Github link: https://github.com/Kavyareddy03/ICP-5

```
from tensorflow.keras.models import Sequential
    from tensorflow.keras.layers import Conv2D, MaxPooling2D, Dropout, Flatten, Dense
    from keras.datasets import cifar10
    from keras.src.utils import to_categorical
    import matplotlib.pyplot as plt
    np.random.seed(seed)
    # load data
    (X_train, y_train), (X_test, y_test) = cifar10.load_data()
# normalize inputs from 0-255 to 0.0-1.0
    X_train = X_train.astype('float32')
    X_test = X_test.astype('float32')
    X_train = X_train / 255.0
    X_test = X_test / 255.0
    # one hot encode outputs
    # use to_categorical directly instead of np_utils.to_categorical
    y_train = to_categorical(y_train)
    y_test = to_categorical(y_test)
    num_classes = y_test.shape[1]
    # Initialize the model
    model = Sequential()
    # 1st Convolutional Layer (32 feature maps, 3x3 kernel, ReLU activation)
    model.add(Conv2D(32, (3, 3), activation='relu', input_shape=(32, 32, 3))) # Adjust input shape for your data
```

```
# Max Pooling Layer (2x2 pool size)
model.add(MaxPooling2D(pool_size=(2, 2)))
# 2nd Convolutional Layer (64 feature maps, 3x3 kernel, ReLU activation)
model.add(Conv2D(64, (3, 3), activation='relu'))
# Max Pooling Layer (2x2 pool size)
model.add(MaxPooling2D(pool_size=(2, 2)))
# Flatten layer
model.add(Flatten())
# Fully Connected Layer (512 units, ReLU activation)
model.add(Dense(512, activation='relu'))
# Dropout (50%)
model.add(Dropout(0.5))
# Output Layer (10 units, Softmax activation)
model.add(Dense(10, activation='softmax'))
model.compile(optimizer='adam', loss='categorical_crossentropy', metrics=['accuracy'])
# Summary of the model
model.summary()
```

Downloading data from https://www.cs.toronto.edu/~kriz/cifar-10-python.tar.gz

Zs dus/step
/usr/local/lib/python3.10/dist-packages/keras/src/layers/convolutional/base_conv.py:107: UserWarning: Do not pass an `input_shape`/`input_dim` argume super().__init__(activity_regularizer=activity_regularizer, **kwargs)
Model: "sequential"

Layer (type)	Output Shape	Param #
conv2d (Conv2D)	(None, 30, 30, 32)	896
max_pooling2d (MaxPooling2D)	(None, 15, 15, 32)	0
conv2d_1 (Conv2D)	(None, 13, 13, 64)	18,496
max_pooling2d_1 (MaxPooling2D)	(None, 6, 6, 64)	0
flatten (Flatten)	(None, 2304)	0
dense (Dense)	(None, 512)	1,180,160
dropout (Dropout)	(None, 512)	9
dense_1 (Dense)	(None, 10)	5,130

Total params: 1,204,682 (4.60 MB) Trainable params: 1,204,682 (4.60 MB) Non-trainable params: 0 (0.00 B)

Explanation:

- 1. Data Preparation: It loads the CIFAR-10 dataset, normalizes the pixel values from 0-255 to 0-1, and applies one-hot encoding to the labels.
- 2. Model Definition: A Sequential model is created with:
- Convolutional Layers: Two layers that apply convolution operations to extract features from the images, each followed by a ReLU activation function.
- Max Pooling Layers: Two layers that downsample the feature maps, reducing spatial dimensions while retaining important features.
 - Flatten Layer: Converts the 2D feature maps into a 1D vector.
- Dense Layer: A fully connected layer with 512 units and ReLU activation for learning complex patterns.
- Dropout Layer: Regularizes the model by randomly dropping 50% of the neurons during training to prevent overfitting.
- Output Layer: A dense layer with 10 units and softmax activation for multi-class classification.
- 3. Model Compilation: The model is compiled with the Adam optimizer, categorical crossentropy loss, and accuracy as the evaluation metric.
- 4. Model Summary: A summary of the model architecture is printed to show the layers and parameters.

```
import matplotlib.pyplot as plt
plt.figure(figsize = (4,4))
for i in range(4):
    plt.subplot(3,5,1+i)
    plt.axis('off')
    plt.imshow(X_train[i], cmap = 'gray')
```

Explanation:

- 1. Figure Setup: It creates a 4x4 inch figure.
- 2. Subplot Loop: It iterates over the first four images to create subplots in a 3-row by 5-column grid.
- 3. Axis Management: The axes are turned off for each subplot to focus solely on the images.
- 4. Image Display: Each image is displayed using a grayscale colormap, though the images are originally colored.

Explanation:

Prediction: It uses the model to predict the labels for the first four images in X_test.

Label Conversion: The predicted probabilities are converted to class labels by using argmax, which identifies the index of the highest probability for each prediction.

Actual Labels: The true labels of the first four images are also converted to class labels using argmax.

Output: Finally, it prints the predicted labels and the actual labels for comparison.

```
classes = ["airplane","automobile","bird","cat","deer","dog","frog","horse","ship","truck"]
L = 2
W = 2
fig, axes = plt.subplots(L, W, figsize = (6,6))
axes = axes.ravel() #

for i in np.arange(0, 4):
    axes[i].imshow(X_test[i])
    axes[i].set_title("Predicted = {}\n Actual = {}\".format(classes[y_predictions[i]], classes[actual_labels[i]]))
    axes[i].axis('off')

plt.subplots_adjust(wspace=1)
```

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Predicted = deer Actual = cat



Predicted = deer Actual = ship



 $\overline{\Rightarrow}$

Predicted = deer Actual = cat



Predicted = deer Actual = ship



Predicted = deer Actual = ship



Predicted = deer Actual = airplane

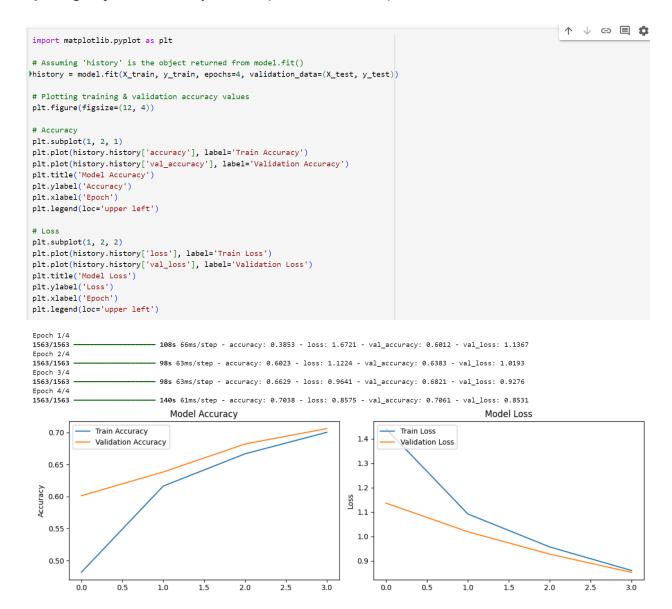


Explanation:

Class Labels: A list of class names for the CIFAR-10 dataset is defined. **Figure Setup**: A 2x2 grid of subplots is created for displaying images. **Image Loop**: It iterates over the first four images:

- Each image is displayed in its corresponding subplot.
- The title for each subplot shows the predicted and actual class labels.
- The axis is turned off for a cleaner look.

Spacing Adjustment: It adjusts the space between subplots for better visualization



Explanation:

Model Training: The model is trained using the training data for four epochs, with validation data also provided to monitor performance.

Figure Setup: A figure is created with a specified size to accommodate two subplots side by side.

Accuracy Plot:

- The first subplot shows the training and validation accuracy over the epochs.
- Lines are plotted for both metrics, and the axes are labeled accordingly.
- A legend is included to differentiate between training and validation accuracy.

Loss Plot:

- The second subplot displays the training and validation loss over the epochs.
- Similar to the accuracy plot, lines are added, and the axes are labeled.

Layout Adjustment: The layout is adjusted for better spacing, and the plots are displayed