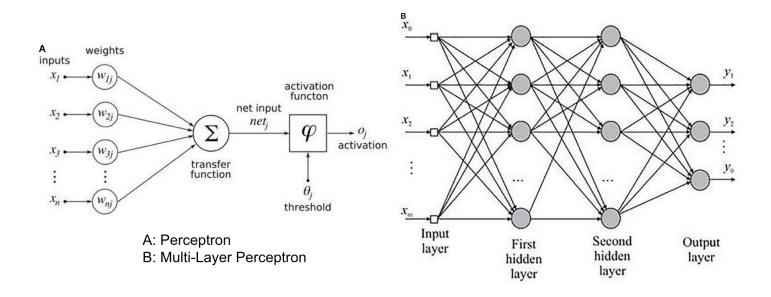
BMEN 4470 - Deep Learning for Biomedical Signal

Processing Homework 1: Multi-Layer Perceptrons and ARIMA Model for Time Series Forecasting

Due 11:59pm on October 4th, 2020

Problem 1a: We've discussed in class about the perceptron, or 'artificial neuron', the simplest artifical neural unit of processing that enables machines to learn linear mappings between inputs and outputs, as shown below. Here, we will implement a Multi-Layer Perceptron (MLP), which enables *non-linear* mappings, and will use it to predict the next number in a sequence.



```
X.append(seq x)
        y.append(seq y)
    return np.array(X), np.array(y)
# Define input sequence
raw seq = [10, 20, 30, 40, 50, 60, 70, 80, 90]
# choose a number of time steps
n \text{ steps} = 3
# split into samples
X, y = split_sequence(raw_seq, n_steps)
# summarize the data
for i in range(len(X)):
    print(X[i], y[i])
     [10 20 30] 40
     [20 30 40] 50
     [30 40 50] 60
     [40 50 60] 70
     [50 60 70] 80
     [60 70 80] 90
```

YOUR TASK: Define a model using Keras Sequential API that correctly predicts the output given a new input sequence of 3 numbers, train your model, plot the model training loss, and demonstrate a sample prediction. Provide comments to explain each line of your model. What is the impact on the accuracy of your prediction when you have no hidden layers (perceptron) vs. at least 1 hidden layer (MLP)?

```
#Some Imports to help you as a start; feel free to add more as needed
from keras.models import Sequential
from keras.layers import Activation, Dense, Input
import tensorflow as tf
from tensorflow import keras
from tensorflow.keras import layers
from tensorflow.keras.optimizers import Adam
from tensorflow.keras import activations
from tgdm import tgdm
# Your code here: define the model
model = Sequential(
    [
        Input(shape=(n steps,)),
        Dense(units=128),
        Activation(activations.relu),
        Dense(units=1),
```

Model: "sequential"

Layer (type)	Output	Shape	Param #
dense (Dense)	(None,	128)	512
activation (Activation)	(None,	128)	0
dense_1 (Dense)	(None,	1)	129
Total params: 641 Trainable params: 641			

Non-trainable params: 0

```
# Your code here: train the model via the Keras fit function
model.compile(
    optimizer=Adam(learning rate=1e-2),
    loss='mse'
    )
history = model.fit(X, y,
                    epochs=300,
                    verbose = False)
# The second model is the same as the first one,
# but with an Early Stopping Callback
callback = tf.keras.callbacks.EarlyStopping(monitor='loss', patience=3)
model.compile(
      optimizer=Adam(learning rate=1e-2),
      loss='mse'
history es = model.fit(X, y,
                      epochs=300,
                      verbose = False,
                      callbacks=[callback])
```

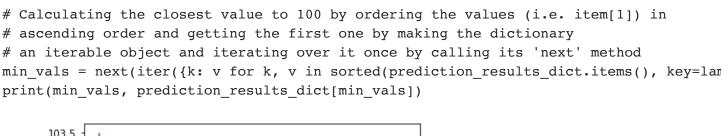
#Plot the training loss using information from the above model training step import matplotlib.pyplot as plt

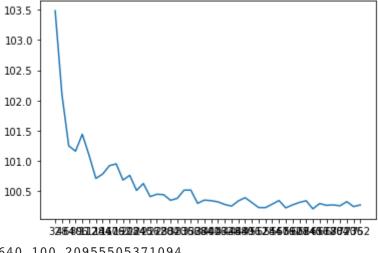
```
loss = history.history['loss']
plt.plot(history.history['loss'], label='Train Loss')
plt.title('Model Loss')
plt.ylabel('Loss')
plt.xlabel('Epochs')
plt.legend(loc='upper right')
plt.show()
# Introducing another plot to show how the graph looks if we introduced
# an early stopping callback during the training
loss es = history es.history['loss']
plt.plot(history_es.history['loss'], label='Train Loss with ES')
plt.title('Model Loss with ES')
plt.ylabel('Loss with ES')
plt.xlabel('Epochs with ES')
plt.legend(loc='upper right')
plt.show()
```

Model Loss # Your code here: Demonstrate Prediction using model.predict $x_{input} = np.array([70, 80, 90]) # predict the output for this input$ x input = x input.reshape((1, n steps)) # With Early Stopping print(model.predict(x input)) [[84.60964]] # Without Early stopping print(model.predict(x input)) WARNING: tensorflow: 5 out of the last 6 calls to <function Model.make predict func [[100.0739]] # Here we'll analyze the model deeper by adjusting the number of # Dense layer perceptrons that we can put into the model dense_layer_neurons_list = [16*i for i in range (2, 48)] print(dense layer neurons list) x input = np.array([70, 80, 90])x input = x input.reshape((1, n_steps)) # Defining a dictionary to keep the predictions prediction results dict = {} for dense layer neurons in tqdm(dense layer neurons list): # Iterating 5 times to get the average so we can be sure that we're not # getting a random value current predict vals = [] for i in range(5): model = Sequential([Input(shape=(n steps,)), Dense(units=dense layer neurons), Activation(activations.relu), Dense(units=1),] model.compile(optimizer=Adam(learning rate=1e-2), loss='mse')

history = model.fit(X, y,

```
epochs=300,
                      verbose = False)
    current predict vals.append(model.predict(x input)[0][0])
  prediction_results_dict[str(dense_layer_neurons)] = sum(current_predict_vals) / 5.0
print(prediction_results_dict)
    [32, 48, 64, 80, 96, 112, 128, 144, 160, 176, 192, 208, 224, 240, 256, 272, 288,
                    0/46 [00:00<?, ?it/s]WARNING:tensorflow:6 out of the last 11 cal
                     46/46 [04:38<00:00, 6.05s/it]{'32': 103.48425598144532, '48':
    100%
x, y = zip(*prediction_results_dict.items())
plt.plot(x, y)
plt.show()
# Calculating the closest value to 100 by ordering the values (i.e. item[1]) in
```





640 100.20955505371094

Here we can see that the best approximation we get at 640 epoxhs and the general trend is downwards until 336 epochs and then the graph becomes to almost oscillate with only just a tiny bit of a trend to go down towards 100. However, we can almost safely assume that after the graph reaches 800+ epochs the neural network starts to overfit.

Problem 1b: What happens when you try to predict the next number for the following test sequence: 1², 2², 3²? Does the model you developed in Part 1a work here? Explain.

```
# Your code here: Demonstrate Prediction using model.predict
x_input = np.array([1, 4, 9]) # predict the output for this input
x_input = x_input.reshape((1, n_steps))
print(model.predict(x_input))

[[13.999823]]
```

Explaination

The model was unable to predict the expected value - 16 - because the initial training has most likely produced a function that behave linear-like and here we need a quadratic function

Having said this, the 'new function' we are seeking requires us to either evaluate the next number by using $(n+1)^2$ where n^2 is the last number in the sequence, or by using 1 + 3 + 5 + 7 + ... + (2n+1) which is essentially the same thing since there is a quadratic representation of this sum which we can obtain if we start grouping every i-th number with every (m-i+1)-th number where m is the total number of sequence members.

The prediction that we got from the model, 13.9998, is actually a good prediction for the training set that we introduced our model to. This is because, we can see that if we analyze the difference between each of the sequence members.

```
1. |4 - 1| = 3
```

$$2. |9 - 4| = 5$$

$$3. |16 - 9| = 7$$

If we are trying to get a linearized function that connects the first two numbers to a third one, we would most likely expect the third number to be some kind of a mean of the first two (either arithmetic mean, geometric mean, quadratic/cubic mean, or maybe even harmonic mean). Given the fact that the answer we got is 13.9998 and the cubic mean (which we know is one of the biggest means) is approx. 4.2358, we can argue that our model is actually predicting as good as we expect it to do since we see that there is still a significant offset - by approx. |13.9998 - (9 + 4.2358)| = 0.764 > 3/4. This tells us that the model has recognized the introduced irregularity (the fact that the differences between the 1st and 2nd number and the 2nd and third number are not the same) and tried to adjust to it.

Problem 2a: Repeat Problem #1, but this time train your model to predict the next number in the sequence of squares (n^2 , where n = 1, 2, 3, ...). Also, this time use Keras Functional API to build your model.

```
raw_seq = [1, 4, 9, 16, 25, 36, 49, 64, 81]
# choose a number of time steps
n \text{ steps} = 3
# split into samples
X, y = split_sequence(raw_seq, n_steps)
# summarize the data
for i in range(len(X)):
   print(X[i], y[i])
    [1 4 9] 16
    [ 4 9 16] 25
    [ 9 16 25] 36
    [16 25 36] 49
    [25 36 49] 64
    [36 49 64] 81
#Some Imports to help you as a start; feel free to add more as needed
from keras.models import Sequential
from keras.layers import Activation, Dense
from keras import optimizers, Input, Model
# Your code here: define the model
input layer = Input(shape=(n steps,))
dense layer = Dense(units=128, activation='relu')(input layer)
output layer = Dense(units=1)(dense layer)
model squares = Model(inputs=input layer,
                       outputs=output layer)
model squares.summary()
```

Model: "model 2"

Layer (type)	Output Shape	Param #
input_4 (InputLayer)	[(None, 3)]	0
dense_6 (Dense)	(None, 128)	512
dense_7 (Dense)	(None, 1)	129
	:============	

Total params: 641 Trainable params: 641 Non-trainable params: 0

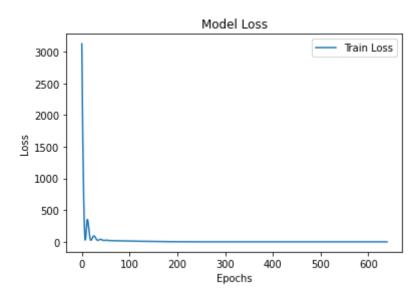
Your code here: train the model via the Keras fit function model squares.compile(optimizer=Adam(learning_rate=1e-2),

```
loss='mse'
)
```

verbose=False)

#Plot the training loss using information from the above model training step import matplotlib.pyplot as plt

```
loss = history_squares.history['loss']
plt.plot(history_squares.history['loss'], label='Train Loss')
plt.title('Model Loss')
plt.ylabel('Loss')
plt.xlabel('Epochs')
plt.legend(loc='upper right')
plt.show()
```



```
# Your code here: Demonstrate Prediction using model.predict; feel free to show
# additional examples than the one already provided here
x_input = np.array([49, 64, 81]) # predict the ooutput for this input
x_input = x_input.reshape((1, n_steps))

print(model_squares.predict(x_input))

[[101.22513]]
```

Since we're expecting a value of a 100 and we got 101.22513, we can safely assume that this value is close enough since the error is of order 10⁽⁻²⁾ which is negligible for a neural network of the

most basic kind (MLP), which has only one deep layer with more than 1 perceptron.

However, let's do a couple of more predictions so we can see how the model performs in different cases:

```
predictions_dict = {}
# Another one whole number test case
x_{input} = np.array([64, 81, 100])
x input = x input.reshape((1, n steps))
predictions_dict['whole-number'] = model_squares.predict(x_input)
# A rational number test case
x input = np.array([1.5**2, 2.5**2, 3.5**2])
x_input = x_input.reshape((1, n_steps))
predictions_dict['rational-number'] = model_squares.predict(x_input)
# Real number test case
x_input = np.array([
    (2**(1/4))**2,
    (2**(1/4) + 1)**2,
    (2**(1/4) + 2)**2
  1)
x_input = x_input.reshape((1, n_steps))
predictions dict['real-number'] = model squares.predict(x input)
expected results = [
                      # the expected whole number result
    121,
    4.5**2,
                     # the expected rational number result
    (2**(1/4) + 3)**2 # the expected real number result
]
for i, (key, value) in enumerate(predictions dict.items()):
  print(f'Predicted val: {key} -> {value}')
  print(f'Expected val: {expected results[i]}\n')
    Predicted val: whole-number -> [[123.75958]]
    Expected val: 121
    Predicted val: rational-number -> [[20.366497]]
    Expected val: 20.25
    Predicted val: real-number -> [[17.744862]]
    Expected val: 17.54945625238942
```

If we analyze the results above, we can see that our model handles every case very well since we have not provided it with training data for the second and third case we evaluated and the predictions are still very close to the expected value.

If we calculate the error:

- 1. 123.75958 / 121 = 1.022806; |1.022806 1| = 0.022806 = 2.2806 %
- 2. 20.366497 / 20.25 = 1.005753; |1.005753 1| = 0.005753 = 0.5753 %
- 3. 17.744862 / 17.54945625238942 = 1.011135; |1.011135 1| = 0.011135 = 1.1135 %

we can see that the error of the 'edge cases' is even lower than the case we would normally expect to train our model for. This means that the model is not overfitted and the error is below 2.5% which is very much acceptable.

Problem 2b: How many perceptrons (neurons) are there in the Multi-Layer Perceptron (MLP) model you defined in the problem 2a? How many weights? And how many learnable parameters? Please show your calculation and explain briefly.

The model we used to predict the sequences in the previous 2 tasks included the following 3 layers:

- 1. An input layer with 3 perceptrons
- 2. A dense layer with 128 perceptrons
- 3. An output dense layer with 1 perceptron

(The ReLu activation has no perceptrons since it's basically only a function)

=> The total number of perceptrons is: 3 + 128 + 1 = 132 perceptrons

Now, the total number of parameters are:

- 1. Graph edges between first 2 layers: Input_layer x Dense_layer_1 = 3 x 128 = 384
- 2. Graph edges between second 2 layers: Dense_layer_1 x Dense_layer_2 = 128 x 1 = 128
- 3. Biases for first dense layer: 128
- 4. Biases for second dense layer: 1
- => The total number of learnable parameters is: 384 + 128 + 128 + 1 = 641

The answer we got is the same as the automatically calculated number at the end of the models' summaries.

Problem 3: Now we will analyze the Medical Expenditures Dataset (dataSheet2.csv). [Refer to https://www.frontiersin.org/articles/10.3389/fdata.2020.00004/full for more details on dataset and problem.] This dataset contains average daily expenditures on a medicine. We will break up this

dataset into train and test sets and design a Multi-Layer Perceptron to forecast average daily

```
#MOUNTING GOOGLE DRIVE WHERE DATA IS STORED
from google.colab import drive
drive.mount('/gdrive', force_remount=True)
```

Mounted at /qdrive

print(train.shape)

#GET THE DATA (Remember to change the path to the location of your saved file!)
import pandas as pd
data = pd.read_csv("/gdrive/MyDrive/BMEN_4470_Fall_2021/HW1/dataSheet2.csv")
data.head()

Date Average Daily Expenditure on Medicine B in Dollars 0 1/2/2011 114.874671 1 1/3/2011 112.041149 2 1/4/2011 167.311618 3 1/5/2011 241.058332 4 1/6/2011 124.520457

```
import numpy as np
#JUST GET THE MEDICINE EXPENDITURE VALUES
data1 = data["Average Daily Expenditure on Medicine B in Dollars"]
data1Arr = np.array(data1)
print(data1.shape)
print(data1Arr.shape)
    (1565,)
    (1565,)
#DIFFERENT WAY OF SPLITTING DATA INTO INPUT/OUTPUT SEQUENCES
def convert2matrix(data arr, look back):
X, Y = [], []
 for i in range(len(data arr)-look back):
  d=i+look back
 X.append(data arr[i:d])
  Y.append(data arr[d])
 return np.array(X), np.array(Y)
#Split data set into testing dataset and train dataset
train size = int(len(data1Arr) * 0.8) #80% training, 20% testing
train = data1Arr[0:train size]
test = data1Arr[train size:len(data1Arr)]
```

```
print(test.shape)
# setup look back window
look\ back = 7
#convert dataset into right shape in order to input into the DNN
trainX, trainY = convert2matrix(train, look back)
testX, testY = convert2matrix(test, look back)
#Print a few of the samples to check format
for i in range(5):
  print(trainX[i], trainY[i])
    (1252,)
    (313,)
    [114.8746706 112.0411489 167.3116175 241.0583317 124.5204571 113.8333722
     149.25554531 159.0992442
    [112.0411489 167.3116175 241.0583317 124.5204571 113.8333722 149.2555453
     159.09924421 187.9188007
    [167.3116175 241.0583317 124.5204571 113.8333722 149.2555453 159.0992442
     187.91880071 138.774122
    [241.0583317 124.5204571 113.8333722 149.2555453 159.0992442 187.9188007
     138.774122 ] 176.4914938
    [124.5204571 113.8333722 149.2555453 159.0992442 187.9188007 138.774122
     176.4914938] 165.71289480000002
```

YOUR TASK: Design an MLP taking into account what you have learned from Problems #1 and #2 that achieves as low RMSE (Root Mean Square Error) as possible. Consider the impact of number of look-back time-steps. Experiment with and comment on results of changing 3 of the following parameters, providing reasons for your observations.

- 1. Number of hidden layers
- 2. Number of neurons per layer
- 3. Activation Function
- 4. Number of epochs
- 5. Batch size
- 6. Optimizer/Learning Rate

You will be graded on your analysis of these factors, not on the accuracy (RMSE) you achieve. Show plots of training/test loss for your 3 chosen experiments and a plot of actual vs. prediction for your best model. Does the same MLP that you designed for Problems #1 and #2 work well for Problem #3?

```
# Imports
from keras.models import Sequential
from keras.layers import Activation, Dense, Dropout
from keras import optimizers
from tgdm import tgdm
```

```
# YOUR CODE HERE: define model & train model
# Since we're restricted to only using Dense layers since we're creating an MLP
# we will create a for loop which will change different hyperparameters
# every time it iterates and we will evaluate the RMSE every time so we can
# find out which models perform better, which models perform worse and try to
# find a trend that explains the results we get
# Defining dictionaries where we'll keep the histories & trained models
# so we can plot them afterwards
histories_dict = {}
models_dict = {}
# Defining the possible hyperparameters -> the number of layers and
# the number of epochs for the model
number_of_layers = [2, 3, 4, 5]
number of epochs = [300, 400, 500]
for num of layers in tqdm(number of layers):
  # Defining the number of neurons per layer
  neurons per layer = [2**i for i in range(5, 5+num of layers)]
 # Defining the model by giving it an Input layer,
 # several dense layers, and a final 1-perceptron dense layer
 model = keras.Sequential(name=f'{num of layers}-layers-model')
  model.add(keras.Input(shape=(look back,)))
  [model.add(layers.Dense(num neurons, activation="relu")) for num neurons in neurons
  model.add(layers.Dense(1))
 model.compile(loss='mse',
                optimizer="adam")
  for num of epochs in number of epochs:
    dict key = f'{num of layers}-layers-{num of epochs}-epochs-model'
   histories dict[dict key]=model.fit(trainX, trainY,
                                       epochs=num of epochs,
                                       verbose=False,
                                       validation data=(testX,
                                                        testY))
   models dict[f'{num of layers}-layers-{num of epochs}-epochs-model'] = model
# Now let's recreate the same neural network but with tanh as the first
# activation function as opposed to just the normal ReLu function
for num of layers in tqdm(number of layers):
  neurons per layer = [2**i for i in range(6, 5+num of layers)]
 model = keras.Sequential(name=f'{num of layers}-layers-model')
  model.add(keras.Input(shape=(look_back,)))
```

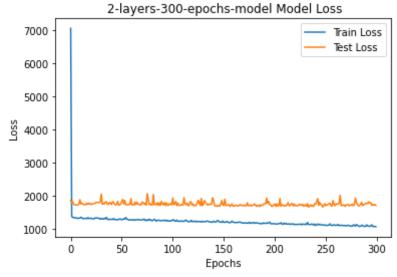
```
# Adding the tanh layer
  model.add(layers.Dense(2**5, activation="tanh"))
  # And continuing to build the model the same way it was previously built
  [model.add(layers.Dense(num neurons, activation="relu")) for num neurons in neurons
  model.add(layers.Dense(1))
  model.compile(loss='mse',
                optimizer="adam")
  for num of epochs in number of epochs:
    dict key = f'tanh-{num of layers}-layers-{num of epochs}-epochs-model'
    histories dict[dict key] = model.fit(trainX, trainY,
                                          epochs=num of_epochs,
                                          verbose=False,
                                          validation data=(testX,
                                                           testY))
    models dict[dict key] = model
# Lastly let's add the model we used for tasks 1 and 2 but with different
# learning rates for the optimizer
# 1) The same model as for tasks 1 and 2
model = Sequential(
    [
        Input(shape=(look back,)),
        Dense(units=128),
        Activation(activations.relu),
        Dense(units=1),
    ]
)
model.compile(
    optimizer=Adam(learning rate=1e-2),
    loss='mse'
    )
history = model.fit(trainX, trainY,
                    epochs=300,
                    verbose=False,
                    validation data=(testX,
                                     testY))
histories dict['previous-model-10**(-2)'] = history
models dict['previous-model-10**(-2)'] = model
# 2) The same model as for tasks 1 and 2 but with a different learning rate
model = Sequential(
        Input(shape=(look back,)),
        Dense(units=128),
        Activation(activations.relu),
```

```
Dense(units=1),
    ]
)
model.compile(
    optimizer=Adam(learning rate=1e-4),
    loss='mse'
    )
history = model.fit(trainX, trainY,
                    epochs=300,
                    verbose=False,
                    validation_data=(testX,
                                     testY))
histories dict['previous-model-10**(-4)'] = history
models dict['previous-model-10**(-4)'] = model
                    4/4 [09:39<00:00, 144.86s/it]
                    || 4/4 [10:05<00:00, 151.37s/it]
# Let's see what we have in our history dictionary finally
[print(key) for key in histories dict.items()]
    ('2-layers-300-epochs-model', <keras.callbacks.History object at 0x7f20e230d3d0>
    ('2-layers-400-epochs-model', <keras.callbacks.History object at 0x7f20e65f0e10>
    ('2-layers-500-epochs-model', <keras.callbacks.History object at 0x7f20e61a6cd0>
    ('3-layers-300-epochs-model', <keras.callbacks.History object at 0x7f20e6460a50>
    ('3-layers-400-epochs-model', <keras.callbacks.History object at 0x7f20e67f5a10>
    ('3-layers-500-epochs-model', <keras.callbacks.History object at 0x7f20e6817cd0>
    ('4-layers-300-epochs-model', <keras.callbacks.History object at 0x7f20e67dce50>
    ('4-layers-400-epochs-model', <keras.callbacks.History object at 0x7f20e62eccd0>
    ('4-layers-500-epochs-model', <keras.callbacks.History object at 0x7f20e68bbad0>
    ('5-layers-300-epochs-model', <keras.callbacks.History object at 0x7f20e6876850>
    ('5-layers-400-epochs-model', <keras.callbacks.History object at 0x7f20e6535510>
    ('5-layers-500-epochs-model', <keras.callbacks.History object at 0x7f20e292d910>
    ('tanh-2-layers-300-epochs-model', <keras.callbacks.History object at 0x7f20e62a
    ('tanh-2-layers-400-epochs-model', <keras.callbacks.History object at 0x7f20e63a
    ('tanh-2-layers-500-epochs-model', <keras.callbacks.History object at 0x7f20e29f
    ('tanh-3-layers-300-epochs-model', <keras.callbacks.History object at 0x7f20e29a
    ('tanh-3-layers-400-epochs-model', <keras.callbacks.History object at 0x7f20e628
    ('tanh-3-layers-500-epochs-model', <keras.callbacks.History object at 0x7f20d8c6
    ('tanh-4-layers-300-epochs-model', <keras.callbacks.History object at 0x7f20d8c8!
    ('tanh-4-layers-400-epochs-model', <keras.callbacks.History object at 0x7f20db02
    ('tanh-4-layers-500-epochs-model', <keras.callbacks.History object at 0x7f20db00
    ('tanh-5-layers-300-epochs-model', <keras.callbacks.History object at 0x7f20db040
    ('tanh-5-layers-400-epochs-model', <keras.callbacks.History object at 0x7f20daebo
    ('tanh-5-layers-500-epochs-model', <keras.callbacks.History object at 0x7f20dae8:
    ('previous-model-10**(-2)', <keras.callbacks.History object at 0x7f20dad9d750>)
    ('previous-model-10**(-4)', <keras.callbacks.History object at 0x7f20d8ec6f50>)
    [None,
     None,
```

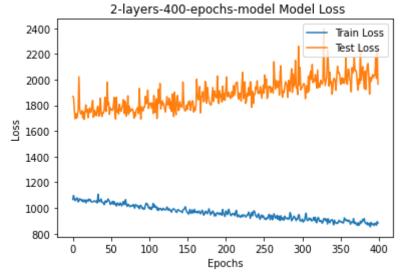
```
None,
None]
```

```
#PLOT THE TRAINING LOSS
import matplotlib.pyplot as plt
for history key in sorted(histories dict.keys()):
  print(f'History for: {history key}')
 history = histories dict[history key]
  loss = history.history['loss']
  val loss = history.history['val loss']
 plt.plot(history.history['loss'], label='Train Loss')
 plt.plot(history.history['val loss'], label='Test Loss')
 plt.title(f' {history key} Model Loss')
 plt.ylabel('Loss')
 plt.xlabel('Epochs')
 plt.legend(loc='upper right')
 plt.show();
```

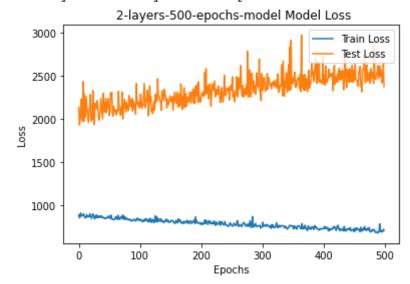
History for: 2-layers-300-epochs-model



History for: 2-layers-400-epochs-model

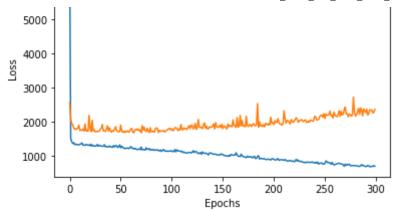


History for: 2-layers-500-epochs-model

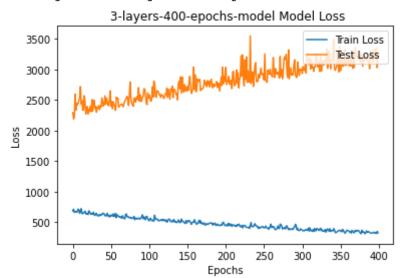


History for: 3-layers-300-epochs-model

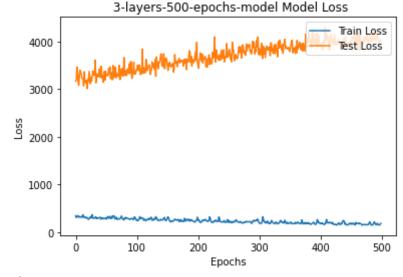




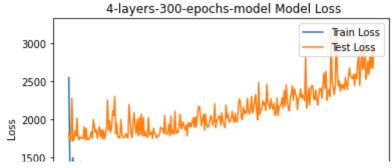
History for: 3-layers-400-epochs-model

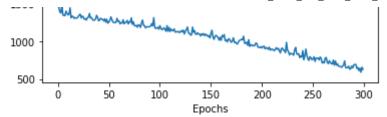


History for: 3-layers-500-epochs-model

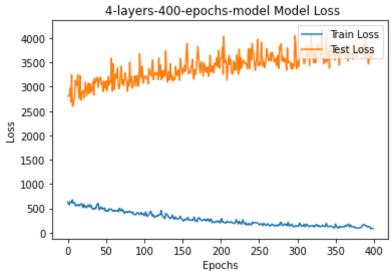


History for: 4-layers-300-epochs-model

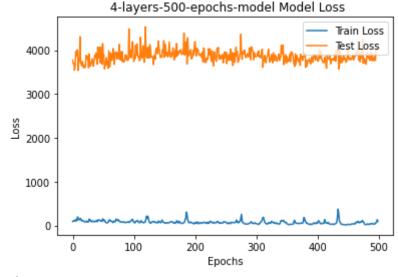




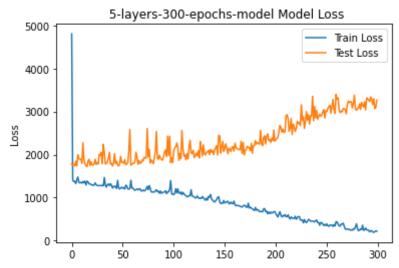
History for: 4-layers-400-epochs-model



History for: 4-layers-500-epochs-model

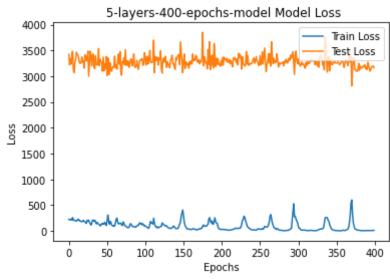


History for: 5-layers-300-epochs-model

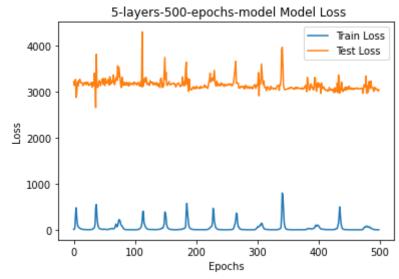


Epochs

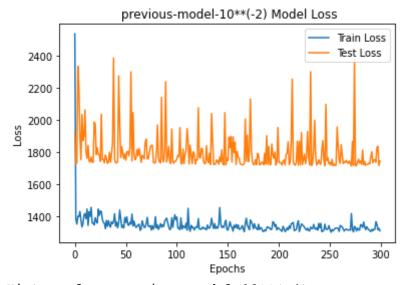
History for: 5-layers-400-epochs-model



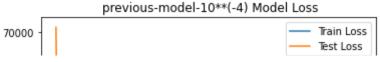
History for: 5-layers-500-epochs-model

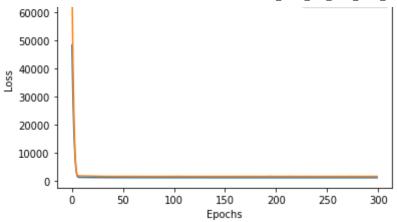


History for: previous-model-10**(-2)

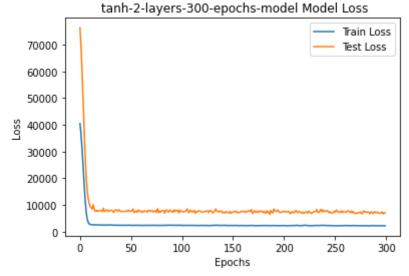


History for: previous-model-10**(-4)

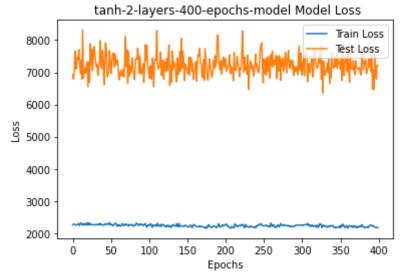




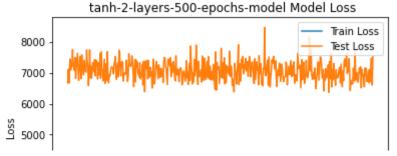
History for: tanh-2-layers-300-epochs-model

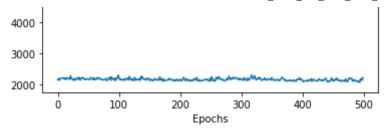


History for: tanh-2-layers-400-epochs-model

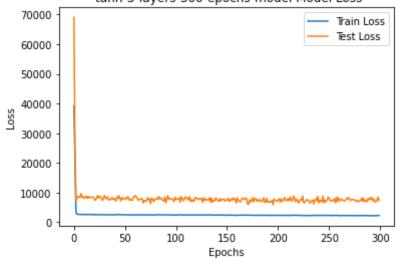


History for: tanh-2-layers-500-epochs-model

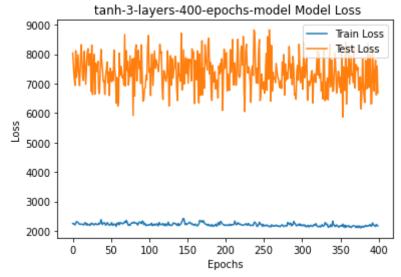




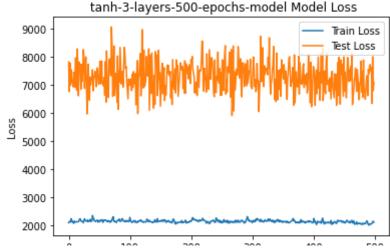
History for: tanh-3-layers-300-epochs-model tanh-3-layers-300-epochs-model Model Loss



History for: tanh-3-layers-400-epochs-model



History for: tanh-3-layers-500-epochs-model

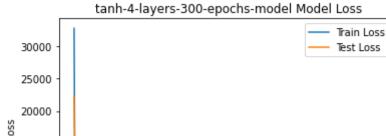


Epochs

History for: tanh-4-layers-300-epochs-model

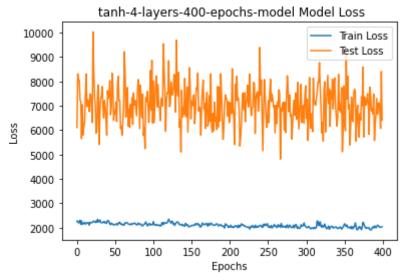
200

100

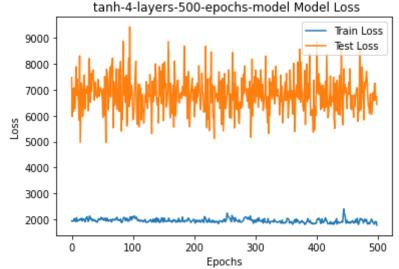


15000 - 10000

History for: tanh-4-layers-400-epochs-model



History for: tanh-4-layers-500-epochs-model

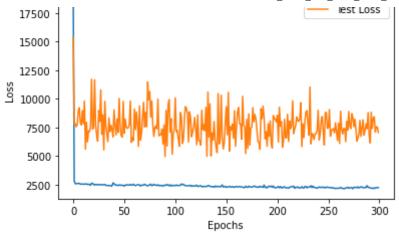


History for: tanh-5-layers-300-epochs-model

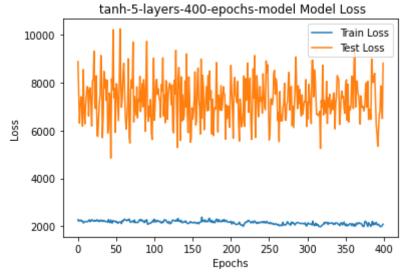
20000

tanh-5-layers-300-epochs-model Model Loss

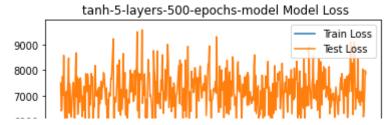
— Train Loss



History for: tanh-5-layers-400-epochs-model



History for: tanh-5-layers-500-epochs-model



#MODEL EVALUATION via RMSE

```
# Now we'll iterate over the all the models and we will sort them by average
# RMSE from their train and test score

# Defining a dict where we'll keep the RMSE for each model
rmse_dict = {}

for model_key in sorted(models_dict.keys()):
   model = models_dict[model_key]

   train_score = model.evaluate(trainX, trainY, verbose=0)
   test_score = model.evaluate(testX, testY, verbose=0)

rmse_dict[model_key] = {
    'train_score': train_score,
```

```
'test score': test score
  }
# Sorting the dictionary and printing out the results
for key, value in sorted(rmse_dict.items(),
                         key=lambda item: item[1]['train score'] + item[1]['test score']
  print(f'Model evaluation for: {key}')
  train score = value['train score']
  test_score = value['test_score']
  print(f'Train Root Mean Squared Error(RMSE): {train score}')
  print(f'Test Root Mean Squared Error(RMSE): {test score}')
  # Empty line for readibility
  print()
    Test Root Mean Squared Error(RMSE): 4025.065673828125
    Model evaluation for: 3-layers-400-epochs-model
    Train Root Mean Squared Error(RMSE): 158.07066345214844
    Test Root Mean Squared Error(RMSE): 4025.065673828125
    Model evaluation for: 3-layers-500-epochs-model
    Train Root Mean Squared Error(RMSE): 158.07066345214844
    Test Root Mean Squared Error(RMSE): 4025.065673828125
    Model evaluation for: tanh-4-layers-300-epochs-model
    Train Root Mean Squared Error(RMSE): 1742.6588134765625
    Test Root Mean Squared Error(RMSE): 6432.33935546875
    Model evaluation for: tanh-4-layers-400-epochs-model
    Train Root Mean Squared Error(RMSE): 1742.6588134765625
    Test Root Mean Squared Error(RMSE): 6432.33935546875
    Model evaluation for: tanh-4-layers-500-epochs-model
    Train Root Mean Squared Error(RMSE): 1742.6588134765625
    Test Root Mean Squared Error(RMSE): 6432.33935546875
    Model evaluation for: tanh-3-layers-300-epochs-model
    Train Root Mean Squared Error(RMSE): 2116.103515625
    Test Root Mean Squared Error(RMSE): 7093.88623046875
    Model evaluation for: tanh-3-layers-400-epochs-model
    Train Root Mean Squared Error(RMSE): 2116.103515625
    Test Root Mean Squared Error(RMSE): 7093.88623046875
    Model evaluation for: tanh-3-layers-500-epochs-model
    Train Root Mean Squared Error(RMSE): 2116.103515625
    Test Root Mean Squared Error(RMSE): 7093.88623046875
    Model evaluation for: tanh-5-layers-300-epochs-model
    Train Root Mean Squared Error(RMSE): 1824.5821533203125
    Test Root Mean Squared Error(RMSE): 7863.2685546875
    Model evaluation for: tanh-5-layers-400-epochs-model
```

```
Train Root Mean Squared Error(RMSE): 1824.5821533203125
Test Root Mean Squared Error(RMSE): 7863.2685546875

Model evaluation for: tanh-5-layers-500-epochs-model
Train Root Mean Squared Error(RMSE): 1824.5821533203125
Test Root Mean Squared Error(RMSE): 7863.2685546875

Model evaluation for: tanh-2-layers-300-epochs-model
Train Root Mean Squared Error(RMSE): 2190.031005859375
Test Root Mean Squared Error(RMSE): 7515.4912109375

Model evaluation for: tanh-2-layers-400-epochs-model
Train Root Mean Squared Error(RMSE): 2190.031005859375
Test Root Mean Squared Error(RMSE): 7515.4912109375

Model evaluation for: tanh-2-layers-500-epochs-model
Train Root Mean Squared Error(RMSE): 2190.031005859375
Test Root Mean Squared Error(RMSE): 7515.4912109375
```

As we can see, there actually is a general trend which we can identify.

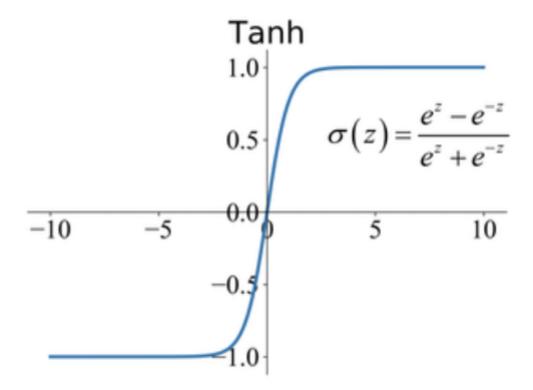
All the models can be distinguished into 3 groups:

```
> a) The 2 models which resemble the model from task 1 and 2> b) The models which have only ReLu activation functions> c) The models which have a Tanh activation function as their first activation
```

There is a clear trend that the **a**) models are doing significantly better than the **b**) and **c**) models. Since simple calculations can easily show that the **a**) models have much less trainable parameters than the other models, we can safely assume that the results are probably because of overfitting. If we look at the dataset, it consists of the simplest possible data architecture -> i.e. a single value input and a single value output. This would make the model more inclined towards overfitting since after a significant amount of epichs, it can just memorize the inputs and outputs as opposed to properly predicting new data.

This theory can be further confirmed by looking at the loss graphs for the models. We can recognise that from group **b**) there is only one model (2 layers, 300 epochs, i.e. the simplest one) for which the loss function does not start to diverge greatly. For all other models from that group we can clearly see the difference in train and test loss values rises, meaning that the model, indeed, overfits. As opposed to the **b**) models, the trend in the **c**) models is such that the average difference in loss values stays the same, but it's in the range [5000; 9000], which is far worse. This means that the tanh activation function makes the prediction less accurate. This is probably connected to the previous reason - i.e. the overfitting. This assumption is plausible because of the nature of the 2

opposing activation functions. The ReLu function "loses" more information because it sets values to 0. The tanh function:



does not nullify any weights (expect in the rare case sigma=0) meaning that it forwards information that will have been set to 0 if the activation was a ReLu. In our case, that "helps the overfitting" since the model is already overfitted and by providing it with even more parameters to work with, it becomes even worse. This can be seen also by the training time, which is (as expected) more for the **c**) models, because the training needs more time to account for the more non-nullified parameters.

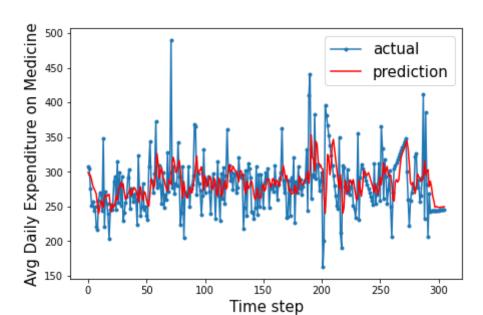
In conclusion, both the graphs and our analysis show that the a) models are the best.

```
# Demonstrate Prediction and Create Plot to Compare Prediction to Actual
short = 313 #how much of the test sequence to plot (currently: all of it)

# Choosing the best model accoring to our previous output
model = models_dict['previous-model-10**(-4)']
test_predict = model.predict(testX)

def prediction_plot(testY, test_predict):
    len_prediction=[x for x in range(len(testY[:short]))]
    plt.plot(len_prediction, testY[:short], marker='.', label="actual")
    plt.plot(len_prediction, test_predict[:short], 'r', label="prediction")
    plt.tight_layout()
    plt.subplots_adjust(left=0.07)
```

```
plt.ylabel('Avg Daily Expenditure on Medicine', size=15)
plt.xlabel('Time step', size=15)
plt.legend(fontsize=15)
plt.show();
prediction plot(testY, test predict)
```



Problem 4: Now let's repeat Problem #3 using a non-DL model for comparison. Use an ARIMA Model for time-series forecasting on the medicine expenditures dataset. Your task is to experiment with the lag order (p), degree of differencing (d), and order of the moving average (q). Discuss the impact of these factors on your final prediction error. How does this model compare with your MLP model from Problem #3? Which one is better? Why? Why are they both still not "good enough"?

```
from statsmodels.tsa.arima_model import ARIMA
from sklearn.metrics import mean_squared_error

#ARIMA model training and prediction (NOTE: training may take ~10min)
history = [x for x in train]
p, d, q = 2, 1, 2

predictions = list()
for t in range(len(test)):
    model = ARIMA(history, order=(p, d, q)) # <--- YOUR TASK HERE: modify p, d, q
    model_fit = model.fit(disp=0)
    output = model_fit.forecast()
    yhat = output[0]
    predictions.append(yhat)
    obs = test[t]
    history.append(obs)
    #print('predicted=%f, expected=%f' % (yhat, obs))</pre>
```

```
error = mean squared error(test, predictions)
print('Test MSE: %.3f' % error)
    /usr/local/lib/python3.7/dist-packages/statsmodels/base/model.py:492: HessianInve
       'available', HessianInversionWarning)
    /usr/local/lib/python3.7/dist-packages/statsmodels/tsa/tsatools.py:668: RuntimeWa
      newparams = ((1-np.exp(-params))/(1+np.exp(-params))).copy()
    /usr/local/lib/python3.7/dist-packages/statsmodels/tsa/tsatools.py:668: RuntimeWa
      newparams = ((1-np.exp(-params))/(1+np.exp(-params))).copy()
    /usr/local/lib/python3.7/dist-packages/statsmodels/tsa/tsatools.py:669: RuntimeWa
      tmp = ((1-np.exp(-params))/(1+np.exp(-params))).copy()
    /usr/local/lib/python3.7/dist-packages/statsmodels/tsa/tsatools.py:669: RuntimeWa
      tmp = ((1-np.exp(-params))/(1+np.exp(-params))).copy()
    /usr/local/lib/python3.7/dist-packages/statsmodels/base/model.py:512: Convergence
       "Check mle_retvals", ConvergenceWarning)
    /usr/local/lib/python3.7/dist-packages/statsmodels/base/model.py:492: HessianInve
       'available', HessianInversionWarning)
    /usr/local/lib/python3.7/dist-packages/statsmodels/base/model.py:492: HessianInv
       'available', HessianInversionWarning)
    /usr/local/lib/python3.7/dist-packages/statsmodels/base/model.py:512: Convergence
      "Check mle retvals", ConvergenceWarning)
    /usr/local/lib/python3.7/dist-packages/statsmodels/base/model.py:492: HessianInve
       'available', HessianInversionWarning)
    /usr/local/lib/python3.7/dist-packages/statsmodels/base/model.py:492: HessianInve
       'available', HessianInversionWarning)
    /usr/local/lib/python3.7/dist-packages/statsmodels/base/model.py:492: HessianInve
       'available', HessianInversionWarning)
    /usr/local/lib/python3.7/dist-packages/statsmodels/base/model.py:492: HessianInve
       'available', HessianInversionWarning)
    Test MSE: 1600.893
from matplotlib import pyplot
```

```
from matplotlib import pyplot
# plotting actual vs. ARIMA prediction
pyplot.plot(test, label='actual')
pyplot.plot(predictions, color='red', label='predictions')
pyplot.ylabel('Avg Daily Expenditure on Medicine', size=15)
pyplot.xlabel('Time step', size=15)
pyplot.legend(fontsize=15)
pyplot.show()
```

 \Box



By trying out different parameters for p, q, and d the values 2, 1, and 2 respectibely seem to give us the best possible combination of results. By running several different examples we can easily see that obtaining an MSE value below 1200 is impossible. However, comparing that to our Test Root Mean Squared Error (RMSE) results from Problem 3, we can see that this is actually a better prediction by a factor of approx. 1.125 (or 12.5%). Having said that, we can see that both models do not predict the expected values well and, instead, output some value which is only up to a standard deviation away from the middle value. Because of this, both models are "not good enough".

P.S. Visually, we can argue that the model from Problem 3 is better since it is not so fixed around the middle value.

WHAT TO SUBMIT: Please download and submit (1) this jupyter notebook (.ipynb file) with your code and comment additions as well as (2) a pdf summarizing your analysis of results (plots/explanations) for each problem on courseworks by **October 4th, 2020 at 11:59pm**.

X