediction task is to predict whether or not the patient has a recurrence of cancer, or not.

Below is a sample of the raw dataset. You can learn more about this dataset at the UCI Machine Learning Repository and download it in CSV format from mldata.org.

```
1 '40-49','premeno','15-19','0-2','yes','3','right','left_up','no','recurrence-events'
2 '50-59','ge40','15-19','0-2','no','1','right','central','no','no-recurrence-events'
3 '50-59','ge40','35-39','0-2','no','2','left','left_low','no','recurrence-events'
4 '40-49','premeno','35-39','0-2','yes','3','right','left_low','yes','no-recurrence-event
5 '40-49','premeno','30-34','3-5','yes','2','left','right_up','no','recurrence-events'
```

We can see that all 9 input variables are categorical and described in string format. The problem is a binary classification prediction problem and the output class values are also described in string format.

We can reuse the same approach from the previous section and convert the string class values to integer values to model the prediction using

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```
1 # encode string class values as integers
```

We can use this same approach on each input feature in X, but this is only a starting point.

```
1 # encode string input values as integers
2 features = []
3 for i in range(0, X.shape[1]):
4    label_encoder = LabelEncoder()
5    feature = label_encoder.fit_transform(X[:,i])
6    features.append(feature)
7 encoded_x = numpy.array(features)
8 encoded_x = encoded_x.reshape(X.shape[0], X.shape[1])
```

XGBoost may assume that encoded integer values for each input variable have an ordinal relationship. For example that 'left-up' encoded as 0 and 'left-low' encoded as 1 for the breast-quad variable have a meaningful relationship as integers. In this case, this assumption is untrue.

Instead, we must map these integer values onto new binary variables, one new variable for each categorical value.

For example, the breast-quad variable has the values:

```
1 left-up
2 left-low
3 right-up
4 right-low
5 central
```

We can model this as 5 binary variables as follows:

```
1 left-up, left-low, right-up, right-low, central
2 1,0,0,0,0
3 0,1,0,0,0
4 0,0,1,0,0
5 0,0,0,1,0
6 0,0,0,0,1
```

This is called one hot encoding. We can one hot encode all of the categorical input variables using the OneHotEncoder class in scikit-learn.

We can one hot encode each feature after we have label encoded it. First we must transform the feature array into a 2-dimensional NumPy array where each integer value is a feature vector with a length 1.

```
1 feature = feature.reshape(X.shape[0], 1)
```

We can then create the OneHotEncoder and Get Your Start in Machine Learning

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```
1 onehot_encoder = OneHotEncoder(sparse=False)
2 feature = onehot_encoder.fit_transform(feature)
```

Finally, we can build up the input dataset by concatenating the one hot encoded features, one by one, adding them on as new columns (axis=2). We end up with an input vector comprised of 43 binary input variables.

```
# encode string input values as integers
   encoded_x = None
3 for i in range(0, X.shape[1]):
4
       label_encoder = LabelEncoder()
5
       feature = label_encoder.fit_transform(X[:,i])
6
       feature = feature.reshape(X.shape[0], 1)
7
       onehot_encoder = OneHotEncoder(sparse=False)
8
       feature = onehot_encoder.fit_transform(feature)
9
       if encoded_x is None:
10
           encoded_x = feature
11
       else:
12
           encoded_x = numpy.concatenate((encoded_x, feature), axis=1)
13 print("X shape: : ", encoded_x.shape)
```

Ideally, we may experiment with not one hot encode some of input attributes as we could encode them with an explicit ordinal relationship, for example the first column age with values like '40-49' and '50-59'. This is left as an exercise, if you are interested in extending this example.

Below is the complete example with label and one hot encoded input variables and label encoded output variable.

```
# binary classification, breast cancer dataset, label and one hot encoded
2 import numpy
3 from pandas import read_csv
4 from xgboost import XGBClassifier
5 from sklearn.model_selection import train_test_split
6 from sklearn.metrics import accuracy_score
7 from sklearn.preprocessing import LabelEncoder
8 from sklearn.preprocessing import OneHotEncoder
9 # load data
10 data = read_csv('datasets-uci-breast-cancer.csv', header=None)
11 dataset = data.values
12 # split data into X and y
13 X = dataset[:,0:9]
14 X = X.astype(str)
15 Y = dataset[:,9]
16 # encode string input values as integers
17 \text{ encoded}_x = \text{None}
18 for i in range(0, X.shape[1]):
19
       label_encoder = LabelEncoder()
20
       feature = label_encoder.fit_transform(X[:,i])
21
       feature = feature.reshape(X.shape[0], 1)
22
       onehot_encoder = OneHotEncoder(sparse=False)
23
       feature = onehot_encoder.fit_transform(feature)
24
       if encoded_x is None:
25
           encoded_x = feature
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26
       else:
```

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```
encoded_x = numpy.concatenate((encoded_x, feature), axis=1)
28 print("X shape: : ", encoded_x.shape)
29 # encode string class values as integers
30 label_encoder = LabelEncoder()
31 label_encoder = label_encoder.fit(Y)
32 label_encoded_y = label_encoder.transform(Y)
33 # split data into train and test sets
34 \text{ seed} = 7
35 test_size = 0.33
36 X_train, X_test, y_train, y_test = train_test_split(encoded_x, label_encoded_y, test_\u00e9
37 # fit model no training data
38 model = XGBClassifier()
39 model.fit(X_train, y_train)
40 print(model)
41 # make predictions for test data
42 y_pred = model.predict(X_test)
43 predictions = [round(value) for value in y_pred]
44 # evaluate predictions
45 accuracy = accuracy_score(y_test, predictions)
46 print("Accuracy: %.2f%%" % (accuracy * 100.0))
```

Running this example we get the following output:

Again we can see that the XGBoost framework chose the 'binary:logistic' objective automatically, the right objective for this binary classification problem.

Support for Missing Data

XGBoost can automatically learn how to best handle missing data.

In fact, XGBoost was designed to work with sparse data, like the one hot encoded data from the previous section, and missing data is handled the same way that sparse or zero values are handled, by minimizing the loss function.

For more information on the technical details for how missing values are handled in XGBoost, see Section 3.4 "Sparsity-aware Split Finding" in the paper XGBoost: A Scalable Tree Boosting System.

The Horse Colic dataset is a good example to demonstrate this capability as it contains a large percentage of missing data, approximately 30%.

You can learn more about the Horse Colic d

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Machine Learning repository.

The values are separated by whitespace and we can easily load it using the Pandas function read_csv.

```
1 dataframe = read_csv("horse-colic.csv", delim_whitespace=True, header=None)
```

Once loaded, we can see that the missing data is marked with a question mark character ('?'). We can change these missing values to the sparse value expected by XGBoost which is the value zero (0).

```
1 # set missing values to 0
2 X[X == '?'] = 0
```

Because the missing data was marked as strings, those columns with missing data were all loaded as string data types. We can now convert the entire set of input data to numerical values.

```
1 # convert to numeric
2 X = X.astype('float32')
```

Finally, this is a binary classification problem although the class values are marked with the integers 1 and 2. We model binary classification problems in XGBoost as logistic 0 and 1 values. We can easily convert the Y dataset to 0 and 1 integers using the LabelEncoder, as we did in the iris flowers example.

```
1 # encode Y class values as integers
2 label_encoder = LabelEncoder()
3 label_encoder = label_encoder.fit(Y)
4 label_encoded_y = label_encoder.transform(Y)
```

The full code listing is provided below for completeness.

```
1 # binary classification, missing data
2 from pandas import read_csv
3 from xgboost import XGBClassifier
4 from sklearn.model_selection import train_test_split
5 from sklearn.metrics import accuracy_score
6 from sklearn.preprocessing import LabelEncoder
7 # load data
8 dataframe = read_csv("horse-colic.csv", delim_whitespace=True, header=None)
9 dataset = dataframe.values
10 # split data into X and y
11 X = dataset[:,0:27]
12 Y = dataset[:,27]
13 # set missing values to 0
14 X[X == '?'] = 0
15 # convert to numeric
16 X = X.astype('float32')
17 # encode Y class values as integers
18 label_encoder = LabelEncoder()
                                          Get Your Start in Machine Learning
19 label_encoder = label_encoder.fit(Y)
```

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```
label_encoded_y = label_encoder.transform(Y)

1  # split data into train and test sets

2  seed = 7

2  test_size = 0.33

24  X_train, X_test, y_train, y_test = train_test_split(X, label_encoded_y, test_size=test)

5  # fit model no training data

6  model = XGBClassifier()

7  model.fit(X_train, y_train)

8  print(model)

9  # make predictions for test data

30  y_pred = model.predict(X_test)

31  predictions = [round(value) for value in y_pred]

32  # evaluate predictions

33  accuracy = accuracy_score(y_test, predictions)

34  print("Accuracy: %.2f%%" % (accuracy * 100.0))
```

Running this example produces the following output.

We can tease out the effect of XGBoost's automatic handling of missing values, by marking the missing values with a non-zero value, such as 1.

```
1 X[X == '?'] = 1
```

Re-running the example demonstrates a drop in accuracy for the model.

```
1 Accuracy: 79.80%
```

We can also impute the missing data with a specific value.

It is common to use a mean or a median for the column. We can easily impute the missing data using the scikit-learn Imputer class.

```
1 # impute missing values as the mean
2 imputer = Imputer()
3 imputed_x = imputer.fit_transform(X)
```

Below is the full example with missing data imputed with the mean value from each column.

```
1  # binary classification, missing data, impute with mean
2  import numpy
3  from pandas import read_csv
4  from xgboost import XGBClassifier
5  from sklearn.model_selection import train_test_split
6  from sklearn.metrics import accuracy_score
7  from sklearn.preprocessing import Lah
8  from sklearn.preprocessing import Imp
```

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```
10 dataframe = read_csv("horse-colic.csv", delim_whitespace=True, header=None)
11 dataset = dataframe.values
12 # split data into X and y
13 X = dataset[:,0:27]
14 Y = dataset[:,27]
15 # set missing values to 0
16 X[X == '?'] = numpy.nan
17 # convert to numeric
18 X = X.astype('float32')
19 # impute missing values as the mean
20 imputer = Imputer()
21 imputed_x = imputer.fit_transform(X)
22 # encode Y class values as integers
23 label_encoder = LabelEncoder()
24 label_encoder = label_encoder.fit(Y)
25 label_encoded_y = label_encoder.transform(Y)
26 # split data into train and test sets
27 \text{ seed} = 7
28 test_size = 0.33
29 X_train, X_test, y_train, y_test = train_test_split(imputed_x, label_encoded_y, test_s
30 # fit model no training data
31 model = XGBClassifier()
32 model.fit(X_train, y_train)
33 print(model)
34 # make predictions for test data
35 y_pred = model.predict(X_test)
36 predictions = [round(value) for value in y_pred]
37 # evaluate predictions
38 accuracy = accuracy_score(y_test, predictions)
39 print("Accuracy: %.2f%%" % (accuracy * 100.0))
```

Running this example we see results equivalent to the fixing the value to one (1). This suggests that at least in this case we are better off marking the missing values with a distinct value of zero (0) rather than a valid value (1) or an imputed value.

```
1 Accuracy: 79.80%
```

It is a good lesson to try both approaches (automatic handling and imputing) on your data when you have missing values.

Summary

In this post you discovered how you can prepare your machine learning data for gradient boosting with XGBoost in Python.

Specifically, you learned:

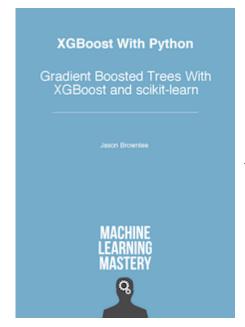
- How to prepare string class values for binary classification using label encoding.
- How to prepare categorical input variables using a one hot encoding to model them as binary variables.
- How XGBoost automatically handles mi
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missing values.

Do you have any questions about how to prepare your data for XGBoost or about this post? Ask your questions in the comments and I will do my best to answer.

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About Jason Brownlee

Dr. Jason Brownlee is a husband, proud father, academic researcher, author, professional developer and a machine learning practitioner. He is dedicated to helping developers get started and get good at applied machine learning. Learn more.

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20 Responses to Data Preparation for Gradient Boosting with XGBoost in Python



Ralph_adu August 28, 2016 at 1:24 am #

REPLY 🦴

hi Jason, the train data for the last example should be imputed_x, but you use the original X which has missing data. I tried with data imputed_x and get the accuracy 79.8%



Jason Brownlee September 12, 2016 at 2:35 pm #

REPLY 🦴

Thanks Ralph. Fixed.



Qichang September 19, 2016 at 9:29 am #

REPLY 🦴

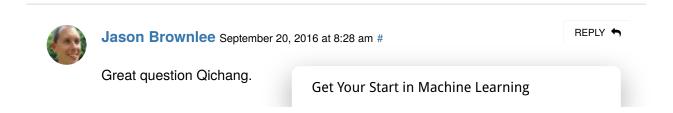
Hi Jason,

Thanks for the tutorial with such useful information! I have one question regarding the label encoding and the one hot encoding you applied on the breast cancer dataset.

You perform label encoding and one hot encoding for the whole dataset and then split into train and test set. This way it can be ensured that all the data are transformed with the same encoding configuration.

However, if we have new unseen data with the raw dataset type, how can we ensure that label encoding and one hot encoding is still transforming the unseen data in the same way? Do we need to save the encoders for the sake of processing unseen data?

Thanks in advance!



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I would prepare the encodings on the training data, store the mappings (or pickle the objects), then reuse the encodings on the test data.

It means we must be confident that the training data is representative of the data we may need to predict in the future.



Qichang September 20, 2016 at 7:47 pm #



Thanks Jason for the prompt reply.

Besides this kinds of data transformation, do we need to consider scaling or normalisation of the input variables before passing to XGBoost? We know that it generally yields better result for SVM especially with kernel function.



Jason Brownlee September 21, 2016 at 8:28 am #



Generally no scaling. You may see some benefit by spreading out a univariate distribution to highlight specific features (e.g. with a square, log, square root, etc.)



JChen October 12, 2016 at 5:26 am #



Hi Jason, your post is very helpful. Thanks a lot!

I had a question around how to treat "default" values of continuous predictors for XGBoost. For example, let's say attributer X may take continuous values as such (say in range of 1 -100). But certain records may have some default value (say 9999) which denotes certain segment of customers for whom that predictor X cannot be calculated or is unavailable. Can we directly use predictor X as input variable for an XGBoost model? Or, should we do some data treatment for X? If so, what would that be?

TIA



Jason Brownlee October 12, 2016 at 9:14 am #

REPLY 🖴

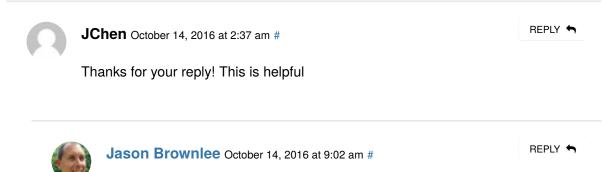
Great question JChen.

I would try modeling the data as-is to sta

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You could then try some feature engineering (maybe add a new flag variable for such cases) and see if you can further list performance.



I'm glad to hear it JChen.



For your missing data part you replaced '?' with 0. But you have not mentioned while defining XGBClassifier model that in your dataset treat 0 as missing value. And by default 'missing' parameter value is none which is equivalent to treating NaN as missing value. So i don't think your model is handling missing values.



Thanks for the note Sargam.

Tursun April 19, 2017 at 11:54 pm #

REPLY 5

REPLY 5

REPLY 🦴

Hi Jason,

I came to this page from another: IRIS,

source: http://machinelearningmastery.com/multi-class-classification-tutorial-keras-deep-learninglibrary/#comment-396630

In that webPage, your code classifies IRIS with 96.6% accuracy, which is very good. In comments section, you told this guy (Abhilash Menon) to use gradient boosting with XGBoost. Which is this tutorial. Here is also IRIS classification.

Well I do not understand:

Here, the IRIS classification with gradient bo

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Why should we use gradient boosting with XGBoost then? Accuracy is too low.



Jason Brownlee April 20, 2017 at 9:28 am #



The tutorial is a demonstration of how to use the algorithm.

Your job as the machine learning practitioner is to try many algorithms on your problem and see what works best.

See this post:

http://machinelearningmastery.com/a-data-driven-approach-to-machine-learning/



daniel May 10, 2017 at 1:43 am #



Hi, Jason. First of all i would like to thank you for the wonderful material. I have been trying the XGBoost algorithm, it seems its acting weird on my pc. The first iris species dataset i got score of 21.2 %. Now this one of the breast cancer my accuracy is 2.1%. I really dont know wats wrong, can you please, please help me



Jason Brownlee May 10, 2017 at 8:51 am #



I'm sorry to hear that. Perhaps the library is not installed correctly?

I would recommend posting a question to stackoverflow or even the xgboost users group.

Let me know how you go.



Giulio June 15, 2017 at 3:50 am #



Hello Jason, I know that for regression models we should drop the first dummy variable to avoid the "dummy trap". I don't see you doing that in this case. Is is because the dummy variable trap only applies to linear regression models and not gradient boosting algorithms?

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Jas	on Brownlee June 15, 2017 at 8:50 am # I believe the dummy variable trap applies with linear models and when the are multicollinear.	REPLY 5 ne variables
	Eran August 25, 2017 at 12:22 am # Hello Jason. Appens if I have a string column with NA's? first put 0's instead of the NA's and then use the OneHotEncounter?	REPLY 🦘
=	Jason Brownlee August 25, 2017 at 6:44 am # XGBoost can handle the NAs. No need to do anything. u do want to impute or similar, see this post: s://machinelearningmastery.com/handle-missing-data-python/	REPLY
eave a	Reply	
	Name (required) Email (will not be published) (required)	

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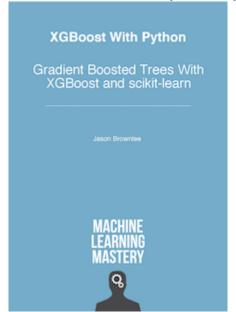
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