### Machine Learning Homework 5

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```
[1]: # toc
     import os
     import matplotlib.pyplot as plt
     import certifi
     import numpy as np
     import seaborn as sns
     import cv2
     from sklearn.metrics import confusion_matrix, accuracy_score
     # noinspection PyUnresolvedReferences
     from tensorflow.keras.datasets import cifar10
     # noinspection PyUnresolvedReferences
     from tensorflow.keras.models import Sequential, load model
     # noinspection PyUnresolvedReferences
     from tensorflow.keras.layers import Conv2D, MaxPooling2D, Flatten, Dense,
      →Input, GlobalAveragePooling2D
     # noinspection PyUnresolvedReferences
     from tensorflow.keras.callbacks import ModelCheckpoint, EarlyStopping
     # noinspection PyUnresolvedReferences
     from tensorflow.keras.applications import MobileNetV2
     # noinspection PyUnresolvedReferences
     from tensorflow.keras.applications.mobilenet_v2 import preprocess_input
     # If you have a raw python installation, you have to set the SSL_CERT_FILE_
     ⇔environment variable
     os.environ['SSL_CERT_FILE'] = certifi.where()
     plt.style.use('../maroon_ipynb.mplstyle')
     # ignore all warnings
     import tensorflow as tf
     # noinspection PyUnresolvedReferences
     tf.get_logger().setLevel('ERROR')
```

## Contents

Problem 1 Solution		 		 				 								 				 		<b>3</b>
Problem 2 Solution		 		 	•			 	•					 •	•	 			 •	 		<b>6</b>
Problem 3 Solution		 		 		 •		 	•		 •		•			 			 •	 		<b>10</b>
Problem 4 Solution		 	•	 			•	 			 •	•				 	•			 		<b>12</b> 12
Problem 5 Solution		 		 				 								 	•	 •		 		<b>14</b> 14
Problem 6 Solution		 		 				 								 				 		18 18

Keras includes the CIFAR-10 dataset, a collection of 60,000 32x32 color images. Each contains 1 of 10 different objects, and the dataset is often used as a benchmark for classification schemes. The 10 different classes are: airplane, car, bird, cat, deer, dog, frog, horse, ship, and truck. In this assignment you will build a CNN to attempt to correctly classify these images.

Check the labels on the data to see how many of each class are present in the training and test datasets. Do you foresee any issues with this data balance? Plot a few of the images as well to get a sense for what the images look like to the human eye.

#### Solution

```
[2]: # load the dataset
  (x_train, y_train), (x_test, y_test) = cifar10.load_data()

# normalize the data
  x_train = x_train/255
  x_test = x_test/255
```

Upon inspection of the data, we see that x\_train is a 50,000x32x32x3 array meaning we have 50,000 images of size 32x32 with 3 color channels. y\_train is a 50,000x1 array of labels. x\_test is a 10,000x32x32x3 array and y\_test is a 10,000x1 array of labels. The target values are represented with integers from 0 to 9, corresponding to the class names below as the class\_names list.

Let's show the first of each class that we see in the training data.

```
[3]: # Plotting the images
     # class names order is seen in the doc strings of cifar10
     class_names = ['Airplane', 'Car', 'Bird', 'Cat', 'Deer', 'Dog', 'Frog', _
      ⇔'Horse', 'Ship', 'Truck']
     classes_seen = set()
     fig, ax = plt.subplots(nrows=5, ncols=2, figsize=(10, 20))
     axes = ax.flatten()
     i = 0
     while len(classes_seen) < len(class_names):</pre>
         label = y_train[i][0]
         if label not in classes seen:
             classes_seen.add(label)
             axes[label].imshow(x train[i])
             axes[label].set_title(class_names[label])
         i += 1
     plt.show()
```



#### Answer

```
Airplane: 5000 training images, 1000 test images
Car: 5000 training images, 1000 test images
Bird: 5000 training images, 1000 test images
Cat: 5000 training images, 1000 test images
Deer: 5000 training images, 1000 test images
Dog: 5000 training images, 1000 test images
Frog: 5000 training images, 1000 test images
Horse: 5000 training images, 1000 test images
Ship: 5000 training images, 1000 test images
Truck: 5000 training images, 1000 test images
```

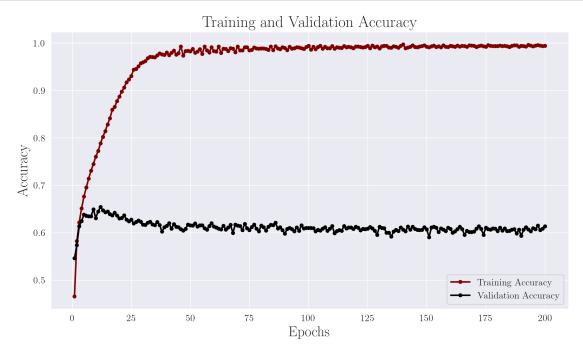
The CIFAR-10 dataset is designed so that each of the 10 classes has the same number of images—specifically, 5000 training images and 1000 test images per class. This balance ensures that models trained on the dataset don't become biased towards any one class and helps provide a fair evaluation during testing. So no, I don't foresee any issues with this data balance.

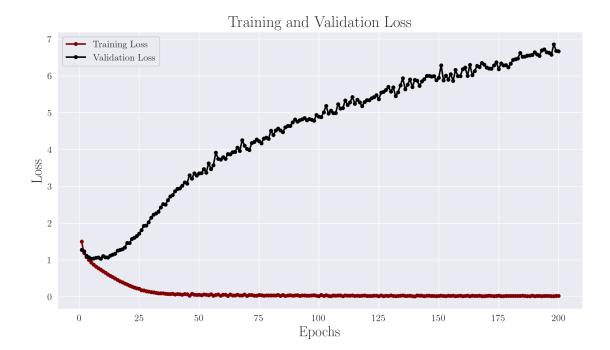
Build a simple CNN with a single convolution layer and a single dense hidden layer and train it over the data. Plot the evolution of the accuracy and the loss as a function of epoch. Show a confusion matrix for the best result from the training.

#### Solution

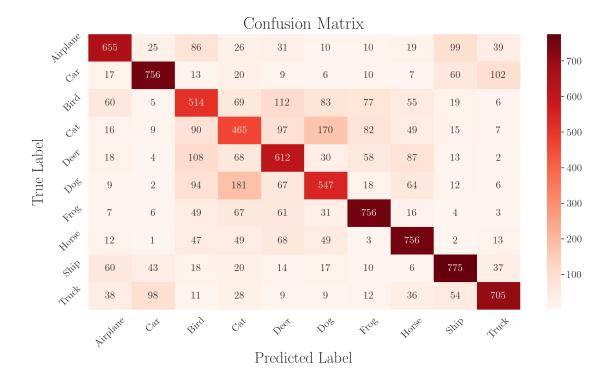
```
[5]: # Create model
    p2_model = Sequential([
         Input(shape=(32, 32, 3)), # input shape
         Conv2D(filters=32, kernel_size=(3, 3), activation='relu'), # convolution_
      \hookrightarrow layer
         MaxPooling2D(pool_size=(2, 2)), # max pooling layer
         Flatten(), # flatten the output
         Dense(128, activation='relu'), # dense hidden layer
         Dense(10, activation='softmax') # output layer with softmax activation
    ])
     # Compile model
     p2_model.compile(
         optimizer='adam', # optimizer
         loss='sparse_categorical_crossentropy', # loss function for multi-class⊔
     \hookrightarrow classification
        metrics=['accuracy'] # metric to track
     # Define callback to save the best model based on validation accuracy
     checkpoint_callback = ModelCheckpoint(
         'best_model_p2.keras', # file to save the model
         monitor='val_acc', # monitor validation accuracy
         save_best_only=True, # save only the best model
         mode='max', # maximize the monitored metric
         verbose=False # suppress output
     # Train the model
     history = p2 model.fit(
         x_train, y_train, # training data
         epochs=200, # number of epochs
         batch size=64, # batch size
         validation_data=(x_test, y_test), # validation data
         callbacks=[checkpoint_callback], # callback to save the best model
         verbose=False # suppress output during training
     )
     train_accuracy = history.history['acc']
     val_accuracy = history.history['val_acc']
```

```
train_loss = history.history['loss']
val_loss = history.history['val_loss']
epochs = np.array(history.epoch) + 1
# Plot the training and validation accuracy
fig, ax = plt.subplots()
ax.plot(epochs, train_accuracy, label='Training Accuracy', marker='.')
ax.plot(epochs, val_accuracy, label='Validation Accuracy', marker='.')
ax.set_title('Training and Validation Accuracy')
ax.set_xlabel('Epochs')
ax.set_ylabel('Accuracy')
ax.legend()
plt.show()
# Plot the training and validation loss
fig, ax = plt.subplots()
ax.plot(epochs, train_loss, label='Training Loss', marker='.')
ax.plot(epochs, val_loss, label='Validation Loss', marker='.')
ax.set_title('Training and Validation Loss')
ax.set_xlabel('Epochs')
ax.set_ylabel('Loss')
ax.legend()
plt.show()
```





```
[6]: # Making the confusion matrix
     p2_best_model = load_model('best_model_p2.keras') # Load the best model
     y_pred = p2_best_model.predict(x_test, verbose=False) # Get predictions on the_
      ⇔test set
     # Convert predictions to class labels
     y_pred_labels = np.argmax(y_pred, axis=1) # finding the highest probability
     cm = confusion_matrix(y_test.flatten(), y_pred_labels)
     # Plot confusion matrix
     fig, ax = plt.subplots()
     sns.heatmap(cm, annot=True, fmt='d', cmap='Reds', xticklabels=class_names,_
      →yticklabels=class_names, ax=ax)
     ax.set_title('Confusion Matrix')
     ax.set_xlabel('Predicted Label')
     ax.set_ylabel('True Label')
     ax.tick_params(axis='x', rotation=45)
     ax.tick_params(axis='y', rotation=45)
     plt.show()
```



[7]: # Accuracy score
accuracy = accuracy\_score(y\_test.flatten(), y\_pred\_labels)
print(f'Accuracy of the best model on the test set: {accuracy:.3%}')

#### Answer

Accuracy of the best model on the test set: 65.410%

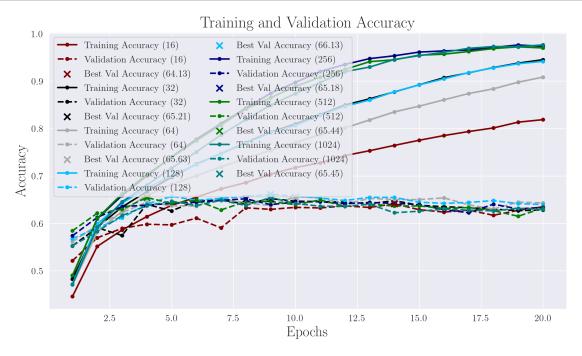
These results show that overfitting typically occurs around epoch 10 for this simple model. The training accuracy continues to improve while the validation accuracy starts to decline. This is even more evident in the loss plots, where the training loss keeps decreasing while the validation loss increases after epoch 10. This is a result of the simple architecture quickly learning the training data but failing to generalize to unseen data.

Measure how the accuracy varies as you change the size of the single convolution layer. Does there appear to be an ideal size for training?

#### Solution

I am going to assume that this means we are changing the filter size of the convolution layer.

```
[8]: fig, ax = plt.subplots()
     filter_sizes = [16, 32, 64, 128, 256, 512, 1024]
     for filter_size in filter_sizes:
         # Create model
         p3 model = Sequential([
             Input(shape=(32, 32, 3)), # input shape
             Conv2D(filters=filter_size, kernel_size=(3, 3), activation='relu'), #_U
      ⇔convolution layer
             MaxPooling2D(pool_size=(2, 2)), # max pooling layer
             Flatten(), # flatten the output
             Dense(128, activation='relu'), # dense hidden layer
             Dense(10, activation='softmax') # output layer with softmax activation
         ])
         # Compile model
         p3_model.compile(
             optimizer='adam', # optimizer
             loss='sparse_categorical_crossentropy', # loss function for_
      \hookrightarrow multi-class classification
             metrics=['acc'] # metric to track
         )
         # Train the model
         history = p3_model.fit(
             x_train, y_train, # training data
             epochs=20, # number of epochs
             batch_size=64, # batch size
             validation_data=(x_test, y_test), # validation data
             verbose=False # suppress output during training
         )
         train_accuracy = history.history['acc']
         val_accuracy = history.history['val_acc']
         epochs = np.array(history.epoch) + 1
         best_accuracy = max(val_accuracy) # find the best validation accuracy
         best_accuracy_epoch = val_accuracy.index(best_accuracy) + 1 # find the__
      \rightarrowepoch at which it occurred
         # Plot the training and validation accuracy
```

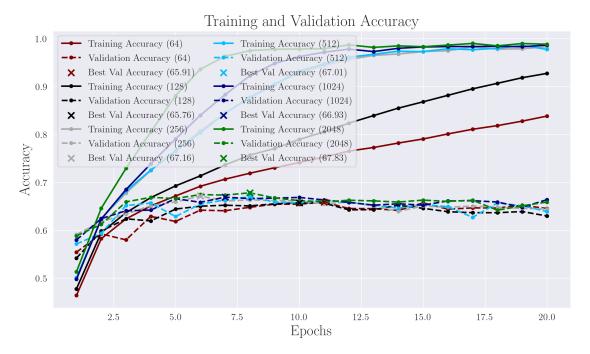


The results show that as the output size of the convolution layer increases, the validation accuracy has little change. The training accuracy, however, will converge quicker as the filter size increases, which means that the validation accuracy converges in fewer epochs, but each epoch takes longer to train with the higher number of filters. Nevertheless, a filter size of 32 or 64 seems to be alright, but I don't see major differences that would warrant an ideal size.

Now vary the size of the dense layer and do the same analysis as above.

#### Solution

```
[9]: fig, ax = plt.subplots()
    dense_sizes = [64, 128, 256, 512, 1024, 2048]
    for dense_size in dense_sizes:
        # Create model
        p4_model = Sequential([
            Input(shape=(32, 32, 3)), # input shape
            Conv2D(filters=64, kernel_size=(3, 3), activation='relu'), #__
      ⇔convolution layer
            MaxPooling2D(pool_size=(2, 2)), # max pooling layer
            Flatten(), # flatten the output
            Dense(dense_size, activation='relu'), # dense hidden layer
            Dense(10, activation='softmax') # output layer with softmax activation
        ])
        # Compile model
        p4_model.compile(
            optimizer='adam', # optimizer
            loss='sparse_categorical_crossentropy', # loss function for_
      \hookrightarrow multi-class classification
            metrics=['acc'] # metric to track
        )
        # Train the model
        history = p4_model.fit(
            x_train, y_train, # training data
            epochs=20, # number of epochs
            batch_size=64, # batch size
            validation_data=(x_test, y_test), # validation data
            verbose=False # suppress output during training
        )
        train_accuracy = history.history['acc']
        val_accuracy = history.history['val_acc']
        epochs = np.array(history.epoch) + 1
        best_accuracy = max(val_accuracy) # find the best validation accuracy
        best_accuracy_epoch = val_accuracy.index(best_accuracy) + 1 # find the_
      →epoch at which it occurred
        # Plot the training and validation accuracy
        line = ax.plot(epochs, train_accuracy, label=f'Training Accuracy_
```



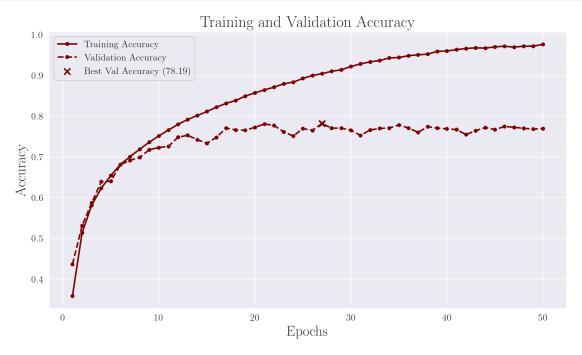
The results show that the validation accuracy increases by about 3% at 2048. There could be further increase in the accuracy, but this analysis takes a really long time. For refining in further problems, I will choose to start at 2048 and work from there.

Using multiple convolution and pooling layers, and a single dense layer, find a combination of hyperparameters that minimizes your validation error. What are the optimum parameters you found? Plot the confusion matrix for this best case.

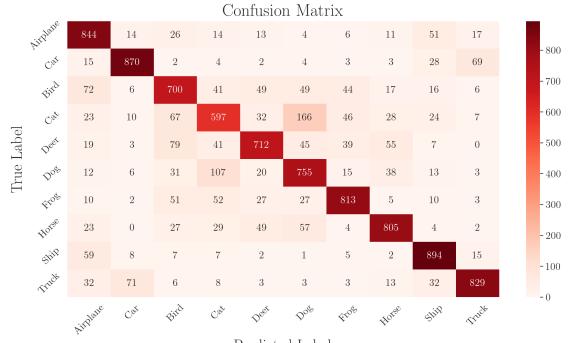
#### Solution

```
[17]: # Revised model with 'same' padding and fewer pooling layers to preserve
       \hookrightarrow dimensions
      p5_model = Sequential([
          Input(shape=(32, 32, 3)),
          Conv2D(filters=32, kernel_size=(3, 3), activation='relu', padding='same'),
          MaxPooling2D(pool_size=(2, 2)),
          Conv2D(filters=64, kernel_size=(3, 3), activation='relu', padding='same'),
          MaxPooling2D(pool_size=(2, 2)),
          Conv2D(filters=128, kernel_size=(3, 3), activation='relu', padding='same'),
          GlobalAveragePooling2D(),
          Dense(2048, activation='relu'),
          Dense(10, activation='softmax')
      ])
      # Compile model
      p5_model.compile(
          optimizer='adam',
          loss='sparse_categorical_crossentropy',
          metrics=['acc']
      )
      # Define callbacks: save the best model and use early stopping
      checkpoint_callback = ModelCheckpoint(
          'best_model_p5.keras',
          monitor='val_acc',
          save_best_only=True,
          mode='max',
          verbose=False
      # early_stopping = EarlyStopping(
      #
          monitor='val_loss',
      #
            patience=3,
      #
            restore_best_weights=True,
            verbose=False
      # )
      # Train the model
      history = p5_model.fit(
          x_train, y_train,
```

```
epochs=50,
    batch_size=64,
   validation_data=(x_test, y_test),
    # callbacks=[checkpoint_callback, early_stopping],
    callbacks=[checkpoint_callback],
   verbose=False
)
train_accuracy = history.history['acc']
val_accuracy = history.history['val_acc']
epochs = np.array(history.epoch) + 1
best_accuracy = max(val_accuracy) # find the best validation accuracy
best_accuracy_epoch = val_accuracy.index(best_accuracy) + 1 # find the epoch_
 →at which it occurred
# Plot the training and validation accuracy
fig, ax = plt.subplots()
line = ax.plot(epochs, train_accuracy, label=f'Training Accuracy', marker='.',u
 ⇒zorder=2) [0]
ax.plot(epochs, val_accuracy, label=f'Validation Accuracy', marker='.', u
 s='--', color=line.get_color(), zorder=2)
ax.scatter(best_accuracy_epoch, best_accuracy, marker='x', color=line.
 Get_color(), zorder=3, label=fr'Best Val Accuracy ({best_accuracy*100:.2f})')
ax.set title('Training and Validation Accuracy')
ax.set_xlabel('Epochs')
ax.set_ylabel('Accuracy')
ax.legend()
plt.show()
```



```
[18]: # Making the confusion matrix
     p5_best_model = load_model('best_model_p5.keras') # Load_the_best_model
     y_pred = p5_best_model.predict(x_test, verbose=False) # Get predictions on the_
       →test set
     # Convert predictions to class labels
     y_pred_labels = np.argmax(y_pred, axis=1) # finding the highest probability
     cm = confusion_matrix(y_test.flatten(), y_pred_labels)
     # Plot confusion matrix
     fig, ax = plt.subplots()
     sns.heatmap(cm, annot=True, fmt='d', cmap='Reds', xticklabels=class_names,_
       ax.set_title('Confusion Matrix')
     ax.set_xlabel('Predicted Label')
     ax.set_ylabel('True Label')
     ax.tick_params(axis='x', rotation=45)
     ax.tick_params(axis='y', rotation=45)
     plt.show()
```



# [19]: # Accuracy score accuracy = accuracy\_score(y\_test.flatten(), y\_pred\_labels) print(f'Accuracy of the best model on the test set: {accuracy:.3%}')

#### Answer

Accuracy of the best model on the test set: 78.190%

In this problem, we built a more complex CNN using multiple convolutional layers with 'same' padding, max pooling, and a global average pooling layer, followed by a large dense layer. This architecture allowed the network to extract richer features while preventing excessive reduction in spatial dimensions. By using callbacks like model checkpointing, the model minimized validation error. I tried adding more layers and adjusting the hyperparameters and found that 32, 64, and 128 produce the best results. The final accuracy of 78% exceeds the single convolution layer by 10%, which is pretty substantial.

One method that could be used to improve these results is to add onto the training data by using the existing images and duplicating with zoom, rotations, translations, and mirroring, but this was left out on purpose due to time constraints.

Now replace your convolution layers with one of the existing pretrained models available in keras and optimize your dense layer to produce the best possible fit. What is the validation accuracy in this case? How does it compare to the convolution layers you trained by hand?

#### Solution

In this approach we use Keras's pretrained MobileNetV2 as a feature extractor (with its convolutional base frozen) and then add our own dense layers on top. Note that because MobileNetV2 was originally trained on ImageNet, its preprocessing expects pixel values in a different range than our normalized CIFAR-10 images; we therefore first rescale the images back to [0, 255] and then use the MobileNetV2 preprocessing function.

```
[4]: # Preprocess CIFAR-10 images for MobileNetV2:
     # Our x train and x test are normalized between O and 1, but MobileNetV2,
     ⇔expects inputs in a different scale.
     # First, rescale back to [0, 255], resize to 224x224, then use the MobileNetV2
     ⇔preprocessing.
     x_train_rescaled = (x_train*255).astype('float32')
     x_test_rescaled = (x_test*255).astype('float32')
     def resize_images(images, size=(224, 224)):
        resized = np.zeros((images.shape[0], size[0], size[1], images.shape[3]),

dtype=np.float32)
        for im in range(images.shape[0]):
             resized[im] = cv2.resize(images[im], size)
        return resized
     x train resized = resize images(x train rescaled)
     x_test_resized = resize_images(x_test_rescaled)
     x_train_preprocessed = preprocess_input(x_train_resized)
     x_test_preprocessed = preprocess_input(x_test_resized)
```

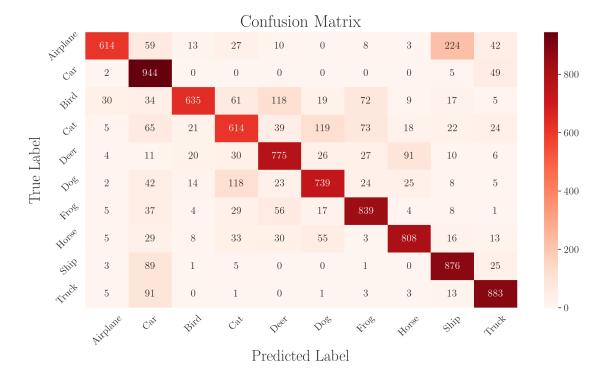
```
[]: # Load the pretrained MobileNetV2 model without its top, and freeze its layers.
base_model = MobileNetV2(include_top=False, weights='imagenet',u
input_shape=(224, 224, 3))
base_model.trainable = False

# Build our new model by stacking our dense layers on top of the frozen base.
p6_model = Sequential([
    base_model,
    GlobalAveragePooling2D(),
    Dense(512, activation='relu'), # optimized dense layer
    Dense(10, activation='softmax')
])
```

```
# Compile the model.
p6_model.compile(
    optimizer='adam',
    loss='sparse_categorical_crossentropy',
    metrics=['acc']
)
# Create callback
checkpoint callback = ModelCheckpoint(
    'best_model_p6.keras',
    monitor='val acc',
    save_best_only=True,
    mode='max',
    verbose=True
)
# Train the model.
history_p6 = p6_model.fit(
    x_train_preprocessed, y_train,
    epochs=10,
    batch_size=64,
    validation_data=(x_test_preprocessed, y_test),
    callbacks=[checkpoint_callback],
    verbose=True
)
# Get accuracy results
train_accuracy = history_p6.history['acc']
val_accuracy = history_p6.history['val_acc']
epochs = np.array(history_p6.epoch) + 1
best_accuracy = max(val_accuracy)
best_accuracy_epoch = val_accuracy.index(best_accuracy) + 1
# Plot the training and validation accuracy
fig, ax = plt.subplots()
line = ax.plot(epochs, train_accuracy, label=f'Training Accuracy', marker='.', u
 ⇒zorder=2)[0]
ax.plot(epochs, val_accuracy, label=f'Validation Accuracy', marker='.', __
⇒ls='--', color=line.get_color(), zorder=2)
ax.scatter(best_accuracy_epoch, best_accuracy, marker='x', color=line.

→get_color(), zorder=3, label=fr'Best Val Accuracy ({best_accuracy*100:.2f})')
ax.set title('Training and Validation Accuracy')
ax.set_xlabel('Epochs')
ax.set_ylabel('Accuracy')
ax.legend()
plt.show()
```

```
[8]: # Making the confusion matrix
     p6_best_model = load_model('best_model_p6.keras')  # Load the best model
     y_pred = p6_best_model.predict(x_test_preprocessed, verbose=False) # Get∟
      ⇔predictions on the test set
     # Convert predictions to class labels
     y_pred_labels = np.argmax(y_pred, axis=1) # finding the highest probability
     cm = confusion_matrix(y_test.flatten(), y_pred_labels)
     # Plot confusion matrix
     fig, ax = plt.subplots()
     sns.heatmap(cm, annot=True, fmt='d', cmap='Reds', xticklabels=class_names,_
      ⇔yticklabels=class_names, ax=ax)
     ax.set title('Confusion Matrix')
     ax.set xlabel('Predicted Label')
     ax.set_ylabel('True Label')
     ax.tick_params(axis='x', rotation=45)
     ax.tick_params(axis='y', rotation=45)
     plt.show()
```



[9]: # Accuracy score
accuracy = accuracy\_score(y\_test.flatten(), y\_pred\_labels)
print(f'Accuracy of the best model on the test set: {accuracy:.3%}')

#### Answer

Accuracy of the best model on the test set: 77.270%