Introduction

One very hot area in the current field of data science is forecasting housing and real-estate prices. The real-estate company Zillow recently made headlines when they announced that they were discontinuing their "iBuying" feature as they were rapidly losing money due to a failure to accurately estimate house prices using algorithms. In general, financial data modeling and financial forecasting is extremely difficult (bordering on "impossible"), and real-estate forecasting has many similar problems.

In light of this, it seemed obvious that a housing dataset would make a good choice for the assignment as most people interact with the real estate market over their lifetimes and the usual features of housing datasets (geographic features, physical features, etc.) are quite fun to play around with. Consequently, a suitable housing dataset was found and used as the basis for the assignment.

Methodology

Data and Data Cleaning

The dataset of interest came from kaggle.com and concerned housing data from the city of Melbourne, Australia in the years 2017 and 2018 (https://www.kaggle.com/anthonypino/melbourne-housing-market? select=Melbourne_housing_FULL.csv). The Kaggle dataset did not originally have a data dictionary and documentation was very sparse in general, but a data dictionary was found on a third-party GitHub repository (https://github.com/njtierney/melb-housing-data), which helped shed light on some of the data attributes. The base dataset had 34.857 rows and 21 columns.

Cleaning the data was a multi-step process. To start with, all rows that did not have pricing information were dropped. A number of columns were also dropped. This was because these were irrelevant or hard to work with (such as the name of the seller or the date sold) or because the number of unique values was prohibitively large (such as the list of unique addresses). In total, 9 features were dropped in this manner.

One tricky thing to consider was the question of area (how large the property is), which was expected to correlate highly with price. In the base dataset there were two features meant to estimate this ("Landsize" and "BuildingArea"), but neither feature appeared in the data dictionary, both had very large numbers of missing values (about 9,200 for Landsize and about 16,600 for BuildingArea), and for cases where neither value was missing one value was not consistently larger than the other. While definitely dirty and lacking in documentation, early exploration showed that both variables had a Pearson's correlation of more than 0.4 with price, meaning that they were too valuable to simply discard.

To reconcile this, a new feature, "Area", was engineered which was equal to the average of Landsize and BuildingArea (for cases where both values existed) or just whichever value was present (if only one value was present for a given datapoint). "Area" was then taken to the 4th root to ensure a sufficiently smooth distribution between 0 and 1 during the normalization phase of the data cleaning process.

To finish cleaning the dataset, two outliers for "Price" and "YearBuilt" were excluded and almost all features were scaled to be in the range of 0 to 1 with missing values indicated as -1. The two remaining nominal variables ("Type" and "RegionName") were converted to a one hot encoding scheme. The final, cleaned dataset consisted of 27,245 rows and 11 features (21 features after one hot encoding). The final list of features in the cleaned dataset and how they were processed can be seen in Table 1.

Table of Features in Cleaned Dataset

Column	Definition	Handling
Rooms	The number of bedrooms in the property	Normalized
Туре	The property type (house, townhouse, or unit)	One Hot Encoded
Price	The sale price in Australian Dollars. Also the label variable to be predicted	Unaltered, rows with missing values dropped
Distance	The property's distance from the Melbourne Central Business District (in km)	Normalized
Bathroom	The number of bathrooms	Normalized
Car	The number of carspots for the property	Normalized
Area	An engineered feature representing the "size" of the property	Engineered, taken to the 4th root
YearBuilt	The year the property was built	Normalized
Lattitude	The latititude of the property	Normalized
Longtitude	The longitude of the property	Normalized
Regionname	The name of the city region the property exists in	One Hot Encoded

Table 1: List of features in the cleaned dataset

```
In [1]:
        import numpy as np
        import pandas as pd
        import matplotlib.pyplot as plt
        import qc
        from sklearn.metrics import mean absolute error
        from tensorflow.keras import models, layers
        from tensorflow.keras.optimizers import Adam, RMSprop
        from tensorflow.random import set seed
In [2]:
        # helper functions.
        def normalize(df, list of cols):
            """Helper function that normalizes the given columns for a df. Returns the new df."""
            for col in list of cols:
                df[col] = (df[col] - np.nanmin(df[col]))/(np.nanmax(df[col]) - np.nanmin(df[col]))
            return (df)
        def split into validate(data array, labels array, validation proportion):
            """Helper function that split train x and train y into train and validate x and y"""
             # Getting train and validation indices. Validation indices are randomly selected from
             # just whichever indices where not chosen as validation indices
            overall indices = range(len(data array))
            val indices = np.random.choice(a = overall indices, size = int(len(data array) * valid
                                             replace = False)
```

```
# Training inidices are the set of overall indices after we take out the validation i
            train indices = list(set(overall indices).difference(val indices))
            # train data partition
            train X = data array[train indices]
            train Y = labels array[train indices]
            # validation data partition
            val X = data array[val indices]
            val Y = labels array[val indices]
            return(train X, train Y, val X, val Y)
In [3]:
        # Setting seeds for replication.
        seed val = 19
        np.random.seed(seed val)
        set seed(seed val)
In [4]:
        # Reading in the base df
        base df = pd.read csv("Melbourne housing FULL.csv")
         # Drop all rows where we don't have price data.
        cleaned df = base df[base df["Price"].notna()]
         # Get rid of the annyoing warning messages for slice assignment.
        pd.options.mode.chained assignment = None # default='warn'
        # Drop some columns. Either too many unique values or just generally irrelvent.
        drop cols = ["Suburb", "Address", "Method", "SellerG", "Date", "Postcode", "Bedroom2",
                     "Propertycount", "CouncilArea"]
        cleaned df.drop(axis = 1, columns=drop cols, inplace=True)
         # Build a new measure, "Area", from the two really dirty columns.
        temp matrix = cleaned df[["Landsize", "BuildingArea"]].values
        cleaned df["Area"] = np.nanmean(temp matrix, axis = 1)
        # Area has an extremely long tail. Take the 4th root to compress the range of value and
         # prevent bunching.
        cleaned df["Area"] = np.sqrt(np.sqrt(cleaned df["Area"]))
         # No longer needed.
        del temp matrix
        # Drop the no longer needed columns relating to area.
        drop cols = ["Landsize", "BuildingArea"]
        cleaned df.drop(axis = 1, columns=drop cols, inplace=True)
         # rename one of the columns.
        cleaned df.rename(columns ={"Regionname": "RegionName"}, inplace =True )
         # Excluding a few outliers we noticed during exploration. We drop them from the dataset.
        cleaned df = cleaned df[cleaned df["Price"] != 11200000.0]
        cleaned df = cleaned df[cleaned df["YearBuilt"]!= 1196]
         # For all these columns, range to [0 - 1]. Missing values will be indicated with -1.
        normalized columns = ['Rooms', 'Distance', 'Bathroom', 'Car', 'YearBuilt', 'Lattitude', 'I
        cleaned df = normalize(cleaned df, normalized columns)
         # For the entire dataframe, fill in NA values with -1 after normalization.
```

```
cleaned df.fillna(value = -1, inplace = True)
         # For all these, convert labels to OHE.
        cleaned df = pd.get dummies(data = cleaned df, columns= ["Type", "RegionName"])
         # Save cleaned dataframe to disk, just to be on the safe side.
        cleaned df.to csv("cleaned melbourne.csv", header=True, index=False)
       C:\Users\Jack\AppData\Local\Temp/ipykernel 35880/3317385550.py:17: RuntimeWarning: Mean of
       empty slice
         cleaned df["Area"] = np.nanmean(temp matrix, axis = 1)
In [5]:
        # Going over everything to make sure our distributions are sensible after normalization.
         # This has been commented out to avoid cluttering things,
         # for key in cleaned df.keys():
             arr = cleaned df[key].unique()
             print(key)
             print("----")
             try:
                 print(np.sort(arr))
            except:
        #
         #
               print(arr)
         # plt.plot(np.sort(cleaned_df[key]))
              plt.show()
In [6]:
        # Shuffle the dataframe to prevent ordering bias.
        cleaned df = cleaned df.sample(frac=1).reset index(drop=True)
         # Taking our dataframe and extracting the relevant data as matrices.
        Y dataset = cleaned df["Price"].values
        cleaned df.drop(axis = 1, columns=["Price"], inplace=True)
        X dataset = cleaned df.values
        # 80/20 train test split
        cutoff value = int(0.8 * len(Y dataset))
        X train = X dataset[:cutoff value]
        X test = X dataset[cutoff value:]
        Y train = Y dataset[:cutoff value]
        Y test = Y dataset[cutoff value:]
        # Convert to numpy arrays.
        train indices = np.array(range(len(X train)))
        test indices = np.array(range(len(X test)))
        X train = np.array(X train)[train indices.astype(int)]
        X test = np.array(X test)[test indices.astype(int)]
        Y train = np.array(Y train)[train indices.astype(int)]
        Y test = np.array(Y test)[test indices.astype(int)]
         # No longer needed.
        del cutoff value, X dataset, Y dataset
```

Network Topologies and Training

To estimate housing prices, three different networks were constructed. The first was simple network that could outperform a pre-selected baseline, the second was a high-capacity network that was deliberately meant to overfit, and the third was a final network that was an optimized and regularized version of the high-capacity network For the baseline measure of success, the simple average of the housing prices in the training dataset was selected.

For all of the layers in every network the only activations used were ReLU. The output in every network was a single neuron with relu activation, as the variable of interest for this assignment was house prices which are positive real numbers that range from 0 to several million, making ReLU a natural choice.

For the loss metric, mean absolute error (MAE) was chosen. This was because the task was estimating asset prices, and in such a task financial loss and gain is completely fungible (i.e. losing 5,000 dollars twice is exactly as bad as losing 10,000 dollars once). Early experimentation also revealed that fitting was very bad with mean squared error, as the network became far too sensitive to outliers (i.e. the smaller number of extremely expensive properties).

For the optimizer, the ADAM (adaptive moment estimation) optimizer was chosen, as it is a more advanced optimizer that builds on the work of previous optimizers such as AdaGrad and RMSProp by making use of both the first and second statistical moments for gradient updates. All parameters for the ADAM optimizer were kept constant at the default values with the exception of the learning rate, which was part of hyperparameter optimization for finding the final model

For hyperparameter optimization, the strategy employed was a simple grid search. The two hyperparameters considered were the batch size (8, 16, 32, or 64) and the learning rate (0.005, 0.001, 0.0005, or 0.0001). The number of epochs was kept constant at 200. The training and testing sets were a simple 80/20 split, meaning that each training set was 64% of the data, each validation set was 16% of the data, and the testing set was 20% of the data.

For validation, simple hold-out validation was used where the the validation set was always 20% of the data in the training set, randomly chosen each time before training. The model's performance on the validation set was used to select the best hyperparameters for training the final mode, and performance on the validation and training sets was also used to identify the ideal number of training

After hyperparameter optimization, the best set of hyperparameters was fed into a final model consisting of two hidden layers of 32 neurons, dropout of 0.2, and relu activation. The final model was trained for 2,400 epochs to try and get the most ideal fit on the training data.

Baseline.

The "baseline" model is simply the average of the house prices on the training dataset.

MAE on testing set if just taking the average: 456321.5907042787

Simple Model.

A simple model that performs better than the baseline. The model consists of two hidden layers, each with 8 neurons. All activations are Relu. The model has a batch size of 32, a learning rate of 0.001, and is trained for 200 epochs. The model is able to beat the baseline score, as demonstrated below.

```
In [ ]:
         simp network = models.Sequential()
         simp network.add(layers.Dense(8, activation='relu', input shape=(20, )))
         simp network.add(layers.Dense(8, activation='relu', input shape=(8, )))
         simp network.add(layers.Dense(1, activation='relu', input shape= (8, )))
         # optimiser, loss and metrics are chosen at compilation
         simp network.compile(optimizer=Adam(lr=0.001),
                         loss='mean absolute error',
                         metrics=["mean absolute error"])
         simp network.fit(X train,
                         Y train,
                         epochs=200,
                                                     # Run for 200 epochs.
                         batch size = 32,
In [10]:
         print("Train MAE: {}".format(mean absolute error(simp network.predict(X train), Y train)))
         print("Test MAE: {}".format(mean absolute error(simp network.predict(X test), Y test)))
```

Train MAE: 280723.8074483635 Test MAE: 288779.00253262854

Model that overfits.

A deliberately over-capacity model that performs better than the baseline. The model consists of five hidden layers, each with 32 neurons. All activations are Relu. The model has a batch size of 32, a learning rate of 0.001, and is trained for 3000 epochs. After extension experimentation, the point where overfitting happens is around epoch 2400.

```
In [ ]:
        overfit network = models.Sequential()
        overfit network.add(layers.Dense(32, activation='relu', input shape=(20, )))
        overfit network.add(layers.Dense(32, activation='relu', input shape=(32, )))
        overfit network.add(layers.Dense(1, activation='relu', input shape= (32, )))
        # optimiser
        overfit network.compile(optimizer=Adam(lr=0.001),
                        loss='mean absolute error',
                        metrics=["mean absolute error"])
        train data, train labels, val data, val labels = split into validate(X train, Y train, 0.2
        learning history = overfit network.fit(
                        x = train data,
                         y = train labels,
                        epochs=3000,
                                                      # Run for 3000 epochs.
                        batch size = 32,
                         validation data = (val data, val labels)
```

```
# No longer needed past this point.
         del train data, train labels, val data, val labels
In [12]:
         print("Train MAE: {}".format(mean absolute error(overfit network.predict(X train), Y train
         print("Test MAE: {}".format(mean absolute error(overfit network.predict(X test), Y test)))
         best epoch = np.argmin(learning history.history['val loss'])
         print("Best Epoch Number: {}".format(best epoch))
        Train MAE: 170737.4840122614
        Test MAE: 193264.34697020095
        Best Epoch Number: 2403
In [13]:
         plt.plot(range(3000), learning history.history['loss'], label = "Train")
         plt.plot(range(3000), learning history.history['val loss'], label = "Validation")
         plt.title('Performance of the over-capacity model.')
         plt.ylabel('MAE')
         plt.xlabel('epoch')
         plt.legend(loc='upper right')
         plt.show()
         # plt.savefig("bias variance.png")
```

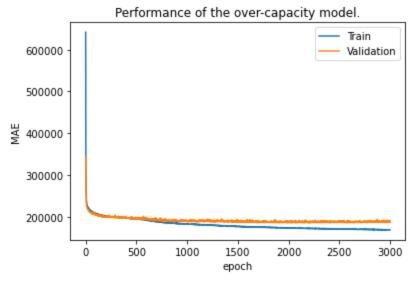


Figure 1: Bias/Variance decomposition for the overfit network.

Final model

The final model that makes use of hyperparameter optimization and regularization. Based on the results from training the overfitting network, the model was trained for 2,400 epochs. The model consisted of two layers of 32 neurons with dropout of 0.2 on each layer.

For hyperparameter optimization, A unique model was trained for each hyperparameter combination (16 overall) for 200 epochs each. The validation and training loss for each of those individual networks was then recorded and used to determine the best hyperparameters for training the final.

Hyperparameter Optimization

```
In [ ]: # Array to hold the train and validation scores for each combination.
    results_array = np.zeros(shape = (4, 4, 2))
```

```
for j, learning rate in enumerate([0.005, 0.001, 0.0005, 0.0001]):
                 # Garbage collection for optimization reasons.
                 gc.collect()
                 # Deliberately over-capacity fully connected network. For the final network, these
                 # with 32 neurons.
                 network = models.Sequential()
                 network.add(layers.Dense(32, activation='relu', input shape=(20, )))
                 network.add(layers.Dropout(.2, input shape=(32,)))
                 network.add(layers.Dense(32, activation='relu', input shape=(32, )))
                 network.add(layers.Dropout(.2, input shape=(32,)))
                 network.add(layers.Dense(1, activation='relu', input shape= (32, )))
                 # optimiser, loss and metrics are chosen at compilation
                 network.compile(optimizer=Adam(lr=learning rate),
                                 loss='mean absolute error',
                                 metrics=["mean absolute error"])
                 train data, train labels, val data, val labels = split into validate(X train, Y ti
                 learning history = network.fit(
                         x = train data,
                         y = train labels,
                         epochs=200,
                                                     # Run for 200 epochs.
                         batch size = 32,
                         validation data = (val data, val labels)
                 results array[i, j, 0] = np.min(learning history.history["loss"])
                 results array[i, j, 1] = np.min(learning history.history["val loss"])
         del train data, train labels, val data, val labels
In [15]:
         # Finding the best hyperparameters based on the validation performance.
         best = np.min(results array[:, :, 1])
         indices of best validation = np.where(results array[:, :, 1] == best)
         print("best batch size indice: {}".format(indices of best validation[0][0]))
         print("best learning rate indice: {}".format(indices of best validation[1][0]))
         # Plotting df to hold our values
         plotting df = pd.DataFrame({"validation": results array[:, :, 1].flatten(),
                                     "training": results array[:, :, 0].flatten()})
         plotting df.sort values(by = "training", ascending=False, inplace=True)
         plotting df.reset index(drop=True, inplace=True)
         # Plot train and validation, by training descending
         plt.plot(range(1, 17), plotting df["training"], '-o', c = "blue", label = "training loss")
         plt.plot(range(1, 17), plotting df["validation"], '-o', c = "red", label = "validation los
         plt.xlabel("")
         plt.ylabel("MAE")
         plt.title("Loss")
         plt.legend()
```

for i, batch size in enumerate([8, 16, 32, 64]):

plt.show()

```
# plt.savefig("train_and_val_plot.png")

# # no longer_needed
del plotting_df, best, indices_of_best_validation
```

```
best batch size indice: 1
best learning rate indice: 0
```

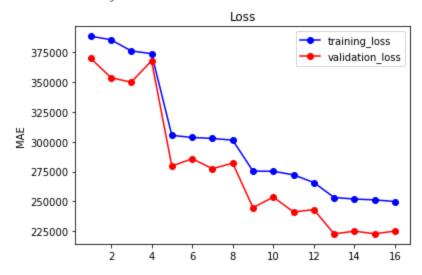


Figure 2: Training and Validation loss for every hyperparameter combination. Notice the serated pattern, implying that one of the hyperparameters is more important than the other.

Final Model training.

```
In [ ]:
         # Real final model here. Learning rate and batch size selected based off of previous hyper
        final network = models.Sequential()
        final network.add(layers.Dense(32, activation='relu', input shape=(20, )))
        final network.add(layers.Dropout(.2, input shape=(32,)))
        final network.add(layers.Dense(32, activation='relu', input shape=(32, )))
        final network.add(layers.Dropout(.2, input shape=(32,)))
        final network.add(layers.Dense(1, activation='relu', input shape= (32, )))
         # optimiser, loss and metrics are chosen at compilation
        final network.compile(optimizer=Adam(1r=0.005),
                         loss='mean absolute error',
                         metrics=["mean absolute error"])
        learning history = final network.fit(
                         x = X train,
                         y = Y train,
                                                      # Run for 2400 epochs.
                         epochs=2400,
                         batch size = 16
```

```
In [17]: print("Final Model Train MAE: {}".format(mean_absolute_error(final_network.predict(X_train print("Final Model Test MAE: {}".format(mean_absolute_error(final_network.predict(X_test),
```

Final Model Train MAE: 200119.21404586852 Final Model Test MAE: 208717.87287231602

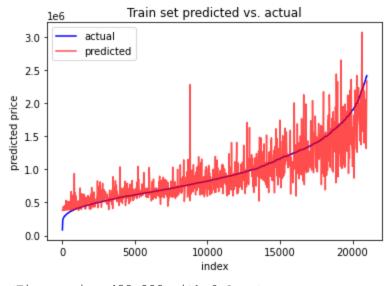
Results

The results of the hyperparameter selection are displayed in Figure 2. The best hyperparameter combination was a learning rate of 0.005 and a batch size of 16, which were forwarded and used to train the final model.

The final network did perform fairly well, as can be seen in Figures 3 and 4, though not quite as well as the overfit model. The final network demonstrated substantial improvement over average imputation, though there was still quite some room for improvement, particulary given the network's noticable inabillity to accurately forecast the lowest value homes in both the training set and the testing set. Finding the "sweet" spot in terms of fitting was also quite tricky, with both the overfit model and the final model scoring better on the training set than the testing set.

Overall, despite a few hiccups the universal workflow of machine learning was followed closely, with substantial improvement seen across multiple networks, though never quite to the desired level of performance. Table 2 shows the performance for every network.

```
In [18]:
         # Make two different dfs: one for the training set and one for the testing set.
         train pred = final network.predict(X train)
         train plot df = pd.DataFrame({"Actual": np.squeeze(Y train), "Predicted": np.squeeze(train
         train plot df.sort values(by = "Actual", ascending=True, inplace=True)
         train plot df.reset index(drop=True, inplace=True)
         test pred = final network.predict(X test)
         test plot df = pd.DataFrame({"Actual": np.squeeze(Y test), "Predicted": np.squeeze(test pr
         test plot df.sort values(by = "Actual", ascending=True, inplace=True)
         test plot df.reset index(drop=True, inplace=True)
         # Plot training
         plt.plot(range(0, 21000, 21), train plot df["Actual"][range(0, 21000, 21)], c = "blue", la
         plt.plot(range(0, 21000, 21), train plot df["Predicted"][range(0, 21000, 21)], c = "red",
                  label = "predicted", alpha = 0.7)
         plt.xlabel("")
         plt.ylabel("predicted price")
         plt.xlabel("index")
         plt.title("Train set predicted vs. actual")
         plt.legend(loc="upper left")
         plt.show()
         # plt.savefig("train performance.png")
```



<Figure size 432x288 with 0 Axes>

Figure 3: Final model performance on the training set

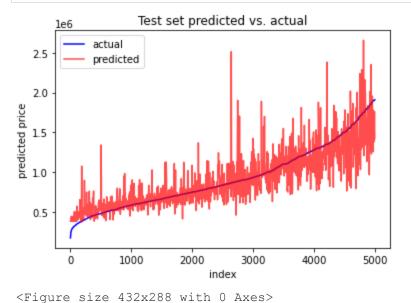


Figure 4: Final model performance on the testing set

Table of Final Results

Model	MAE Train Loss	MAE Test Loss
Average Price	449500	456321
Simple Model	280723	288779
Overfit Model	170737	193264
Final Model	200119	208717

Table 2: Table of all train/test model performance

Conclusions

The assignment proved to be more difficult than expected, as a truly archetypal example of overfitting was never achieved and the process of making sure all details were right for for all of the different training runs (and avoiding information leaks) was quite difficult. As the old saying goes, "programming is the art of a thousand details."

One lesson to take away from this assignment is the importance of feature engineering. Part of the final network's underperformance can probably be attributed to the fact that it was very hard to get useful

geographic data from the dataset (location, location, location...). There were also very important questions of feature engineering and feature inclusion that did not have any obvious answers, and there is a lot of potential for ruining a project before any modeling actually happens just by choosing features badly.

Another important lesson I was reminded of was the sheer complexity of hyperparameter optimization. Selecting from among just 16 hyperparameters and doing validation took many minutes, even with a top-of-the-line graphics card. In real-life, "try all possibilities" (the "responsible" thing to do) is almost never practical, and data scientists have to be judicious with the ways their spend time and decide what areas are worthy of investigation and experimentation. It introduces an import element of intuition into the workflow of machine learning that cannot be easily automated away.

In the future, one better approach than deep learning might be a high-dimensional version of KNN with both temporal and chronological data, assuming the appropriate data can found. Another possible approach be an ensemble model of simple regressors or simpler neural networks, each trained on different feature subsets.

Code Attribution

All code in this notebook was my own. No code was taken from the module notebooks.