

# GA\_HPO\_1\_Code

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[ ]: import numpy as np
import matplotlib.pyplot as plt
import random
import tensorflow as tf
from deap import base, creator, tools
from functools import partial
from tensorflow.keras.models import Sequential
from tensorflow.keras.layers import Conv2D, MaxPooling2D, Flatten, Dense,
    ↳Dropout, BatchNormalization
from tensorflow.keras.optimizers import Adam
from tensorflow.keras.callbacks import EarlyStopping
from sklearn.metrics import classification_report, confusion_matrix,
    ↳roc_auc_score
import time

# Load and preprocess the CIFAR-10 dataset
(x_train, y_train), (x_test, y_test) = tf.keras.datasets.cifar10.load_data()
x_train, x_test = x_train / 255.0, x_test / 255.0
y_train, y_test = tf.keras.utils.to_categorical(y_train), tf.keras.utils.
    ↳to_categorical(y_test)

# Split the training data into train and validation sets
validation_split = 0.1
split_index = int(len(x_train) * validation_split)
x_val, y_val = x_train[:split_index], y_train[:split_index]
x_train, y_train = x_train[split_index:], y_train[split_index:]

sample_size = 5000 # Adjust this value as needed
sample_indices = np.random.choice(np.arange(x_train.shape[0]), sample_size,
    ↳replace=False)

x_train_small = x_train[sample_indices]
y_train_small = y_train[sample_indices]

# Create fitness and individual classes for DEAP
creator.create("FitnessMin", base.Fitness, weights=(-1.0,))
creator.create("Individual", list, fitness=creator.FitnessMin)
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# Initialize the individual and population functions for DEAP
def init_individual():
    num_filters1 = random.randint(16, 128)
    num_filters2 = random.randint(16, 128)
    dense_units = random.randint(128, 1024)
    learning_rate = random.uniform(1e-5, 1e-2)
    return creator.Individual([num_filters1, num_filters2, dense_units,
↪learning_rate])

def init_population(n):
    return [init_individual() for _ in range(n)]

def fitness_function_data(individual):
    num_filters1, num_filters2, dense_units, learning_rate = individual
    num_filters1 = int(num_filters1)
    num_filters2 = int(num_filters2)
    dense_units = int(dense_units)

    print(f"Training with hyperparameters: num_filters1={num_filters1},
↪num_filters2={num_filters2}, "
          f"dense_units={dense_units}, learning_rate={learning_rate}")

    model = Sequential([
        Conv2D(num_filters1, (3, 3), activation='relu', padding='same',
↪input_shape=(32, 32, 3)),
        BatchNormalization(),
        Conv2D(num_filters1, (3, 3), activation='relu', padding='same'),
        BatchNormalization(),
        MaxPooling2D((2, 2)),
        Dropout(0.25),

        Conv2D(num_filters2, (3, 3), activation='relu', padding='same'),
        BatchNormalization(),
        Conv2D(num_filters2, (3, 3), activation='relu', padding='same'),
        BatchNormalization(),
        MaxPooling2D((2, 2)),
        Dropout(0.25),

        Flatten(),
        Dense(dense_units, activation='relu'),
        BatchNormalization(),
        Dropout(0.5),
        Dense(10, activation='softmax')
    ])

    model.compile(optimizer=Adam(learning_rate=learning_rate),

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        loss='categorical_crossentropy',
        metrics=['accuracy'])

    early_stopping = EarlyStopping(monitor='val_loss', patience=5,
↪ restore_best_weights=True)

    history = model.fit(x_train_small, y_train_small, epochs=5, batch_size=256,
                        validation_data=(x_val, y_val),
                        callbacks=[early_stopping],
                        verbose=0)

    val_loss = min(history.history['val_loss'])
    return val_loss,

# Define the evaluation, crossover, and mutation functions for DEAP
toolbox = base.Toolbox()
toolbox.register("individual", init_individual)
toolbox.register("population", init_population, n=10)
toolbox.register("mate", tools.cxTwoPoint)
toolbox.register("mutate", tools.mutGaussian, mu=0, sigma=0.2, indpb=0.1)
toolbox.register("select", tools.selBest)
toolbox.register("evaluate", fitness_function_data)

# Run the GA optimizer
population = toolbox.population()
NGEN = 20

for gen in range(NGEN):
    print(f"Generation {gen + 1} of {NGEN}")
    offspring = tools.selBest(population, len(population) // 2)
    offspring = list(offspring)
    for child1, child2 in zip(offspring[::2], offspring[1::2]):
        if random.random() < 0.5:
            toolbox.mate(child1, child2)
            del child1.fitness.values
            del child2.fitness.values

    for mutant in offspring:
        if random.random() < 0.2:
            toolbox.mutate(mutant)
            del mutant.fitness.values

    invalid_ind = [ind for ind in offspring if not ind.fitness.valid]
    fitnesses = map(toolbox.evaluate, invalid_ind)
    for ind, fit in zip(invalid_ind, fitnesses):
        ind.fitness.values = fit

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    # Sort the population based on their fitness values
    population = sorted(population, key=lambda ind: ind.fitness.values[0] if
↳ ind.fitness.valid else float('inf'))

    # Replace the least fit individuals with offspring
    population[:len(offspring)] = offspring

    # Print the best fitness value in the current generation
    best_fitness = min(ind.fitness.values[0] for ind in population if ind.
↳ fitness.valid)
    print(f"Best fitness in generation {gen + 1}: {best_fitness:.4f}")

best_individual = tools.selBest(population, 1)[0]

# Extract the best hyperparameters
best_num_filters1, best_num_filters2, best_dense_units, best_learning_rate =
↳ best_individual
best_num_filters1 = int(best_num_filters1)
best_num_filters2 = int(best_num_filters2)
best_dense_units = int(best_dense_units)

print(f"Best hyperparameters found by GA: num_filters1={best_num_filters1}, "
      f"num_filters2={best_num_filters2}, dense_units={best_dense_units}, "
      f"learning_rate={best_learning_rate}")

# Train the model with the best hyperparameters
model = Sequential([
    Conv2D(best_num_filters1, (3, 3), activation='relu', padding='same',
↳ input_shape=(32, 32, 3)),
    BatchNormalization(),
    Conv2D(best_num_filters1, (3, 3), activation='relu', padding='same'),
    BatchNormalization(),
    MaxPooling2D((2, 2)),
    Dropout(0.25),

    Conv2D(best_num_filters2, (3, 3), activation='relu', padding='same'),
    BatchNormalization(),
    Conv2D(best_num_filters2, (3, 3), activation='relu', padding='same'),
    BatchNormalization(),
    MaxPooling2D((2, 2)),
    Dropout(0.25),

    Flatten(),
    Dense(best_dense_units, activation='relu'),
    BatchNormalization(),
    Dropout(0.5),

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        Dense(10, activation='softmax')
    ])

model.compile(optimizer=Adam(learning_rate=best_learning_rate),
              loss='categorical_crossentropy',
              metrics=['accuracy'])

early_stopping = EarlyStopping(monitor='val_loss',
                               patience=5, restore_best_weights=True)

start_time = time.time()
history = model.fit(x_train, y_train, epochs=50, batch_size=64,
                   validation_data=(x_val, y_val),
                   callbacks=[early_stopping],
                   verbose=1)

end_time = time.time()

training_time = end_time - start_time
print(f'Total training time: {training_time:.2f} seconds')

test_loss, test_acc = model.evaluate(x_test, y_test, verbose=0)
y_pred = model.predict(x_test)
y_pred_classes = np.argmax(y_pred, axis=1)
y_test_classes = np.argmax(y_test, axis=1)

print(f'Test accuracy: {test_acc}')

print("Classification Report:")
print(classification_report(y_test_classes, y_pred_classes))

print("Confusion Matrix:")
print(confusion_matrix(y_test_classes, y_pred_classes))

# Calculate ROC-AUC for multi-class classification
roc_auc = roc_auc_score(y_test, y_pred, multi_class='ovr')
print(f'ROC-AUC Score: {roc_auc}')

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

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[ ]: # Using these hardcoded values as obtained from confusion matrix above inorder
      ↳to make a plot
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns

# Confusion matrix data
confusion_matrix = np.array([
    [805, 18, 31, 31, 7, 4, 6, 15, 52, 31],
    [5, 933, 1, 2, 1, 1, 3, 1, 11, 42],
    [55, 5, 621, 88, 66, 53, 56, 37, 12, 7],
    [13, 5, 29, 695, 33, 139, 51, 25, 6, 4],
    [18, 3, 26, 65, 721, 55, 52, 51, 4, 5],
    [6, 2, 10, 144, 22, 773, 8, 32, 1, 2],
    [4, 2, 18, 54, 17, 25, 868, 8, 2, 2],
    [10, 1, 5, 32, 22, 58, 3, 859, 1, 9],
    [41, 19, 2, 11, 1, 5, 4, 4, 898, 15],
    [15, 57, 3, 11, 0, 5, 2, 9, 18, 880]
])

# Class names
class_names = ['airplane', 'automobile', 'bird', 'cat', 'deer', 'dog', 'frog',
               ↳'horse', 'ship', 'truck']

# Plot confusion matrix
plt.figure(figsize=(10, 8))
sns.heatmap(confusion_matrix, annot=True, fmt='d', cmap='Blues',
            ↳xticklabels=class_names, yticklabels=class_names)
plt.xlabel('Predicted')
plt.ylabel('True')
plt.title('Confusion Matrix')
plt.show()
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