# C2W2\_Assignment

July 14, 2021

# 1 Breast Cancer Prediction

In this exercise, you will train a neural network on the Breast Cancer Dataset to predict if the tumor is malignant or benign.

If you get stuck, we recommend that you review the ungraded labs for this week.

## 1.1 Imports

```
[1]: import tensorflow as tf
from tensorflow.keras.models import Model
from tensorflow.keras.layers import Dense, Input

import numpy as np
import matplotlib.pyplot as plt
import matplotlib.ticker as mticker
import pandas as pd
from sklearn.model_selection import train_test_split
from sklearn.metrics import confusion_matrix
import itertools
from tqdm import tqdm
import tensorflow_datasets as tfds

tf.get_logger().setLevel('ERROR')
```

## 1.2 Load and Preprocess the Dataset

We first load the dataset and create a data frame using pandas. We explicitly specify the column names because the CSV file does not have column headers.

```
[3]: df.head()
[3]:
                                                                       marginal_adheshion
              id
                   clump_thickness
                                      un_cell_size
                                                      un_cell_shape
     0
         1000025
         1002945
                                   5
                                                   4
                                                                    4
                                                                                           5
     1
         1015425
                                   3
                                                   1
                                                                    1
     2
                                                                                            1
     3
         1016277
                                   6
                                                   8
                                                                    8
                                                                                            1
         1017023
                                   4
                                                   1
                                                                    1
                                                                                            3
         single_eph_cell_size bare_nuclei
                                                bland_chromatin
                                                                   normal_nucleoli
     0
                               2
                                                                                   1
                              7
                                                                                   2
     1
                                           10
                                                                3
     2
                               2
                                            2
                                                                3
                                                                                   1
                                                                3
     3
                               3
                                            4
                                                                                   7
     4
                                            1
                                                                3
                                                                                   1
         mitoses
                   class
     0
               1
                        2
     1
               1
                        2
                        2
     2
               1
     3
               1
                        2
     4
                        2
               1
```

We have to do some preprocessing on the data. We first pop the id column since it is of no use for our problem at hand.

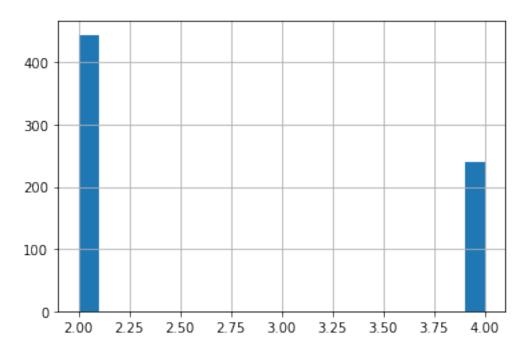
```
df.pop("id")
[4]: 0
             1000025
     1
             1002945
     2
             1015425
     3
             1016277
     4
             1017023
     694
              776715
     695
              841769
     696
              888820
     697
              897471
     698
              897471
     Name: id, Length: 699, dtype: int64
```

Upon inspection of data, you can see that some values of the **bare\_nuclei** column are unknown. We drop the rows with these unknown values. We also convert the **bare\_nuclei** column to numeric. This is required for training the model.

```
[5]: df = df[df["bare_nuclei"] != '?']
df.bare_nuclei = pd.to_numeric(df.bare_nuclei)
```

We check the class distribution of the data. You can see that there are two classes, 2.0 and 4.0 According to the dataset: \* 2.0 = benign \* 4.0 = malignant

- [6]: df['class'].hist(bins=20)
- [6]: <matplotlib.axes.\_subplots.AxesSubplot at 0x7feeaea1c190>



We are going to model this problem as a binary classification problem which detects whether the tumor is malignant or not. Hence, we change the dataset so that: \* benign(2.0) = 0 \* malignant(4.0) = 1

```
[7]: df['class'] = np.where(df['class'] == 2, 0, 1)
```

We then split the dataset into training and testing sets. Since the number of samples is small, we will perform validation on the test set.

```
[8]: train, test = train_test_split(df, test_size = 0.2)
```

We get the statistics for training. We can look at statistics to get an idea about the distribution of plots. If you need more visualization, you can create additional data plots. We will also be using the mean and standard deviation from statistics for normalizing the data

```
[11]: train_stats = train.describe()
    train_stats.pop('class')
    train_stats = train_stats.transpose()
```

```
[12]: train_stats
```

```
[12]:
                                               std min 25%
                                                             50%
                                                                   75%
                          count
                                                                         max
                                    mean
     clump_thickness
                          546.0 4.408425 2.814745
                                                   1.0
                                                        2.0
                                                             4.0
                                                                  6.00
                                                                        10.0
     un cell size
                          546.0 3.108059 3.039986 1.0 1.0
                                                             1.0
                                                                  4.00
                                                                        10.0
     un_cell_shape
                          546.0 3.183150 3.003877
                                                   1.0 1.0
                                                             1.0
                                                                  5.00
                                                                        10.0
     marginal adheshion
                          546.0 2.800366 2.875789 1.0 1.0
                                                             1.0
                                                                  3.75
                                                                        10.0
     single_eph_cell_size
                          546.0 3.214286 2.209209 1.0
                                                        2.0
                                                             2.0
                                                                  4.00
                                                                        10.0
     bare nuclei
                          546.0 3.543956 3.655195 1.0 1.0
                                                             1.0
                                                                  6.00
                                                                       10.0
     bland_chromatin
                          546.0 3.421245 2.401496 1.0 2.0
                                                             3.0
                                                                  4.00
                                                                        10.0
     normal_nucleoli
                                                             1.0
                                                                  4.00
                          546.0 2.882784 3.086074 1.0 1.0
                                                                       10.0
     mitoses
                          546.0 1.575092 1.717381 1.0 1.0
                                                             1.0
                                                                  1.00
                                                                       10.0
```

We pop the class column from the training and test sets to create train and test outputs.

```
[13]: train_Y = train.pop("class")
test_Y = test.pop("class")
```

Here we normalize the data by using the formula: X = (X - mean(X)) / StandardDeviation(X)

```
[14]: def norm(x):
    return (x - train_stats['mean']) / train_stats['std']
```

```
[17]: norm_train_X = norm(train)
norm_test_X = norm(test)
```

We now create Tensorflow datasets for training and test sets to easily be able to build and manage an input pipeline for our model.

We shuffle and prepare a batched dataset to be used for training in our custom training loop.

```
[21]: batch_size = 32
train_dataset = train_dataset.shuffle(buffer_size=len(train)).batch(batch_size)
test_dataset = test_dataset.batch(batch_size=batch_size)
```

```
[22]: a = enumerate(train_dataset)
print(len(list(a)))
```

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#### 1.3 Define the Model

Now we will define the model. Here, we use the Keras Functional API to create a simple network of two Dense layers. We have modelled the problem as a binary classification problem and hence we add a single layer with sigmoid activation as the final layer of the model.

```
[23]: def base_model():
    inputs = tf.keras.layers.Input(shape=(len(train.columns)))

    x = tf.keras.layers.Dense(128, activation='relu')(inputs)
    x = tf.keras.layers.Dense(64, activation='relu')(x)
    outputs = tf.keras.layers.Dense(1, activation='sigmoid')(x)
    model = tf.keras.Model(inputs=inputs, outputs=outputs)
    return model

model = base_model()
```

## 1.4 Define Optimizer and Loss

We use RMSprop optimizer and binary crossentropy as our loss function.

```
[24]: optimizer = tf.keras.optimizers.RMSprop(learning_rate=0.001)
loss_object = tf.keras.losses.BinaryCrossentropy()
```

#### 1.5 Evaluate Untrained Model

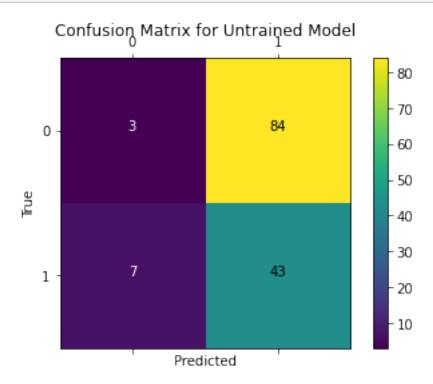
We calculate the loss on the model before training begins.

```
[25]: outputs = model(norm_test_X.values)
  loss_value = loss_object(y_true=test_Y.values, y_pred=outputs)
  print("Loss before training %.4f" % loss_value.numpy())
```

Loss before training 0.7284

We also plot the confusion matrix to visualize the true outputs against the outputs predicted by the model.

```
[27]: def plot_confusion_matrix(y_true, y_pred, title='', labels=[0,1]):
    cm = confusion_matrix(y_true, y_pred)
    fig = plt.figure()
    ax = fig.add_subplot(111)
    cax = ax.matshow(cm)
    plt.title(title)
    fig.colorbar(cax)
    ax.set_xticklabels([''] + labels)
    ax.set_yticklabels([''] + labels)
    plt.xlabel('Predicted')
```



# 1.6 Define Metrics (Please complete this section)

#### 1.6.1 Define Custom F1Score Metric

In this example, we will define a custom F1Score metric using the formula.

We use confusion\_matrix defined in tf.math to calculate precision and recall.

Here you can see that we have subclassed tf.keras.Metric and implemented the three required methods update\_state, result and reset\_states.

## 1.6.2 Please complete the result() method:

```
[29]: class F1Score(tf.keras.metrics.Metric):
          def __init__(self, name='f1_score', **kwargs):
              '''initializes attributes of the class'''
              # call the parent class init
              super(F1Score, self).__init__(name=name, **kwargs)
              # Initialize Required variables
              # true positives
              self.tp = tf.Variable(0, dtype = 'int32')
              # false positives
              self.fp = tf.Variable(0, dtype = 'int32')
              # true negatives
              self.tn = tf.Variable(0, dtype = 'int32')
              # false negatives
              self.fn = tf.Variable(0, dtype = 'int32')
          def update_state(self, y_true, y_pred, sample_weight=None):
              Accumulates statistics for the metric
              Arqs:
                  y_true: target values from the test data
                  y\_pred: predicted values by the model
              # Calulcate confusion matrix.
              conf_matrix = tf.math.confusion_matrix(y_true, y_pred, num_classes=2)
              # Update values of true positives, true negatives, false positives and_{f \sqcup}
       → false negatives from confusion matrix.
              self.tn.assign_add(conf_matrix[0][0])
              self.tp.assign_add(conf_matrix[1][1])
              self.fp.assign_add(conf_matrix[0][1])
              self.fn.assign_add(conf_matrix[1][0])
          def result(self):
              '''Computes and returns the metric value tensor.'''
              # Calculate precision
```

```
if (self.tp + self.fp == 0):
        precision = 1.0
    else:
        precision = self.tp / (self.tp + self.fp)
    # Calculate recall
    if (self.tp + self.fn == 0):
        recall = 1.0
    else:
        recall = self.tp / (self.tp + self.fn)
    # Return F1 Score
    ### START CODE HERE ###
    f1_score = 2 * ((precision * recall) / (precision + recall))
    ### END CODE HERE ###
    return f1_score
def reset_states(self):
    '''Resets all of the metric state variables.'''
    # The state of the metric will be reset at the start of each epoch.
    self.tp.assign(0)
    self.tn.assign(0)
    self.fp.assign(0)
    self.fn.assign(0)
```

```
[30]: # Test Code:

test_F1Score = F1Score()

test_F1Score.tp = tf.Variable(2, dtype = 'int32')
test_F1Score.fp = tf.Variable(5, dtype = 'int32')
test_F1Score.tn = tf.Variable(7, dtype = 'int32')
test_F1Score.fn = tf.Variable(9, dtype = 'int32')
test_F1Score.result()
```

[30]: <tf.Tensor: shape=(), dtype=float64, numpy=0.22222222222222222

## **Expected Output:**

```
<tf.Tensor: shape=(), dtype=float64, numpy=0.222222222222222>
```

We initialize the seprate metrics required for training and validation. In addition to our custom F1Score metric, we are also using BinaryAccuracy defined in tf.keras.metrics

```
[31]: train_f1score_metric = F1Score()
val_f1score_metric = F1Score()
```

```
train_acc_metric = tf.keras.metrics.BinaryAccuracy()
val_acc_metric = tf.keras.metrics.BinaryAccuracy()
```

# 1.7 Apply Gradients (Please complete this section)

The core of training is using the model to calculate the logits on specific set of inputs and compute the loss(in this case **binary crossentropy**) by comparing the predicted outputs to the true outputs. We then update the trainable weights using the optimizer algorithm chosen. The optimizer algorithm requires our computed loss and partial derivatives of loss with respect to each of the trainable weights to make updates to the same.

We use gradient tape to calculate the gradients and then update the model trainable weights using the optimizer.

## 1.7.1 Please complete the following function:

```
[32]: def apply_gradient(optimizer, loss_object, model, x, y):
          applies the gradients to the trainable model weights
          Arqs:
              optimizer: optimizer to update model weights
              loss_object: type of loss to measure during training
              model: the model we are training
              x: input data to the model
              y: target values for each input
          ,,,
          with tf.GradientTape() as tape:
          ### START CODE HERE ###
              logits = model(x)
              loss_value = loss_object(y_true=y, y_pred=logits)
          gradients = tape.gradient(loss_value, model.trainable_weights)
          optimizer.apply_gradients(zip(gradients, model.trainable_weights))
          ### END CODE HERE ###
          return logits, loss_value
```

```
print(test_logits.numpy()[:8])
print(test_loss.numpy())

del test_model
del test_logits
del test_loss
```

```
[[0.52413803]
[0.5602368]
[0.549544]
[0.54723155]
[0.48679572]
[0.5337864]
[0.4776442]
[0.5239992]]
```

## **Expected Output:**

The output will be close to these values:

```
[[0.5516499]
[0.52124363]
[0.5412698]
[0.54203206]
[0.50022954]
[0.5459626]
[0.47841492]
[0.54381996]]
0.7030578
```

# 1.8 Training Loop (Please complete this section)

This function performs training during one epoch. We run through all batches of training data in each epoch to make updates to trainable weights using our previous function. You can see that we also call update\_state on our metrics to accumulate the value of our metrics.

We are displaying a progress bar to indicate completion of training in each epoch. Here we use tqdm for displaying the progress bar.

## 1.8.1 Please complete the following function:

```
[34]: def train_data_for_one_epoch(train_dataset, optimizer, loss_object, model, train_acc_metric, train_f1score_metric, ...

→verbose=True):
```

```
Computes the loss then updates the weights and metrics for one epoch.
   Arqs:
       train_dataset: the training dataset
       optimizer: optimizer to update model weights
       loss_object: type of loss to measure during training
       model: the model we are training
       train_acc_metric: calculates how often predictions match labels
       train_f1score_metric: custom metric we defined earlier
   losses = []
   #Iterate through all batches of training data
   for step, (x_batch_train, y_batch_train) in enumerate(train_dataset):
       #Calculate loss and update trainable variables using optimizer
       ### START CODE HERE ###
       logits, loss_value = apply_gradient(optimizer, loss_object, model,_u
→x_batch_train, y_batch_train)
       losses.append(loss_value)
       ### END CODE HERE ###
       #Round off logits to nearest integer and cast to integer for calulating
\rightarrowmetrics
       logits = tf.round(logits)
       logits = tf.cast(logits, 'int64')
       #Update the training metrics
       ### START CODE HERE ###
       train_acc_metric.update_state(y_true=y_batch_train, y_pred=logits)
       train_f1score_metric.update_state(y_true=y_batch_train, y_pred=logits)
       ### END CODE HERE ###
       #Update progress
       if verbose:
           print("Training loss for step %s: %.4f" % (int(step),
→float(loss_value)))
   return losses
```

```
[36]: # TEST CODE

test_model = tf.keras.models.load_model('./test_model')

test_losses = train_data_for_one_epoch(train_dataset, optimizer, loss_object, u

→test_model,
```

```
train_acc_metric, train_f1score_metric,
verbose=False)

for test_loss in test_losses:
    print(test_loss.numpy())

del test_model
del test_losses
```

- 0.7552004
- 0.6243548
- 0.5939014
- 0.49887848
- 0.47701666
- 0.40541014
- 0.36220235
- 0.40997204
- 0.3567308
- 0.29308903
- 0.30118644
- 0.27259445
- 0.35707676
- 0.2417258
- 0.20194711
- 0.2440404
- 0.17027119
- 0.19259492

## **Expected Output:**

The losses should generally be decreasing and will start from around 0.75. For example:

- 0.7600615
- 0.6092045
- 0.5525634
- 0.4358902
- 0.4765755
- 0.43327087
- 0.40585428
- 0.32855004
- 0.35755336
- 0.3651728
- 0.33971977
- 0.27372319
- 0.25026917
- 0.29229593
- 0.242178
- 0.20602849

- 0.15887335
- 0.090397514

At the end of each epoch, we have to validate the model on the test dataset. The following function calculates the loss on test dataset and updates the states of the validation metrics.

```
[37]: def perform_validation():
    losses = []

#Iterate through all batches of validation data.
for x_val, y_val in test_dataset:

#Calculate validation loss for current batch.
    val_logits = model(x_val)
    val_loss = loss_object(y_true=y_val, y_pred=val_logits)
    losses.append(val_loss)

#Round off and cast outputs to either or 1
    val_logits = tf.cast(tf.round(model(x_val)), 'int64')

#Update validation metrics
    val_acc_metric.update_state(y_val, val_logits)
    val_f1score_metric.update_state(y_val, val_logits)

return losses
```

Next we define the training loop that runs through the training samples repeatedly over a fixed number of epochs. Here we combine the functions we built earlier to establish the following flow: 1. Perform training over all batches of training data. 2. Get values of metrics. 3. Perform validation to calculate loss and update validation metrics on test data. 4. Reset the metrics at the end of epoch. 5. Display statistics at the end of each epoch.

**Note**: We also calculate the training and validation losses for the whole epoch at the end of the epoch.

```
#Perform validation on all batches of test data
  losses_val = perform_validation()
   # Get results from validation metrics
  val_acc = val_acc_metric.result()
  val_f1score = val_f1score_metric.result()
  #Calculate training and validation losses for current epoch
  losses train mean = np.mean(losses train)
  losses_val_mean = np.mean(losses_val)
   epochs val losses.append(losses val mean)
   epochs_train_losses.append(losses_train_mean)
  print('\n Epcoh %s: Train loss: %.4f Validation Loss: %.4f, Train Accuracy:
→ %.4f, Validation Accuracy %.4f, Train F1 Score: %.4f, Validation F1 Score: ⊔
→%.4f' % (epoch, float(losses_train_mean), float(losses_val_mean),
→float(train_acc), float(val_acc), train_f1score, val_f1score))
   #Reset states of all metrics
  train_acc_metric.reset_states()
  val acc metric.reset states()
  val f1score metric.reset states()
  train_f1score_metric.reset_states()
```

```
Start of epoch 0
Training loss for step 0: 0.7127
Training loss for step 1: 0.5764
Training loss for step 2: 0.4708
Training loss for step 3: 0.4578
Training loss for step 4: 0.4218
Training loss for step 5: 0.3714
Training loss for step 6: 0.3532
Training loss for step 7: 0.3666
Training loss for step 8: 0.2898
Training loss for step 9: 0.3062
Training loss for step 10: 0.2359
Training loss for step 11: 0.1876
Training loss for step 12: 0.2370
Training loss for step 13: 0.1951
Training loss for step 14: 0.2294
Training loss for step 15: 0.1303
Training loss for step 16: 0.1295
Training loss for step 17: 0.1334
```

Epcoh 0: Train loss: 0.3225 Validation Loss: 0.1413, Train Accuracy: 0.9074, Validation Accuracy 0.9750, Train F1 Score: 0.8816, Validation F1 Score: 0.9600 Start of epoch 1

```
Training loss for step 0: 0.1034
Training loss for step 1: 0.2242
Training loss for step 2: 0.1379
Training loss for step 3: 0.1278
Training loss for step 4: 0.1505
Training loss for step 5: 0.0914
Training loss for step 6: 0.1202
Training loss for step 7: 0.1210
Training loss for step 8: 0.1361
Training loss for step 9: 0.0579
Training loss for step 10: 0.0493
Training loss for step 11: 0.0478
Training loss for step 12: 0.2459
Training loss for step 13: 0.1078
Training loss for step 14: 0.1651
Training loss for step 15: 0.0796
Training loss for step 16: 0.0617
Training loss for step 17: 0.0187
Epcoh 1: Train loss: 0.1137 Validation Loss: 0.0847, Train Accuracy: 0.9722,
Validation Accuracy 0.9750, Train F1 Score: 0.9583, Validation F1 Score: 0.9600
Start of epoch 2
Training loss for step 0: 0.1586
Training loss for step 1: 0.1552
Training loss for step 2: 0.0875
Training loss for step 3: 0.0727
Training loss for step 4: 0.1333
Training loss for step 5: 0.1190
Training loss for step 6: 0.0243
Training loss for step 7: 0.0489
Training loss for step 8: 0.1300
Training loss for step 9: 0.0766
Training loss for step 10: 0.0375
Training loss for step 11: 0.0951
Training loss for step 12: 0.0406
Training loss for step 13: 0.0317
Training loss for step 14: 0.0338
Training loss for step 15: 0.0336
Training loss for step 16: 0.1517
Training loss for step 17: 0.0118
Epcoh 2: Train loss: 0.0801 Validation Loss: 0.0731, Train Accuracy: 0.9722,
Validation Accuracy 0.9750, Train F1 Score: 0.9581, Validation F1 Score: 0.9600
Start of epoch 3
Training loss for step 0: 0.0176
Training loss for step 1: 0.0200
Training loss for step 2: 0.2108
Training loss for step 3: 0.0958
```

```
Training loss for step 5: 0.0339
Training loss for step 6: 0.0630
Training loss for step 7: 0.1484
Training loss for step 8: 0.1139
Training loss for step 9: 0.0108
Training loss for step 10: 0.0241
Training loss for step 11: 0.0618
Training loss for step 12: 0.1229
Training loss for step 13: 0.0706
Training loss for step 14: 0.0096
Training loss for step 15: 0.1235
Training loss for step 16: 0.0393
Training loss for step 17: 0.0176
Epcoh 3: Train loss: 0.0709 Validation Loss: 0.0706, Train Accuracy: 0.9722,
Validation Accuracy 0.9812, Train F1 Score: 0.9581, Validation F1 Score: 0.9703
Start of epoch 4
Training loss for step 0: 0.0235
Training loss for step 1: 0.0391
Training loss for step 2: 0.0369
Training loss for step 3: 0.0217
Training loss for step 4: 0.0193
Training loss for step 5: 0.2683
Training loss for step 6: 0.0583
Training loss for step 7: 0.0919
Training loss for step 8: 0.0791
Training loss for step 9: 0.0810
Training loss for step 10: 0.0166
Training loss for step 11: 0.1420
Training loss for step 12: 0.0626
Training loss for step 13: 0.0212
Training loss for step 14: 0.0748
Training loss for step 15: 0.0520
Training loss for step 16: 0.0876
Training loss for step 17: 0.0023
```

Epcoh 4: Train loss: 0.0655 Validation Loss: 0.0708, Train Accuracy: 0.9757, Validation Accuracy 0.9812, Train F1 Score: 0.9634, Validation F1 Score: 0.9703

#### 1.9 Evaluate the Model

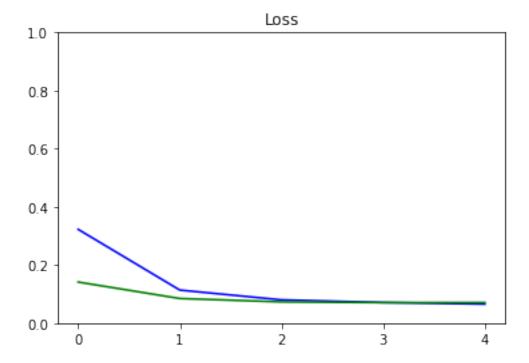
Training loss for step 4: 0.0931

#### 1.9.1 Plots for Evaluation

We plot the progress of loss as training proceeds over number of epochs.

```
[39]: def plot_metrics(train_metric, val_metric, metric_name, title, ylim=5):
    plt.title(title)
    plt.ylim(0,ylim)
    plt.gca().xaxis.set_major_locator(mticker.MultipleLocator(1))
    plt.plot(train_metric,color='blue',label=metric_name)
    plt.plot(val_metric,color='green',label='val_' + metric_name)

plot_metrics(epochs_train_losses, epochs_val_losses, "Loss", "Loss", ylim=1.0)
```



We plot the confusion matrix to visualize the true values against the values predicted by the model.

