```
This notebook is an exercise in the <u>Intermediate Machine Learning</u> course. You can
         reference the tutorial at this link.
         By encoding categorical variables, you'll obtain your best results thus far!
         Setup
         The questions below will give you feedback on your work. Run the following cell to set up the
         feedback system.
In [ ]: |# Set up code checking
         import os
         if not os.path.exists("../input/train.csv"):
              os.symlink("../input/home-data-for-ml-course/train.csv", "../input/t
              os.symlink("../input/home-data-for-ml-course/test.csv", "../input/te
         st.csv")
         from learntools.core import binder
         binder.bind(globals())
         from learntools.ml_intermediate.ex3 import *
         print("Setup Complete")
         In this exercise, you will work with data from the Housing Prices Competition for Kaggle Learn
         Users.
         Ames Housing dataset image
         Run the next code cell without changes to load the training and validation sets in X_train,
         X_{valid}, y_{train}, and y_{valid}. The test set is loaded in X_{test}.
In [ ]: | import pandas as pd
         from sklearn.model_selection import train_test_split
         # Read the data
         X = pd.read_csv('../input/train.csv', index_col='Id')
         X_test = pd.read_csv('../input/test.csv', index_col='Id')
         # Remove rows with missing target, separate target from predictors
         X.dropna(axis=0, subset=['SalePrice'], inplace=True)
         y = X.SalePrice
         X.drop(['SalePrice'], axis=1, inplace=True)
         # To keep things simple, we'll drop columns with missing values
         cols_with_missing = [col for col in X.columns if X[col].isnull().any()]
         X.drop(cols_with_missing, axis=1, inplace=True)
         X_test.drop(cols_with_missing, axis=1, inplace=True)
         # Break off validation set from training data
         X_train, X_valid, y_train, y_valid = train_test_split(X, y,
                                                                      train_size=0.8, te
         st_size=0.2,
                                                                      random_state=0)
         Use the next code cell to print the first five rows of the data.
In [ ]: | X_train.head()
         Notice that the dataset contains both numerical and categorical variables. You'll need to encode
         the categorical data before training a model.
         To compare different models, you'll use the same <code>score_dataset()</code> function from the tutorial.
         This function reports the mean absolute error (MAE) from a random forest model.
In [ ]: from sklearn.ensemble import RandomForestRegressor
         from sklearn.metrics import mean_absolute_error
         # function for comparing different approaches
         def score_dataset(X_train, X_valid, y_train, y_valid):
             model = RandomForestRegressor(n_estimators=100, random_state=0)
             model.fit(X_train, y_train)
             preds = model.predict(X_valid)
              return mean_absolute_error(y_valid, preds)
         Step 1: Drop columns with categorical data
         You'll get started with the most straightforward approach. Use the code cell below to preprocess
         the data in X_train and X_valid to remove columns with categorical data. Set the
         preprocessed DataFrames to drop_X_train and drop_X_valid, respectively.
In [ ]: | # Fill in the lines below: drop columns in training and validation data
         drop_X_train = X_train.select_dtypes(exclude=['object'])
         drop_X_valid = X_valid.select_dtypes(exclude=['object'])
         # Check your answers
         step_1.check()
In [ ]: | # Lines below will give you a hint or solution code
         #step_1.hint()
         #step_1.solution()
         Run the next code cell to get the MAE for this approach.
In [ ]: print("MAE from Approach 1 (Drop categorical variables):")
         print(score_dataset(drop_X_train, drop_X_valid, y_train, y_valid))
         Before jumping into label encoding, we'll investigate the dataset. Specifically, we'll look at the
         'Condition2' column. The code cell below prints the unique entries in both the training and
         validation sets.
In [ ]: print("Unique values in 'Condition2' column in training data:", X_train[
         'Condition2'].unique())
         print("\nUnique values in 'Condition2' column in validation data:", X_va
         lid['Condition2'].unique())
         Step 2: Label encoding
         Part A
         If you now write code to:

    fit a label encoder to the training data, and then

    use it to transform both the training and validation data,

         you'll get an error. Can you see why this is the case? (You'll need to use the above output to
         answer this question.)
In [ ]: # Check your answer (Run this code cell to receive credit!)
         step_2.a.check()
In [ ]: |#step_2.a.hint()
         This is a common problem that you'll encounter with real-world data, and there are many
         approaches to fixing this issue. For instance, you can write a custom label encoder to deal with
         new categories. The simplest approach, however, is to drop the problematic categorical columns.
         Run the code cell below to save the problematic columns to a Python list bad label cols.
         Likewise, columns that can be safely label encoded are stored in good_label_cols.
In [ ]: # All categorical columns
         object_cols = [col for col in X_train.columns if X_train[col].dtype ==
         "object"]
         # Columns that can be safely label encoded
         good_label_cols = [col for col in object_cols if
                               set(X_train[col]) == set(X_valid[col])]
         # Problematic columns that will be dropped from the dataset
         bad_label_cols = list(set(object_cols)-set(good_label_cols))
         print('Categorical columns that will be label encoded:', good_label_cols
         print('\nCategorical columns that will be dropped from the dataset:', ba
         d_label_cols)
         Part B
         Use the next code cell to label encode the data in X_train and X_valid . Set the
         preprocessed DataFrames to label_X_train and label_X_valid, respectively.

    We have provided code below to drop the categorical columns in bad_label_cols from

             the dataset.

    You should label encode the categorical columns in good_label_cols.

In [ ]: from sklearn.preprocessing import LabelEncoder
         # Drop categorical columns that will not be encoded
         label_X_train = X_train.drop(bad_label_cols, axis=1)
         label_X_valid = X_valid.drop(bad_label_cols, axis=1)
         # Apply label encoder
         label_encoder = LabelEncoder()
         for col in good_label_cols:
             label_X_train[col] = label_encoder.fit_transform(label_X_train[col])
             label_X_valid[col] = label_encoder.transform(label_X_valid[col])
         # Check your answer
         step_2.b.check()
In [ ]: # Lines below will give you a hint or solution code
         #step_2.b.hint()
         #step_2.b.solution()
         Run the next code cell to get the MAE for this approach.
In [ ]: print("MAE from Approach 2 (Label Encoding):")
         print(score_dataset(label_X_train, label_X_valid, y_train, y_valid))
         So far, you've tried two different approaches to dealing with categorical variables. And, you've
         seen that encoding categorical data yields better results than removing columns from the dataset.
         Soon, you'll try one-hot encoding. Before then, there's one additional topic we need to cover.
         Begin by running the next code cell without changes
In [ ]: # Get number of unique entries in each column with categorical data
         object_nunique = list(map(lambda col: X_train[col].nunique(), object_col
         d = dict(zip(object_cols, object_nunique))
         # Print number of unique entries by column, in ascending order
         sorted(d.items(), key=lambda x: x[1])
         Step 3: Investigating cardinality
         Part A
         The output above shows, for each column with categorical data, the number of unique values in
         the column. For instance, the 'Street' column in the training data has two unique values:
         'Grv1' and 'Pave', corresponding to a gravel road and a paved road, respectively.
         We refer to the number of unique entries of a categorical variable as the cardinality of that
         categorical variable. For instance, the 'Street' variable has cardinality 2.
         Use the output above to answer the questions below.
In [ ]: | # Fill in the line below: How many categorical variables in the training
         # have cardinality greater than 10?
         high_cardinality_numcols = 3
         # Fill in the line below: How many columns are needed to one-hot encode
         # 'Neighborhood' variable in the training data?
         num_cols_neighborhood = 25
         # Check your answers
         step_3.a.check()
In [ ]: # Lines below will give you a hint or solution code
         #step_3.a.hint()
         #step_3.a.solution()
         Part B
         For large datasets with many rows, one-hot encoding can greatly expand the size of the dataset.
         For this reason, we typically will only one-hot encode columns with relatively low cardinality. Then,
         high cardinality columns can either be dropped from the dataset, or we can use label encoding.
         As an example, consider a dataset with 10,000 rows, and containing one categorical column with
         100 unique entries.

    If this column is replaced with the corresponding one-hot encoding, how many entries are

             added to the dataset?
           If we instead replace the column with the label encoding, how many entries are added?
         Use your answers to fill in the lines below.
In [ ]: # Fill in the line below: How many entries are added to the dataset by
         # replacing the column with a one-hot encoding?
         OH_entries_added = 990000
         # Fill in the line below: How many entries are added to the dataset by
         # replacing the column with a label encoding?
         label_entries_added = 0
         # Check your answers
         step_3.b.check()
In [ ]: # Lines below will give you a hint or solution code
         #step_3.b.hint()
         #step_3.b.solution()
         Next, you'll experiment with one-hot encoding. But, instead of encoding all of the categorical
         variables in the dataset, you'll only create a one-hot encoding for columns with cardinality less
         than 10.
         Run the code cell below without changes to set low_cardinality_cols to a Python list
         containing the columns that will be one-hot encoded. Likewise, high_cardinality_cols
         contains a list of categorical columns that will be dropped from the dataset.
In [ ]: # Columns that will be one-hot encoded
         low_cardinality_cols = [col for col in object_cols if X_train[col].nuniq
         ue() < 10
         # Columns that will be dropped from the dataset
         high_cardinality_cols = list(set(object_cols)-set(low_cardinality_cols))
         print('Categorical columns that will be one-hot encoded:', low_cardinali
         ty_cols)
         print('\nCategorical columns that will be dropped from the dataset:', hi
         gh_cardinality_cols)
         Step 4: One-hot encoding
         Use the next code cell to one-hot encode the data in X_train and X_valid . Set the
         preprocessed DataFrames to OH_X_train and OH_X_valid, respectively.

    The full list of categorical columns in the dataset can be found in the Python list

             object_cols.

    You should only one-hot encode the categorical columns in low_cardinality_cols. All

             other categorical columns should be dropped from the dataset.
In [ ]: from sklearn.preprocessing import OneHotEncoder
         # Use as many lines of code as you need!
         one_hot_encoder = OneHotEncoder(handle_unknown = 'ignore', sparse = Fals
         OH_X_train = pd.DataFrame(one_hot_encoder.fit_transform(X_train[low_card
         inality_cols]))
         OH_X_valid = pd.DataFrame(one_hot_encoder.transform(X_valid[low_cardinal
         ity_cols]))
         OH_X_train.index = X_train.index
         OH_X_valid.index = X_valid.index
         X_train_num_cols = X_train.drop(object_cols, axis = 1, inplace = False)
         X_valid_num_cols = X_valid.drop(object_cols, axis = 1, inplace = False)
         OH_X_train = pd.concat([X_train_num_cols, OH_X_train], axis = 1)
         OH_X_valid = pd.concat([X_valid_num_cols, OH_X_valid], axis = 1)
         # Check your answer
         step_4.check()
In [ ]: # Lines below will give you a hint or solution code
         #step_4.hint()
         #step_4.solution()
```

print(score_dataset(OH_X_train, OH_X_valid, y_train, y_valid)) Generate test predictions and submit your

results

Run the next code cell to get the MAE for this approach.

In []: print("MAE from Approach 3 (One-Hot Encoding):")

After you complete Step 4, if you'd like to use what you've learned to submit your results to the leaderboard, you'll need to preprocess the test data before generating predictions. This step is completely optional, and you do not need to submit results to the leaderboard to successfully complete the exercise.

Check out the previous exercise if you need help with remembering how to join the competition or save your results to CSV. Once you have generated a file with your results, follow the instructions below: 1. Begin by clicking on the blue **Save Version** button in the top right corner of the window. This

will generate a pop-up window. 2. Ensure that the **Save and Run All** option is selected, and then click on the blue **Save** button. 3. This generates a window in the bottom left corner of the notebook. After it has finished

running, click on the number to the right of the Save Version button. This pulls up a list of versions on the right of the screen. Click on the ellipsis (...) to the right of the most recent version, and select **Open in Viewer**. This brings you into view mode of the same page. You

will need to scroll down to get back to these instructions. 4. Click on the **Output** tab on the right of the screen. Then, click on the file you would like to

submit, and click on the blue Submit button to submit your results to the leaderboard.

You have now successfully submitted to the competition!

If you want to keep working to improve your performance, select the blue **Edit** button in the top right of the screen. Then you can change your code and repeat the process. There's a lot of room to improve, and you will climb up the leaderboard as you work.

In []: # (Optional) Your code here

Keep going

With missing value handling and categorical encoding, your modeling process is getting complex. This complexity gets worse when you want to save your model to use in the future. The key to managing this complexity is something called **pipelines**.

<u>Learn to use pipelines</u> to preprocess datasets with categorical variables, missing values and any other messiness your data throws at you.

Have questions or comments? Visit the <u>Learn Discussion forum</u> to chat with other Learners.