MANE 4962 HW5

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Problem 1

```
In [ ]: import numpy as np
          from tensorflow.keras.models import Sequential
          from tensorflow.keras.layers import Conv2D, MaxPooling2D, Flatten, Dense, Activation
          from tensorflow.keras.datasets import cifar10
          from tensorflow.keras.utils import to categorical
           (x train, y train), (x test, y test) = cifar10.load data()
          x train, x test = x train / 255.0, x test / 255.0
          y train, y test = to categorical(y train, 10), to categorical(y test, 10)
          model = Sequential([
              Conv2D(64, (5, 5), input shape=(32, 32, 3), activation='relu'),
              MaxPooling2D(pool size=(2, 2)),
              Conv2D(32, (3, 3), activation='relu'),
              MaxPooling2D(pool size=(2, 2)),
              Conv2D(32, (3, 3), activation='relu'),
              Flatten(),
              Dense (64, activation='relu'),
              Dense(10, activation='softmax')
          ])
```

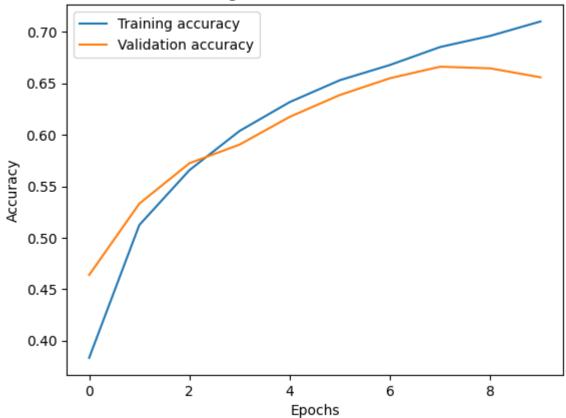
```
In [ ]: model.compile(optimizer='adam', loss='categorical crossentropy', metrics=['accuracy'])
   history = model.fit(x train, y train, epochs=10, batch size=64, validation data=(x test, y test))
   Epoch 1/10
   782/782 [============] - 91s 114ms/step - loss: 1.6773 - accuracy: 0.3832 - val loss: 1.4454 - val ac
   curacy: 0.4638
   Epoch 2/10
   curacy: 0.5331
   Epoch 3/10
   curacy: 0.5724
   Epoch 4/10
   curacy: 0.5905
   Epoch 5/10
   curacy: 0.6175
   Epoch 6/10
   curacy: 0.6387
   Epoch 7/10
   curacy: 0.6550
   Epoch 8/10
   curacy: 0.6662
   Epoch 9/10
   curacy: 0.6646
   Epoch 10/10
```

curacy: 0.6559

```
In [ ]: import matplotlib.pyplot as plt

plt.plot(history.history['accuracy'], label='Training accuracy')
plt.plot(history.history['val_accuracy'], label='Validation accuracy')
plt.title('Training and Validation Accuracies')
plt.xlabel('Epochs')
plt.ylabel('Accuracy')
plt.legend()
plt.show()
```

Training and Validation Accuracies



Problem 2

2(a)

```
In [ ]: from sklearn import datasets
          from sklearn.preprocessing import StandardScaler
          from sklearn. model selection import train test split
          from sklearn import svm
          from sklearn. metrics import accuracy score
          # Load dataset
          breast cancer = datasets.load breast cancer()
          X = breast cancer.data
          y = breast cancer. target
          # Select features - indices may vary based on documentation
          X = X[:, [23, 24, 28]] # Assuming these are 'worst area', 'worst compactness', 'worst concavity'
          # Split dataset into training and testing set
          X train, X test, y train, y test = train test split(X, y, test size=0.3, random state=42)
          # Standardize features
          scaler = StandardScaler()
          X train = scaler.fit transform(X train)
          X test = scaler.transform(X test)
          # Create SVM classifier with a linear kernel
          clf linear = svm. SVC(kernel='linear')
          clf linear.fit(X train, y train)
          # Predict and evaluate the model
          y pred linear = clf linear.predict(X test)
          accuracy linear = accuracy score(y test, y pred linear)
          print(f'Accuracy of SVM with linear kernel: {accuracy linear}')
```

Accuracy of SVM with linear kernel: 0.9707602339181286

Accuracy of SVM with RBF kernel and C=2: 0.9473684210526315