```
In [1]: import keras
import numpy as np
from keras.models import Sequential
```

Using TensorFlow backend.

Sentiment Classification

In this problem we will use Kera's imdb sentiment dataset. You will take in sequences of words and use an RNN to try to classify the sequences sentiment. These sentences are movie reviews, so the sentiment reflects whether its a positive review (sentiment of 1) or a negative review (sentiment of 0).

First we have to process the data a little bit, so that we have fixed length sequences.

The data is given to us in integer form, so each integer represents a unique word, 0 represents a PAD character, 1 represents a START character and 2 represents a character that is unknown because it is not in the top num_words. Thus 3 represents the first real word.

Also the words are in decreasing order of commonness, so the word that 3 represents is the most common word in the dataset. (It happens to be the)

We want to process the data into arrays of sequences that are all length 200. If a given sequence is shorter than 200 tokens we want to pad the rest of the sequence out with zeros so that the sequence is 200 long.

```
In [3]: def process_data(data):
    processed = np.zeros(len(data) * 200).reshape((len(data), 200))
    for i, seq in enumerate(data):
        if len(seq) < 200:
            processed[i] = np.hstack((np.array(seq), np.zeros(200 - len(seq)))))
        else:
            processed[i] = np.array(seq)
        return processed</pre>
```

```
In [4]: x_train_proc = process_data(x_train)
x_test_proc = process_data(x_test)
```

```
In [5]: imdb_model = Sequential()
```

Now we want to add an embedding layer. The purpose of an embedding layer is to take a sequence of integers representing words in our case and turn each integer into a dense vector in some embedding space. (This is essentially the idea of Word2Vec). We want to create an embedding layer with vocab size equal to the max num words we allowed when we loaded the data (in this case 1000), and a fixed dense vector of size 32. Then we have to specify the max length of our sequences and we want to mask out zeros in our sequence since we used zero to pad. Use the docs for embedding layer to fill out the missing entries: https://keras.io/layers/embeddings/ (https://keras.io/layers/embeddings/)

```
In [6]: from keras.layers.embeddings import Embedding
imdb_model.add(Embedding(2000, 32, input_length=200, mask_zero=True))
```

WARNING:tensorflow:From C:\Users\anika\Anaconda3\lib\site-packages\tensorflow\python\framework\op_def_library.py:263: colocate_with (from tensorflow.python.framework.ops) is deprecated and will be removed in a future version. Instructions for updating:

Colocations handled automatically by placer.

(a) Add an LSTM layer with 32 outputs, then a Dense layer with 16 neurons, then a relu activation, then a dense layer with 1 neuron, then a sigmoid activation. Then you should print out the model summary. The Keras documentation is here: https://keras.io/ (https://keras.io/)

```
In [7]: from keras.layers.recurrent import LSTM
    from keras.layers import Dense, Activation
    imdb_model.add(LSTM(32))

In [8]: imdb_model.add(Dense(16))
    imdb_model.add(Activation('relu'))

In [9]: imdb_model.add(Dense(1))
    imdb_model.add(Activation('relu'))
```

```
In [10]: imdb_model.summary()
```

| Layer (type) | Output Shape | Param # |
|---------------------------|-----------------|---------|
| embedding_1 (Embedding) | (None, 200, 32) | 64000 |
| lstm_1 (LSTM) | (None, 32) | 8320 |
| dense_1 (Dense) | (None, 16) | 528 |
| activation_1 (Activation) | (None, 16) | 0 |
| dense_2 (Dense) | (None, 1) | 17 |
| activation_2 (Activation) | (None, 1) | 0 |

Total params: 72,865 Trainable params: 72,865 Non-trainable params: 0

If you did the above parts correctly running imdb_model.summary() should give you the following output.

```
In [11]: #from IPython.display import Image
#Image(filename='/home/james/Desktop/Screenshot from 2018-08-20 21-04-15.png')
```

(b) Now compile the model with binary cross entropy, and the adam optimizer. Also include accuracy as a metric in the compile. Then train the model on the processed data (no need to worry about class weights this time)

```
In [12]: imdb_model.compile(loss='binary_crossentropy', optimizer='adam', metrics=['ac c'])
In [13]: imdb_model.fit(x_train_proc, y_train)
WARNING:tensorflow:From C:\Users\anika\Anaconda3\lib\site-packages\tensorflow
```

\warning:tensorflow:from C:\Users\anika\Anaconda3\lib\site-packages\tensorflow
\python\ops\math_ops.py:3066: to_int32 (from tensorflow.python.ops.math_ops)
is deprecated and will be removed in a future version.

Instructions for updating:

Use tf.cast instead.

Epoch 1/1

acc: 0.6848

Out[13]: <keras.callbacks.History at 0x1b25d53b940>

Now we can look at our predictions and the sentences they correspond to.

```
In [15]: y pred = imdb model.predict(x test proc)
In [16]: | word to id = keras.datasets.imdb.get word index()
          word to id = \{k:(v+3) \text{ for } k,v \text{ in word to id.items()}\}
          word to id["\langle PAD \rangle"] = 0
          word_to_id["<START>"] = 1
          word to id["\langle UNK \rangle"] = 2
          id_to_word = {value:key for key,value in word_to_id.items() if value < 2000}</pre>
          def get words(token sequence):
              return ' '.join(id to word[token] for token in token sequence)
          def get sentiment(y pred, index):
              return 'Positive' if y_pred[index] else 'Negative'
In [17]: y pred = np.vectorize(lambda x: int(x >= 0.5))(y pred)
          correct = []
          incorrect = []
          for i, pred in enumerate(v pred):
              if y_test[i] == pred:
                   correct.append(i)
              else:
                   incorrect.append(i)
```

Now we print out one of the sequences we got correct.

```
In [18]: print(get_sentiment(y_pred, correct[10]))
    print(get_words(x_test[correct[10]]))
```

Positive

<START> i went through <UNK> camp at <UNK> island in <UNK> and this film is a bout as accurate a <UNK> of what <UNK> went through in that era even to <UNK> that <UNK> <UNK> many of the actors in the film were <UNK> <UNK> <UNK> this f ilm may be more entertaining to <UNK> than others but i feel the film itself is very well done and jack <UNK> made a good <UNK> fi

And one we got wrong.

```
In [19]: print(get_sentiment(y_pred, incorrect[10]))
    print(get_words(x_test[incorrect[10]]))
```

Positive

<START> this is the worst film i have ever seen bar none from the <UNK> looking poorly <UNK> sets to the laughable acting to the <UNK> plot and <UNK> drawn out action sequences this film is so bad its hilarious for about ten minutes after which you will be <UNK> for the <UNK> or the power <UNK> to end this film non experience although it was obviously made with the entire production and acting <UNK> <UNK> <UNK> in <UNK> please god i found jack <UNK> 2 so <UNK> as to be <UNK> for more than a <UNK> of an hour if you have not had enough of it after this time you must be <UNK> in drug <UNK>

As you can see the amount of UNKNOWN characters in the sequence cause by having only 1000 vocab words is hurting our performance. See if you can go through and increase the number of vocab words to 2000. HINT: you have to change two places in the above code.

```
In [20]: from keras import backend as K
```

Embedding Exploration

Another interesting thing to do is see if our learned embeddings mean anything reasonable.

Since we used an embedding layer with an output size of 32, our embeddings are going to be 32-dimensional vectors. Humans can't effectively visualize beyond 3 (maybe 4) dimensions so we want to use a dimensionality reduction technique to make our embeddings more visualizable. One such technique is Principal Component Analysis or PCA. The library scikit-learn provides an easy to use API for this technique.

```
In [25]: import sklearn
from sklearn import decomposition
```

using the documentation for scikit-learn's PCA <u>here (http://scikit-learn.org/stable/modules/generated/sklearn.decomposition.PCA.html</u>): create a PCA object with n components=2

```
In [26]: pca = sklearn.decomposition.PCA(n_components=2)
```

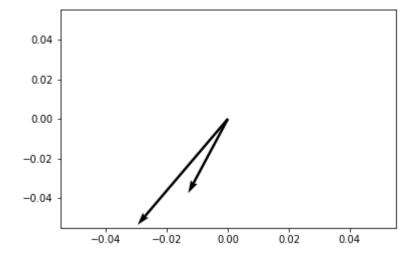
using the same documentation find the function to fit the PCA transform to the provided embedding vectors. This step essentially. For the curious, this step essentially finds the 2 dimensions (since we specified n_components=2 that explain the most variance of the dataset, in other words the two dimensions that are most representative of the deviations of any one sample to another. So these 2 dimensions are the most important and therefore the best to visualize.

Now we want to visualize our embeddings in these new PCA dimensions, so using the same documentation from above fill out the missing spots in the code below to transform the embeddings into the pca dimensions.

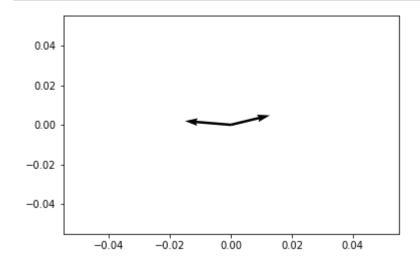
```
In [28]:
          import matplotlib.pyplot as plt
          %matplotlib inline
          def get pca words(words):
              embeddings = [valid word embeddings[word] for word in words]
              pcas = [pca.transform(embedding.reshape(1, -1)) for embedding in embedding
          s]
              return pcas
          def plot pca words(words, scale=1):
              pcas = get pca words(words)
              zeros = [0 for _ in pcas]
              x start = zeros
              y start = zeros
              xs = [p[0, 0] \text{ for } p \text{ in } pcas]
              ys = [p[0, 1]  for p  in pcas
              plt.quiver(x_start, y_start, xs, ys, scale=scale)
              plt.show()
```

Now using the above functions we can plot the corresponding pca vectors of any words we like. Below are some good examples of pairs of words that are similar within the movie review context and their corresponding vectors are also similar. This is a good sign. This means the embedding we have learned is likely doing something somewhat reasonable.

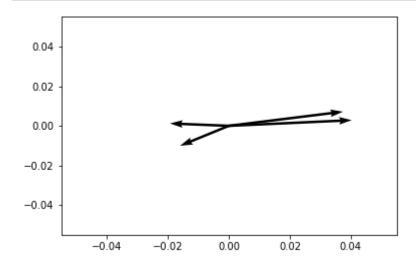
```
In [29]: plot_pca_words(['film', 'entertainment'], scale=0.5)
```



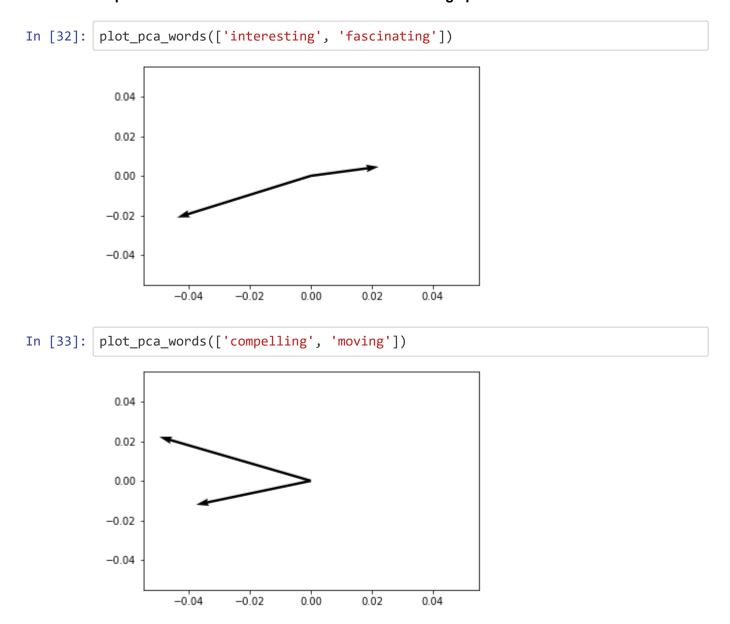
In [30]: plot_pca_words(['man', 'woman'])



In [31]: plot_pca_words(['good', 'bad', 'horrible', 'great'], scale=2)



Now find 2 more pairs of words that are similar in PCA'd embedding space.



Given that the task we learned these embeddings for was sentiment classification, the embeddings are typically more meaningful for adjectives. Write a sentence or two about why you think this last statement makes sense intuitively.

```
In [34]: # Embeddings like adjectives work best to divert results: e.g. good vs. bad, i
    nteresting vs. boring --
# adjectives define the reaction to an event, in this case movies.
```

Now just for fun we can write a function that gives us the 10 closest words to a provided word.

```
In [35]:
         def word to angle(word):
             p = pca.transform(valid word embeddings[word].reshape(-1, 1))
             return np.arctan(p[0, 1] / p[0, 0])
         valid word angles = [word to angle(word) for word in valid words]
In [36]:
         def find_closest_n(value, n):
             indices = np.argsort(np.abs(np.array(valid_word_angles) - value))
             return [(valid words[ind], valid word angles[ind]) for ind in indices[:n]]
In [38]: find_closest_n(word_to_angle('terrible'), 10)
Out[38]: [('terrible', -0.6383390212912412),
          ('sucks', -0.6361462254806416),
          ('avoid', -0.6407294239813976),
          ('skip', -0.644470388637383),
          ('instead', -0.631978946745676),
          ('redeeming', -0.631675834597059),
          ('d', -0.6453269230869975),
          ('annoying', -0.645878152746699),
          ('save', -0.6298893756519642),
          ('zero', -0.6290363901351929)]
In [ ]:
```