Exercise sheet 10 - Introduction - Due date: July 6th

Submitted to:

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Exercise 1 - Variational Autoencoders (VAEs) (12 points)

- 1. Explain how far a VAE's lower dimension representation differs from that learned by a traditional autoencoder. How is that achieved? How far does the training objective differ? (2 points)
- An autoencoder accepts input, compresses it, and then recreates the original input. This is
 an unsupervised technique because all you need is the original data, without any labels of
 known, correct results. The two main uses of an autoencoder are to compress data to two
 (or three) dimensions so it can be graphed, and to compress and decompress images or
 documents, which removes noise in the data.
- A variational autoencoder assumes that the source data has some sort of underlying probability distribution (such as Gaussian) and then attempts to find the parameters of the distribution. The one main use of a variational autoencoder is to generate new data that's related to the original source data.
- Source: https://jamesmccaffrey.wordpress.com/2018/07/03/the-difference-between-an-autoencoder-and-a-variational-autoencoder/
- 1. Inform yourself about the applications of autoencoders in the biomedical field. Explore the literature, then mention one application and explain how it works. (2 points)

There have been lots of other use-cases of autoencoder in biology. It can be used as a dimension reduction method for unsupervised clustering and visualization, similar to principal component analysis (PCA).

Autoencoder has many connections in biology. One example is the L1000 platform developed in the Connectivity Map project [1], in which 1,000 landmark genes are sufficient to recover 81% of the information in the full transcriptome, and significantly lower the cost of transcriptome profiling. These 1,000 genes can be loosely considered as the dimension-reduced latent variables.

Source: https://encodebox.medium.com/auto-encoder-in-biology-9264da118b83

1. Inform yourself about VAE variants, then explain the modifications and uses of the following

variants:

a. Beta-VAE (1 point)

Beta-VAE is a type of variational autoencoder that seeks to discovered disentangled latent factors. It modifies VAEs with an adjustable hyperparameter that balances latent channel capacity and independence constraints with reconstruction accuracy. The idea is to maximize the probability of generating the real data while keeping the distance between the real and estimated distributions small, under a threshold

Source: https://paperswithcode.com/method/beta-vae

b. Vector Quantised-VAE (VQ-VAE) (1 point)

VQ-VAE is a type of variational autoencoder that uses vector quantisation to obtain a discrete latent representation. It differs from VAEs in two key ways: the encoder network outputs discrete, rather than continuous, codes; and the prior is learnt rather than static.

Using the VQ method allows the model to circumvent issues of posterior collapse - where the latents are ignored when they are paired with a powerful autoregressive decoder - typically observed in the VAE framework.

Source: https://paperswithcode.com/method/vq-vae

1. Load the MNIST digits dataset from Keras datasets API. Normalize all your values between 0 and 1, and flatten your images into vectors of size 784. Build a simple VAE model using the following architecture:

```
import tensorflow as tf
from tensorflow import keras
import numpy as np
from keras import layers
from keras.models import Model
from keras.layers import Input, Dense
from keras.losses import binary_crossentropy
import matplotlib.pyplot as plt
```

```
In [2]: #Loading the dataset
    mnist = tf.keras.datasets.mnist #28*28 image
    (x_train,y_train), (x_test,y_test) = mnist.load_data()

#Normalizing values
    x_train = tf.keras.utils.normalize(x_train, axis = 1)
    x_test = tf.keras.utils.normalize(x_test,axis = 1)

#Flattening images into a vector size of 784
    x_train = x_train.reshape(x_train.shape[0], 784)
    x_test = x_test.reshape(x_test.shape[0], 784)
```

- a. Encoder: 1 hidden layer with an input, using ReLU activation function (1 point)
- b. Decoder: 1 hidden layer using sigmoid activation function (1 point)

```
In [3]: encoding_dim = 32
   input_img =Input(shape=(784,)) #input image
   encoded = Dense(encoding_dim, activation='relu')(input_img) #encoder with 1 hidden L
   decoded = Dense(784, activation='sigmoid')(encoded) #decoder with 1 hidden Layer
   #building the VAE model
```

```
vae = Model(input_img, decoded)

#compiling encoder model
encoder = Model(input_img, encoded)
encoded_input = Input(shape=(encoding_dim,))

decoder_layer = vae.layers[-1]
#compiling decoder model
decoder = Model(encoded_input, decoder_layer(encoded_input))
```

Your encoder should take an image input of 784 floats and encode it to 32 floats, while the decoder should take the encoded input and reconstruct an image of 784 floats. Compile your model using the following:

- 1. Adam optimizer (0.5 point)
- 1. Binary cross entropy loss function (0.5 point)
- 1. Batch size of 256 (0.5 point)

```
In [4]: #compiling VAE model
    vae.compile(optimizer='adam', loss='binary_crossentropy')
```

Train your model using the following parameters:

1. 50 epochs (0.5 point)

Batch size of 256 (0.5 point) Use your model to predict 10 digits from the MNIST dataset, then plot the original and reconstructed images for reference. (2 points)

```
#training the model with 50 epochs and a batch size of 256
In [5]:
    vae.fit(x_train, x_train, epochs=50, batch_size=256, shuffle=True, validation_data=(
    Epoch 1/50
    0.1648
    Epoch 2/50
    0.1496
    Epoch 3/50
    235/235 [============= ] - 3s 13ms/step - loss: 0.1477 - val loss:
    0.1424
    Epoch 4/50
    235/235 [============= ] - 4s 16ms/step - loss: 0.1408 - val loss:
    0.1374
    Epoch 5/50
    0.1333
    Epoch 6/50
    235/235 [============= ] - 3s 12ms/step - loss: 0.1323 - val loss:
    0.1300
    Epoch 7/50
    0.1275
    Epoch 8/50
    0.1255
    Epoch 9/50
    0.1240
    Epoch 10/50
    235/235 [===============] - 3s 11ms/step - loss: 0.1235 - val_loss:
    0.1228
```

```
Epoch 11/50
235/235 [================== ] - 3s 13ms/step - loss: 0.1221 - val_loss:
0.1218
Epoch 12/50
1210
Epoch 13/50
1204
Epoch 14/50
1199
Epoch 15/50
0.1195
Epoch 16/50
1191
Epoch 17/50
1187
Epoch 18/50
1185
Epoch 19/50
1182
Epoch 20/50
1180
Epoch 21/50
1179
Epoch 22/50
1178
Epoch 23/50
1177
Epoch 24/50
235/235 [================== ] - 2s 8ms/step - loss: 0.1170 - val_loss: 0.
1176
Epoch 25/50
Epoch 26/50
1175
Epoch 27/50
1174
Epoch 28/50
1174
Epoch 29/50
1173
Epoch 30/50
1173
Epoch 31/50
1173
Epoch 32/50
1172
Epoch 33/50
235/235 [================== ] - 2s 8ms/step - loss: 0.1166 - val_loss: 0.
1172
```

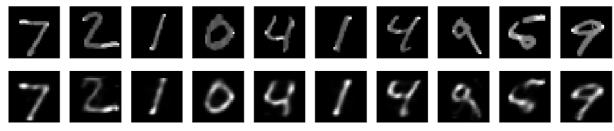
```
Epoch 34/50
  235/235 [================== ] - 2s 8ms/step - loss: 0.1166 - val_loss: 0.
  1172
  Epoch 35/50
  1172
  Epoch 36/50
  1172
  Epoch 37/50
  1171
  Epoch 38/50
  1171
  Epoch 39/50
  1171
  Epoch 40/50
  1171
  Epoch 41/50
  1171
  Epoch 42/50
  1171
  Epoch 43/50
  235/235 [================== ] - 2s 8ms/step - loss: 0.1164 - val_loss: 0.
  1171
  Epoch 44/50
  1171
  Epoch 45/50
  1171
  Epoch 46/50
  1170
  Epoch 47/50
  1170
  Epoch 48/50
  Epoch 49/50
  1170
  Epoch 50/50
  1170
Out[5]: <keras.callbacks.History at 0x7fb09c2b8460>
```

Use your model to predict 10 digits from the MNIST dataset, then plot the original and reconstructed images for reference. (2 points)

```
In [6]:
         encoded_imgs = encoder.predict(x_test)
         decoded_imgs = decoder.predict(encoded_imgs)
In [7]:
        n = 10 #Number of digits
         plt.figure(figsize=(20, 4))
         for i in range(n):
             #Original images
             ax = plt.subplot(2, n, i + 1)
             plt.imshow(x_test[i].reshape(28, 28))
             plt.grav()
             ax.get_xaxis().set_visible(False)
```

```
ax.get_yaxis().set_visible(False)

# Reconstructed images
ax = plt.subplot(2, n, i + 1 + n)
plt.imshow(decoded_imgs[i].reshape(28, 28))
plt.gray()
ax.get_xaxis().set_visible(False)
ax.get_yaxis().set_visible(False)
plt.show()
```



Exercise 2 - Generative Adversarial Networks (GANs) (6 points)

- 1. What are the specific properties of GANs in comparison to VAEs? (2 points)
 - Although VAE generative model and an inference model, and it also learns the underlying data distribution similar to GANS using unsupervised learning,

GANs yield better results as compared to VAE as:

- **1.**There is no assumption and optimizing the the lower variational bound in GAN.
- **2.**No explicit probability density estimation.
- **3.**GAN learns the true posterior distribution and so it can generate sharp images (high quality image and video generation, drug discovery) which is a failure in case of VAE. GAN are better at producing realistic images
- 2. Familiarize yourself with the following type of GANS and briefly explain how each technique differs from vanilla GANs and give an application example for each type.(4 points)
- a. Deep Convolutional GANs (DCGANs):
 - similar to GAN but uses Deep Convolutional networks in place of those fully-connected networks.
 - deep convolutional neural networks for both the generator and discriminator models and configurations for the models and training that result in the stable training of a generator model.
 - DCGAN will be more fitting for image/video data, but GAN can be applied to wider domains
 - generate images with high resolution and with less noise.
- b. Wasserstein GANs (WGANs):

- uses the 1-Wasserstein distance, rather than the JS-Divergence, to measure the difference between the model and target distributions.
- overcomes undesirable behavior of the JSD in the presence of singular measure.
- it is possible to focus on particular image features of interest.

c. Self-Attention GANs (SAGANs):

- GANs generate high-resolution details as a function of only spatially local points in lowerresolution feature maps but in SAGAN, details can be generated using cues from all feature locations.
- incorporate a selfattention mechanism into the GAN framework which is effective in modeling long-range dependencies.

d. BigGANs:

- trains GANs at large scales
- used for scaling generation to high-resolution, high-fidelity images

Exercise 3 - LSTMs & Transformers (7 points)

1. General architecture of LSTM unit. (1 point)

A common LSTM unit is composed of a cell, an input gate, an output gate and a forget gate. The cell remembers values over arbitrary time intervals and the three gates regulate the flow of information into and out of the cell.

2. What is a Bidirectional LSTM? And what are its advantages over a classical LSTM? (2 points)

- Bidirectional LSTMs are an extension of traditional LSTMs that can improve model performance on sequence classification problems.
- In problems where all timesteps of the input sequence are available, Bidirectional LSTMs train two instead of one LSTMs on the input sequence. This can provide additional context to the network and result in faster and even fuller learning on the problem.
- Using bidirectional LSTM will manage your inputs in two ways, one from past to future and
 one from future to past and it differs this approach from unidirectional is that in the LSTM
 which runs backward you preserve information from the future and using the two hidden
 states combined, you will be able at any point in time to preserve information from both
 past and future.
- 3. Both transformers and LSTMs retain memory of the generated output. What are the differences between transformers and LSTMs, and what are the advantages of transformers compared to LSTMs? (2 points)

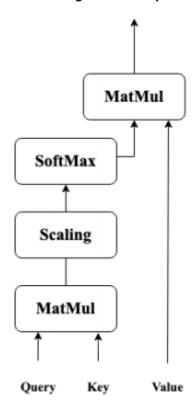
Differences between transformers and LSTMs:

• Transformer model is based on a self-attention mechanism.

• Transformers are non sequential: sentences are processed as a whole rather than word by word but LSTMs are sequential processing: sentences must be processed words by words.

Advantages of transformers compared to LSTMs:

- Transformers avoid recursion in order to allow parallel computation, LSTMs can't be trained in parallel.
- Transformers do not suffer from long dependency issues.
- 4. The image below depicts the architecture of a self-attention model.



Using the following arrays as the Query, Key and the Value vectors, calculate the attention matrix A(Q, K, V) (2 points)

• Query Vector: [0.8, 0.5, 0.21]

• Key Vector: [0.1, 0.62, 0.73]

• Value Vector: [0.6, 0.2, 0.9]

```
Output, attention weight
             matmul_qk = tf.matmul(q, k, transpose_b=True) # (..., seq_len_q, seq_len_k)
             # Zoom matmul qk
             dk = tf.cast(tf.shape(k)[-1], tf.float32)
             scaled_attention_logits = matmul_qk / tf.math.sqrt(dk)
             # Add mask to the scaled tensor.
             if mask is not None:
                 scaled_attention_logits += (mask * -1e9)
             # softmax is normalized on the last axis (seq_len_k), so the score
             # Add is equal to 1.
             attention weights = tf.nn.softmax(scaled attention logits, axis=-1) # (..., seq
             output = tf.matmul(attention_weights, v) # (..., seq_Len_q, depth_v)
             return output, attention_weights
In [9]:
         import tensorflow as tf
         Query = [0.8, 0.5, 0.21]
         Key = [ 0.1, 0.62, 0.73 ]
         Value = [ 0.6, 0.2, 0.9 ]
         scaled_dot_product_attention(Query, Key, Value, None)
        InvalidArgumentError
                                                   Traceback (most recent call last)
        <ipython-input-9-a1196ba4d211> in <module>
              4 Key = [ 0.1, 0.62, 0.73 ]
              5 Value = [ 0.6, 0.2, 0.9 ]
        ----> 6 scaled_dot_product_attention(Query, Key, Value, None)
        <ipython-input-8-19be3e4a69fb> in scaled dot product attention(q, k, v, mask)
             17
             18
                    matmul qk = tf.matmul(q, k, transpose b=True) # (..., seq len q, seq le
        ---> 19
        n_k)
             20
             21
                    # Zoom matmul qk
        /opt/anaconda3/lib/python3.8/site-packages/tensorflow/python/util/dispatch.py in wra
        pper(*args, **kwargs)
                    """Call target, and fall back on dispatchers if there is a TypeError."""
            204
            205
                    try:
        --> 206
                      return target(*args, **kwargs)
            207
                    except (TypeError, ValueError):
            208
                      # Note: convert_to_eager_tensor currently raises a ValueError, not a
        /opt/anaconda3/lib/python3.8/site-packages/tensorflow/python/ops/math_ops.py in matm
        ul(a, b, transpose_a, transpose_b, adjoint_a, adjoint_b, a_is_sparse, b_is_sparse, n
        ame)
                      return ret
           3487
           3488
                    else:
        -> 3489
                      return gen_math_ops.mat_mul(
           3490
                          a, b, transpose_a=transpose_a, transpose_b=transpose_b, name=name)
           3491
        /opt/anaconda3/lib/python3.8/site-packages/tensorflow/python/ops/gen_math_ops.py in
        mat_mul(a, b, transpose_a, transpose_b, name)
           5698
                      return _result
                    except _core._NotOkStatusException as e:
           5699
        -> 5700
                      _ops.raise_from_not_ok_status(e, name)
           5701
                    except _core._FallbackException:
           5702
                      pass
```

```
/opt/anaconda3/lib/python3.8/site-packages/tensorflow/python/framework/ops.py in rai
se_from_not_ok_status(e, name)
        6895    message = e.message + (" name: " + name if name is not None else "")
        6896    # pylint: disable=protected-access
-> 6897    six.raise_from(core._status_to_exception(e.code, message), None)
        6898    # pylint: enable=protected-access
        6899

/opt/anaconda3/lib/python3.8/site-packages/six.py in raise_from(value, from_value)
InvalidArgumentError: In[0] and In[1] ndims must be == 2: 1 [Op:MatMul]

In [ ]:
```