```
import pandas as pd
import numpy as np
import os
import matplotlib.pyplot as plt
import seaborn as sns
import matplotlib.patches as patches
from sklearn.metrics import precision recall curve,
average precision score
from sklearn.preprocessing import LabelEncoder
from torchvision import transforms
from PIL import Image
from tqdm import tqdm
import shutil
from sklearn.metrics import roc curve, auc
import torch
import torch.nn as nn
import torch.optim as optim
from torchvision import transforms, models
from torch.utils.data import Dataset, DataLoader, random split
from sklearn.metrics import confusion matrix, ConfusionMatrixDisplay
test base dir ="/kaggle/input/gtsrb-german-traffic-sign"
# File paths for CSV files
test_csv_path = '/kaggle/input/gtsrb-german-traffic-sign/Test.csv'
# Load CSV files
test df = pd.read csv(test csv path)
# Load CSV files
test df = pd.read csv(test csv path)
# Custom Dataset Class
class CustomDataset(Dataset):
    def init (self, dataframe, base dir, transform=None):
        self.dataframe = dataframe
        self.base dir = base dir
        self.transform = transform
    def __len__(self):
        return len(self.dataframe)
    def getitem (self, idx):
        # Concatenate base directory with relative image path
        img path = os.path.join(self.base dir,
str(self.dataframe.iloc[idx, -1]))
        label = int(self.dataframe.iloc[idx, -2]) # Ensure this is
the correct column for labels
        try:
```

```
image = Image.open(img path).convert("RGB")
        except FileNotFoundError:
            raise ValueError(f"Image not found: {img path}")
        if self.transform:
            image = self.transform(image)
        return image, label
test transforms = transforms.Compose([
    transforms.Resize((224, 224)),
    transforms.ToTensor(),
    transforms.Normalize(mean=[0.485, 0.456, 0.406], std=[0.229,
0.224, 0.225]),
1)
# Create Datasets
test dataset = CustomDataset(test df, test base dir,
transform=test transforms)
# Create Data Loaders
test loader = DataLoader(test dataset, batch size=1, shuffle=False)
print(f"Total images in test dataset: {len(test dataset)}")
Total images in test dataset: 12630
# Load the trained model
model path =
"/kaggle/input/theiss model/pytorch/default/1/mobilenet v2 traffic sig
ns.pth" # Replace with the path to your saved model
device = torch.device("cuda" if torch.cuda.is available() else "cpu")
# Reconstruct the model architecture
model =
models.mobilenet v2(weights=models.MobileNet V2 Weights.IMAGENET1K V1)
model.classifier[1] = nn.Linear(model.last_channel, 43) # 43 classes
in vour dataset
model.load state dict(torch.load(model path))
model = model.to(device)
model.eval() # Set the model to evaluation mode
<ipython-input-10-29859e955a0a>:8: FutureWarning: You are using
`torch.load` with `weights only=False` (the current default value),
which uses the default pickle module implicitly. It is possible to
construct malicious pickle data which will execute arbitrary code
during unpickling (See
https://github.com/pytorch/pytorch/blob/main/SECURITY.md#untrusted-
models for more details). In a future release, the default value for
`weights_only` will be flipped to `True`. This limits the functions
that could be executed during unpickling. Arbitrary objects will no
longer be allowed to be loaded via this mode unless they are
```

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explicitly allowlisted by the user via
`torch.serialization.add safe globals`. We recommend you start setting
`weights_only=True` for any use case where you don't have full control
of the loaded file. Please open an issue on GitHub for any issues
related to this experimental feature.
  model.load state dict(torch.load(model path))
MobileNetV2(
  (features): Sequential(
    (0): Conv2dNormActivation(
      (0): Conv2d(3, 32, kernel size=(3, 3), stride=(2, 2),
padding=(1, 1), bias=False)
      (1): BatchNorm2d(32, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
      (2): ReLU6(inplace=True)
    (1): InvertedResidual(
      (conv): Sequential(
        (0): Conv2dNormActivation(
          (0): Conv2d(32, 32, kernel size=(3, 3), stride=(1, 1),
padding=(1, 1), groups=32, bias=False)
          (1): BatchNorm2d(32, eps=1e-05, momentum=0.1, affine=True,
track_running_stats=True)
          (2): ReLU6(inplace=True)
        (1): Conv2d(32, 16, kernel size=(1, 1), stride=(1, 1),
bias=False)
        (2): BatchNorm2d(16, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
    (2): InvertedResidual(
      (conv): Sequential(
        (0): Conv2dNormActivation(
          (0): Conv2d(16, 96, kernel size=(1, 1), stride=(1, 1),
bias=False)
          (1): BatchNorm2d(96, eps=1e-05, momentum=0.1, affine=True,
track_running_stats=True)
          (2): ReLU6(inplace=True)
        (1): Conv2dNormActivation(
          (0): Conv2d(96, 96, kernel size=(3, 3), stride=(2, 2),
padding=(1, 1), groups=96, bias=False)
          (1): BatchNorm2d(96, eps=1e-05, momentum=0.1, affine=True,
track_running_stats=True)
          (2): ReLU6(inplace=True)
        (2): Conv2d(96, 24, \text{ kernel size}=(1, 1), \text{ stride}=(1, 1),
bias=False)
        (3): BatchNorm2d(24, eps=1e-05, momentum=0.1, affine=True,
```

```
track_running stats=True)
    (3): InvertedResidual(
      (conv): Sequential(
        (0): Conv2dNormActivation(
          (0): Conv2d(24, 144, kernel size=(1, 1), stride=(1, 1),
bias=False)
          (1): BatchNorm2d(144, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
          (2): ReLU6(inplace=True)
        (1): Conv2dNormActivation(
          (0): Conv2d(144, 144, kernel size=(3, 3), stride=(1, 1),
padding=(1, 1), groups=144, bias=False)
          (1): BatchNorm2d(144, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
          (2): ReLU6(inplace=True)
        (2): Conv2d(144, 24, kernel size=(1, 1), stride=(1, 1),
bias=False)
        (3): BatchNorm2d(24, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
    (4): InvertedResidual(
      (conv): Sequential(
        (0): Conv2dNormActivation(
          (0): Conv2d(24, 144, kernel size=(1, 1), stride=(1, 1),
bias=False)
          (1): BatchNorm2d(144, eps=1e-05, momentum=0.1, affine=True,
track_running_stats=True)
          (2): ReLU6(inplace=True)
        (1): Conv2dNormActivation(
          (0): Conv2d(144, 144, kernel size=(3, 3), stride=(2, 2),
padding=(1, 1), groups=144, bias=False)
          (1): BatchNorm2d(144, eps=1e-05, momentum=0.1, affine=True,
track_running_stats=True)
          (2): ReLU6(inplace=True)
        (2): Conv2d(144, 32, kernel size=(1, 1), stride=(1, 1),
bias=False)
        (3): BatchNorm2d(32, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
    (5): InvertedResidual(
      (conv): Sequential(
```

```
(0): Conv2dNormActivation(
          (0): Conv2d(32, 192, kernel size=(1, 1), stride=(1, 1),
bias=False)
          (1): BatchNorm2d(192, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
          (2): ReLU6(inplace=True)
        (1): Conv2dNormActivation(
          (0): Conv2d(192, 192, kernel size=(3, 3), stride=(1, 1),
padding=(1, 1), groups=192, bias=False)
          (1): BatchNorm2d(192, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
          (2): ReLU6(inplace=True)
        (2): Conv2d(192, 32, kernel size=(1, 1), stride=(1, 1),
bias=False)
        (3): BatchNorm2d(32, eps=1e-05, momentum=0.1, affine=True,
track_running_stats=True)
    (6): InvertedResidual(
      (conv): Sequential(
        (0): Conv2dNormActivation(
          (0): Conv2d(32, 192, kernel size=(1, 1), stride=(1, 1),
bias=False)
          (1): BatchNorm2d(192, eps=1e-05, momentum=0.1, affine=True,
track_running_stats=True)
          (2): ReLU6(inplace=True)
        (1): Conv2dNormActivation(
          (0): Conv2d(192, 192, kernel size=(3, 3), stride=(1, 1),
padding=(1, 1), groups=192, bias=False)
          (1): BatchNorm2d(192, eps=1e-05, momentum=0.1, affine=True,
track_running_stats=True)
          (2): ReLU6(inplace=True)
        (2): Conv2d(192, 32, kernel size=(1, 1), stride=(1, 1),
bias=False)
        (3): BatchNorm2d(32, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
      )
    (7): InvertedResidual(
      (conv): Sequential(
        (0): Conv2dNormActivation(
          (0): Conv2d(32, 192, kernel size=(1, 1), stride=(1, 1),
bias=False)
          (1): BatchNorm2d(192, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
```

```
(2): ReLU6(inplace=True)
        )
        (1): Conv2dNormActivation(
          (0): Conv2d(192, 192, kernel size=(3, 3), stride=(2, 2),
padding=(1, 1), groups=192, bias=False)
          (1): BatchNorm2d(192, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
          (2): ReLU6(inplace=True)
        (2): Conv2d(192, 64, kernel size=(1, 1), stride=(1, 1),
bias=False)
        (3): BatchNorm2d(64, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
    (8): InvertedResidual(
      (conv): Sequential(
        (0): Conv2dNormActivation(
          (0): Conv2d(64, 384, \text{kernel size}=(1, 1), \text{stride}=(1, 1),
bias=False)
          (1): BatchNorm2d(384, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
          (2): ReLU6(inplace=True)
        (1): Conv2dNormActivation(
          (0): Conv2d(384, 384, kernel size=(3, 3), stride=(1, 1),
padding=(1, 1), groups=384, bias=False)
          (1): BatchNorm2d(384, eps=1e-05, momentum=0.1, affine=True,
track_running_stats=True)
          (2): ReLU6(inplace=True)
        (2): Conv2d(384, 64, kernel size=(1, 1), stride=(1, 1),
bias=False)
        (3): BatchNorm2d(64, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
      )
    (9): InvertedResidual(
      (conv): Sequential(
        (0): Conv2dNormActivation(
          (0): Conv2d(64, 384, \text{kernel size}=(1, 1), \text{stride}=(1, 1),
bias=False)
          (1): BatchNorm2d(384, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
          (2): ReLU6(inplace=True)
        (1): Conv2dNormActivation(
          (0): Conv2d(384, 384, kernel size=(3, 3), stride=(1, 1),
padding=(1, 1), groups=384, bias=False)
```

```
(1): BatchNorm2d(384, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
          (2): ReLU6(inplace=True)
        (2): Conv2d(384, 64, kernel size=(1, 1), stride=(1, 1),
bias=False)
        (3): BatchNorm2d(64, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
    (10): InvertedResidual(
      (conv): Sequential(
        (0): Conv2dNormActivation(
          (0): Conv2d(64, 384, \text{kernel size}=(1, 1), \text{stride}=(1, 1),
bias=False)
          (1): BatchNorm2d(384, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
          (2): ReLU6(inplace=True)
        (1): Conv2dNormActivation(
          (0): Conv2d(384, 384, kernel size=(3, 3), stride=(1, 1),
padding=(1, 1), groups=384, bias=False)
          (1): BatchNorm2d(384, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
          (2): ReLU6(inplace=True)
        (2): Conv2d(384, 64, kernel size=(1, 1), stride=(1, 1),
bias=False)
        (3): BatchNorm2d(64, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
      )
    (11): InvertedResidual(
      (conv): Sequential(
        (0): Conv2dNormActivation(
          (0): Conv2d(64, 384, \text{kernel size}=(1, 1), \text{stride}=(1, 1),
bias=False)
          (1): BatchNorm2d(384, eps=1e-05, momentum=0.1, affine=True,
track_running_stats=True)
          (2): ReLU6(inplace=True)
        (1): Conv2dNormActivation(
          (0): Conv2d(384, 384, kernel_size=(3, 3), stride=(1, 1),
padding=(1, 1), groups=384, bias=False)
          (1): BatchNorm2d(384, eps=1e-05, momentum=0.1, affine=True,
track_running_stats=True)
          (2): ReLU6(inplace=True)
        (2): Conv2d(384, 96, kernel size=(1, 1), stride=(1, 1),
```

```
bias=False)
        (3): BatchNorm2d(96, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
    (12): InvertedResidual(
      (conv): Sequential(
        (0): Conv2dNormActivation(
          (0): Conv2d(96, 576, \text{kernel size}=(1, 1), \text{stride}=(1, 1),
bias=False)
          (1): BatchNorm2d(576, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
          (2): ReLU6(inplace=True)
        (1): Conv2dNormActivation(
          (0): Conv2d(576, 576, kernel size=(3, 3), stride=(1, 1),
padding=(1, 1), groups=576, bias=False)
          (1): BatchNorm2d(576, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
          (2): ReLU6(inplace=True)
        (2): Conv2d(576, 96, kernel size=(1, 1), stride=(1, 1),
bias=False)
        (3): BatchNorm2d(96, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
      )
    (13): InvertedResidual(
      (conv): Sequential(
        (0): Conv2dNormActivation(
          (0): Conv2d(96, 576, \text{kernel size}=(1, 1), \text{stride}=(1, 1),
bias=False)
          (1): BatchNorm2d(576, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
          (2): ReLU6(inplace=True)
        (1): Conv2dNormActivation(
          (0): Conv2d(576, 576, kernel size=(3, 3), stride=(1, 1),
padding=(1, 1), groups=576, bias=False)
          (1): BatchNorm2d(576, eps=1e-05, momentum=0.1, affine=True,
track_running_stats=True)
          (2): ReLU6(inplace=True)
        (2): Conv2d(576, 96, kernel size=(1, 1), stride=(1, 1),
        (3): BatchNorm2d(96, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
    )
```

```
(14): InvertedResidual(
      (conv): Sequential(
        (0): Conv2dNormActivation(
          (0): Conv2d(96, 576, \text{kernel size}=(1, 1), \text{stride}=(1, 1),
bias=False)
          (1): BatchNorm2d(576, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
          (2): ReLU6(inplace=True)
        (1): Conv2dNormActivation(
          (0): Conv2d(576, 576, kernel_size=(3, 3), stride=(2, 2),
padding=(1, 1), groups=576, bias=False)
          (1): BatchNorm2d(576, eps=1e-05, momentum=0.1, affine=True,
track_running_stats=True)
          (2): ReLU6(inplace=True)
        (2): Conv2d(576, 160, kernel size=(1, 1), stride=(1, 1),
bias=False)
        (3): BatchNorm2d(160, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
      )
    (15): InvertedResidual(
      (conv): Sequential(
        (0): Conv2dNormActivation(
          (0): Conv2d(160, 960, kernel size=(1, 1), stride=(1, 1),
bias=False)
          (1): BatchNorm2d(960, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
          (2): ReLU6(inplace=True)
        (1): Conv2dNormActivation(
          (0): Conv2d(960, 960, kernel size=(3, 3), stride=(1, 1),
padding=(1, 1), groups=960, bias=False)
          (1): BatchNorm2d(960, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
          (2): ReLU6(inplace=True)
        (2): Conv2d(960, 160, kernel size=(1, 1), stride=(1, 1),
bias=False)
        (3): BatchNorm2d(160, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
    (16): InvertedResidual(
      (conv): Sequential(
        (0): Conv2dNormActivation(
          (0): Conv2d(160, 960, kernel size=(1, 1), stride=(1, 1),
bias=False)
```

```
(1): BatchNorm2d(960, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
          (2): ReLU6(inplace=True)
        (1): Conv2dNormActivation(
          (0): Conv2d(960, 960, kernel_size=(3, 3), stride=(1, 1),
padding=(1, 1), groups=960, bias=False)
          (1): BatchNorm2d(960, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
          (2): ReLU6(inplace=True)
        (2): Conv2d(960, 160, kernel size=(1, 1), stride=(1, 1),
bias=False)
        (3): BatchNorm2d(160, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
    (17): InvertedResidual(
      (conv): Sequential(
        (0): Conv2dNormActivation(
          (0): Conv2d(160, 960, kernel size=(1, 1), stride=(1, 1),
bias=False)
          (1): BatchNorm2d(960, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
          (2): ReLU6(inplace=True)
        (1): Conv2dNormActivation(
          (0): Conv2d(960, 960, kernel size=(3, 3), stride=(1, 1),
padding=(1, 1), groups=960, bias=False)
          (1): BatchNorm2d(960, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
          (2): ReLU6(inplace=True)
        (2): Conv2d(960, 320, kernel size=(1, 1), stride=(1, 1),
bias=False)
        (3): BatchNorm2d(320, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
    (18): Conv2dNormActivation(
      (0): Conv2d(320, 1280, kernel size=(1, 1), stride=(1, 1),
bias=False)
      (1): BatchNorm2d(1280, eps=1e-05, momentum=0.1, affine=True,
track running stats=True)
      (2): ReLU6(inplace=True)
    )
  (classifier): Sequential(
    (0): Dropout(p=0.2, inplace=False)
    (1): Linear(in features=1280, out features=43, bias=True)
```

```
)
def test model(model, test loader, device):
    Test the model and return accuracy, labels, and predictions.
   Args:
        model (torch.nn.Module): The model to test.
        test loader (torch.utils.data.DataLoader): DataLoader for the
test dataset.
        device (torch.device): The device to perform testing on (CPU
or GPU).
    Returns:
        tuple: A tuple containing accuracy, all true labels, and all
predicted labels.
    model.eval()
    correct = 0
    total = 0
    all labels = []
    all predictions = []
    with torch.no grad():
        for images, labels in tgdm(test loader, desc="Testing"):
            images, labels = images.to(device), labels.to(device)
            outputs = model(images)
            _, predicted = outputs.max(1)
total += labels.size(0)
            correct += predicted.eq(labels).sum().item()
            # Collect all labels and predictions for confusion matrix
            all labels.extend(labels.cpu().numpy())
            all predictions.extend(predicted.cpu().numpy())
    accuracy = 100. * correct / total
    print(f"Test Accuracy: {accuracy:.2f}%")
    return accuracy, all labels, all predictions
ccuracy, all labels, all predictions = test model(model, test loader,
device)
Testing: 100% | 12630/12630 [04:06<00:00, 51.28it/s]
Test Accuracy: 95.61%
```

```
def predict(model, image, device):
    image = image.to(device)
    with torch.no grad():
        output = model(image.unsqueeze(0)) # Add batch dimension
        probabilities = torch.nn.functional.softmax(output, dim=1)
        confidence, predicted = probabilities.max(1)
    result = {
        "predicted class": predicted.item(),
        "confidence": confidence.item()
    }
    return result
image path = "/kaggle/input/gtsrb-german-traffic-sign/Test/00000.png"
# Replace with your image path
image = Image.open(image path).convert("RGB")
transform = transforms.Compose([
    transforms.Resize((224, 224)),
    transforms.ToTensor(),
    transforms.Normalize(mean=[0.485, 0.456, 0.406], std=[0.229,
0.224, 0.225]),
1)
image tensor = transform(image)
# Perform prediction
result = predict(model, image tensor, device)
# Print the result
print("predicted_class:", result["predicted_class"])
print("confidence:", result["confidence"])
predicted class: 16
confidence: 1.0
import pandas as pd
from sklearn.metrics import confusion matrix, ConfusionMatrixDisplay
import matplotlib.pyplot as plt
def plot and save confusion_matrix(all_labels, all_predictions,
filename="confusion matrix.csv"):
    Plot and save the confusion matrix using true and predicted
labels.
    Args:
        all labels (list): True labels.
        all predictions (list): Predicted labels.
        filename (str): Filename to save the confusion matrix.
```

```
class names = [str(i) for i in range(43)] # Generate class names
from 0 to 42
    cm = confusion_matrix(all_labels, all_predictions)
    # Save confusion matrix to CSV
    pd.DataFrame(cm).to csv(filename, index=False, header=False)
    print(f"Confusion matrix saved as {filename}")
    # Plotting the confusion matrix
    disp = ConfusionMatrixDisplay(confusion matrix=cm,
display labels=class names)
    fig, ax = plt.subplots(figsize=(12, 12))
    disp.plot(cmap=plt.cm.Blues, ax=ax, xticks_rotation='vertical')
    plt.title("Confusion Matrix", fontsize=16)
plt.xlabel("Predicted label", fontsize=12)
    plt.ylabel("True label", fontsize=12)
    plt.tight layout()
    plt.show()
plot and save confusion matrix(all labels, all predictions)
Confusion matrix saved as confusion matrix.csv
```

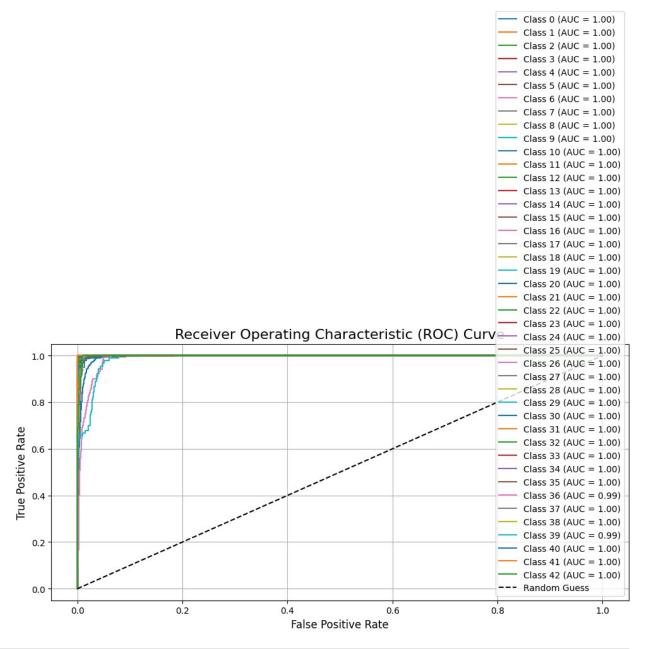
1 0 0 0 1 0 4834 0 0 0

100

```
def plot_roc_curve(model, test_loader, device, num_classes=43):
    Plot the ROC curve for a multi-class classification model.

Args:
    model (torch.nn.Module): The trained model.
    test_loader (torch.utils.data.DataLoader): DataLoader for the test dataset.
    device (torch.device): The device to perform testing on (CPU or GPU).
```

```
num classes (int): The number of classes in the dataset.
    model.eval()
    all labels = []
    all outputs = []
    # Collect predictions and true labels
    with torch.no grad():
        for images, labels in tqdm(test loader, desc="Calculating
ROC"):
            images, labels = images.to(device), labels.to(device)
            outputs = model(images)
            all labels.extend(labels.cpu().numpy())
            all outputs.extend(outputs.cpu().numpy())
    all_labels = np.array(all_labels)
    all outputs = np.array(all outputs)
    # Compute ROC curve and AUC for each class
    fpr = \{\}
    tpr = {}
    roc auc = \{\}
    for i in range(num classes):
        true binary = (all labels == i).astype(int) # Convert labels
to binary for each class
        fpr[i], tpr[i], _ = roc_curve(true_binary, all_outputs[:, i])
        roc auc[i] = auc(fpr[i], tpr[i])
    # Plot ROC curve for each class
    plt.figure(figsize=(10, 8))
    for i in range(num classes):
        plt.plot(fpr[i], tpr[i], label=f"Class {i} (AUC =
{roc auc[i]:.2f})")
    # Plot the diagonal line
    plt.plot([0, 1], [0, 1], 'k--', label="Random Guess")
    plt.title("Receiver Operating Characteristic (ROC) Curve",
fontsize=16)
    plt.xlabel("False Positive Rate", fontsize=12)
    plt.ylabel("True Positive Rate", fontsize=12)
    plt.legend(loc="lower right")
    plt.grid()
    plt.tight_layout()
    plt.show()
plot roc curve(model, test loader, device, num classes=43)
Calculating ROC: 100% | 12630/12630 [02:21<00:00, 89.07it/s]
```



```
from sklearn.metrics import precision_recall_curve,
average_precision_score
import matplotlib.pyplot as plt
import numpy as np

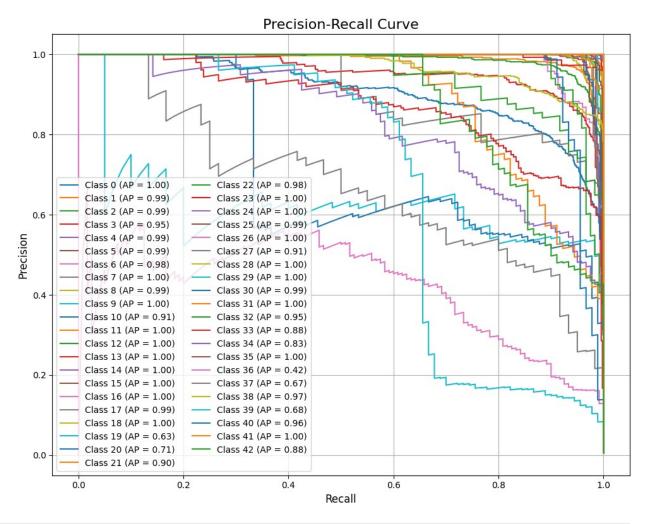
def plot_precision_recall_curve(model, test_loader, device,
num_classes=43):
    Plot the Precision-Recall curve for a multi-class classification
model.

Args:
```

```
model (torch.nn.Module): The trained model.
        test loader (torch.utils.data.DataLoader): DataLoader for the
test dataset.
        device (torch.device): The device to perform testing on (CPU
or GPU).
        num classes (int): The number of classes in the dataset.
    model.eval()
    all labels = []
    all outputs = []
    # Collect predictions and true labels
    with torch.no grad():
        for images, labels in tgdm(test loader, desc="Calculating
Precision-Recall"):
            images, labels = images.to(device), labels.to(device)
            outputs = model(images)
            all labels.extend(labels.cpu().numpy())
            all outputs.extend(outputs.cpu().numpy())
    all labels = np.array(all labels)
    all outputs = np.array(all outputs)
    # Compute Precision-Recall curve and Average Precision for each
class
    precision = {}
    recall = {}
    average precision = {}
    for i in range(num classes):
        true binary = (all labels == i).astype(int) # Convert labels
to binary for each class
        precision[i], recall[i], =
precision_recall_curve(true_binary, all_outputs[:, i])
        average precision[i] = average precision score(true binary,
all outputs[:, i])
    # Plot Precision-Recall curve for each class
    plt.figure(figsize=(10, 8))
    for i in range(num classes):
        plt.plot(recall[i], precision[i], label=f"Class {i} (AP =
{average precision[i]:.2f})")
    plt.title("Precision-Recall Curve", fontsize=16)
    plt.xlabel("Recall", fontsize=12)
    plt.ylabel("Precision", fontsize=12)
    plt.legend(loc="lower left", fontsize=10, ncol=2)
    plt.grid()
    plt.tight_layout()
    plt.show()
```

```
plot_precision_recall_curve(model, test_loader, device,
num_classes=43)
```

Calculating Precision-Recall: 100%| 12630/12630 [02:15<00:00, 93.00it/s]

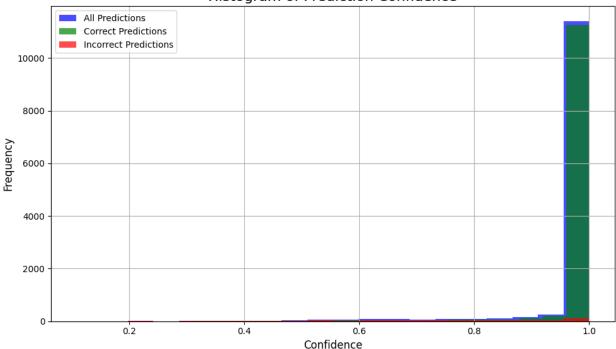


```
def plot_prediction_confidence(model, test_loader, device,
num_bins=10):
    Plot a histogram of prediction confidence for the test dataset.

Args:
        model (torch.nn.Module): The trained model.
        test_loader (torch.utils.data.DataLoader): DataLoader for the test dataset.
        device (torch.device): Device to perform testing on (CPU or GPU).
        num_bins (int): Number of bins for the histogram.
"""
```

```
model.eval()
    confidences = []
    correct predictions = []
    incorrect predictions = []
    with torch.no grad():
        for images, labels in tqdm(test_loader, desc="Calculating")
Confidence"):
            images, labels = images.to(device), labels.to(device)
            outputs = model(images)
            probabilities = torch.nn.functional.softmax(outputs,
dim=1)
            max confidence, predicted = probabilities.max(1)
            confidences.extend(max confidence.cpu().numpy())
            correct predictions.extend(max confidence[labels ==
predicted].cpu().numpy())
            incorrect predictions.extend(max confidence[labels !=
predicted].cpu().numpy())
    # Plot histogram
    plt.figure(figsize=(10, 6))
    plt.hist(confidences, bins=num bins, alpha=0.7, label="All
Predictions", color="blue")
    plt.hist(correct_predictions, bins=num bins, alpha=0.7,
label="Correct Predictions", color="green")
    plt.hist(incorrect predictions, bins=num bins, alpha=0.7,
label="Incorrect Predictions", color="red")
    plt.title("Histogram of Prediction Confidence", fontsize=16)
    plt.xlabel("Confidence", fontsize=12)
    plt.ylabel("Frequency", fontsize=12)
    plt.legend()
    plt.grid(True)
    plt.tight layout()
    plt.show()
plot prediction confidence(model, test loader, device, num bins=20)
Calculating Confidence: 100% | 12630/12630 [02:19<00:00,
90.33it/sl
```

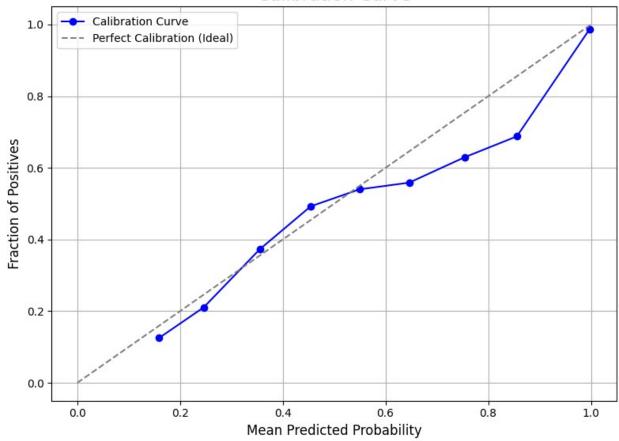




```
from sklearn.calibration import calibration curve
import matplotlib.pyplot as plt
def plot calibration curve(model, test loader, device, num bins=10):
    Plot a calibration curve to evaluate the reliability of the
model's confidence scores.
    Args:
        model (torch.nn.Module): The trained model.
        test loader (torch.utils.data.DataLoader): DataLoader for the
test dataset.
        device (torch.device): Device to perform testing on (CPU or
GPU).
        num bins (int): Number of bins for calibration.
    model.eval()
    all labels = []
    all confidences = []
    all predictions = []
    # Collect predictions and confidences
    with torch.no grad():
        for images, labels in tqdm(test loader, desc="Calculating
Calibration Data"):
            images, labels = images.to(device), labels.to(device)
            outputs = model(images)
```

```
probabilities = torch.nn.functional.softmax(outputs,
dim=1)
           max confidence, predicted = probabilities.max(1)
           all confidences.extend(max confidence.cpu().numpy())
           all labels.extend(labels.cpu().numpy())
           all predictions.extend(predicted.cpu().numpy())
   # Binarize the predictions for calibration curve
   true labels = (np.array(all labels) ==
np.array(all predictions)).astype(int)
    confidences = np.array(all confidences)
   # Compute calibration curve
    prob true, prob pred = calibration curve(true labels, confidences,
n bins=num bins, strategy='uniform')
   # Plot calibration curve
   plt.figure(figsize=(8, 6))
   plt.plot(prob_pred, prob true, marker='o', label="Calibration")
Curve", color="blue")
   plt.plot([0, 1], [0, 1], 'k--', label="Perfect Calibration
(Ideal)", color="gray")
   plt.title("Calibration Curve", fontsize=16)
   plt.xlabel("Mean Predicted Probability", fontsize=12)
   plt.ylabel("Fraction of Positives", fontsize=12)
   plt.legend()
   plt.grid()
   plt.tight_layout()
   plt.show()
plot calibration curve(model, test loader, device, num bins=10)
Calculating Calibration Data: 100%
[02:16<00:00, 92.28it/s]
<ipython-input-23-035aff4a45d5>:41: UserWarning: color is redundantly
defined by the 'color' keyword argument and the fmt string "k--" (->
color='k'). The keyword argument will take precedence.
  plt.plot([0, 1], [0, 1], 'k--', label="Perfect Calibration (Ideal)",
color="gray")
```

## Calibration Curve



```
from sklearn.metrics import classification report
import seaborn as sns
import matplotlib.pyplot as plt
import pandas as pd
def plot_classification_report(model, test_loader, device,
class names):
    Visualize the classification report as a heatmap.
   Args:
        model (torch.nn.Module): The trained model.
        test loader (torch.utils.data.DataLoader): DataLoader for the
test dataset.
        device (torch.device): Device to perform testing on (CPU or
GPU).
        class names (list): List of class names corresponding to the
dataset labels.
    0.00\,0
    model.eval()
    all labels = []
```

```
all predictions = []
   # Collect predictions and true labels
   with torch.no grad():
        for images, labels in tgdm(test loader, desc="Calculating
Classification Report Data"):
            images, labels = images.to(device), labels.to(device)
            outputs = model(images)
            _, predicted = outputs.max(1)
            all labels.extend(labels.cpu().numpy())
            all predictions.extend(predicted.cpu().numpy())
   # Generate classification report
    report = classification_report(all_labels, all_predictions,
target names=class names, output dict=True)
   # Convert to DataFrame for better visualization
    report df = pd.DataFrame(report).transpose()
   # Plot heatmap
   plt.figure(figsize=(10, len(class names) // 2))
    sns.heatmap(report df.iloc[:-1, :-1], annot=True, cmap="Blues",
fmt=".2f", cbar=False, linewidths=0.5)
   plt.title("Classification Report Heatmap", fontsize=16)
   plt.xlabel("Metrics", fontsize=12)
   plt.ylabel("Classes", fontsize=12)
   plt.tight_layout()
   plt.show()
# Example class names (replace with your actual class names)
class_names = [str(i) for i in range(43)]
# Plot the classification report heatmap
plot classification report(model, test loader, device, class names)
Calculating Classification Report Data: 100% | 12630/12630
[02:18<00:00, 91.04it/s]
```

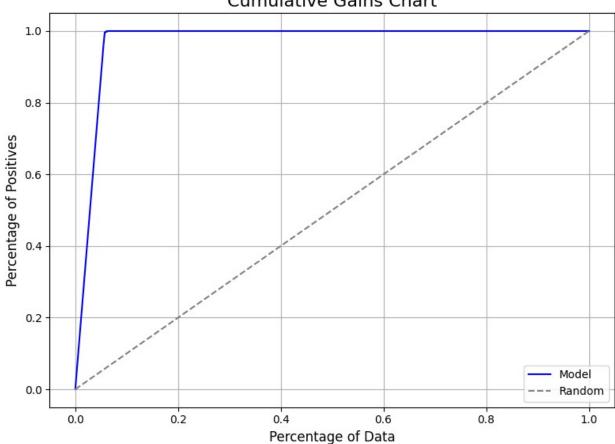
Classification Report Heatmap

Classification Report Heatmap			
0 -	1.00	1.00	1.00
1 -	0.98	1.00	0.99
2 -	0.99	1.00	0.99
3 -	0.97	0.94	0.95
4 -	1.00	0.99	0.99
5 -	0.98	0.96	0.97
6 -	0.96	0.91	0.93
7 -	1.00	1.00	1.00
8 -	0.99	0.95	0.97
9 -	0.98	1.00	0.99
10 -	1.00	0.97	0.99
11 -	0.95	0.99	0.97
12 -	1.00	0.96	0.98
13 -	0.99	1.00	0.99
14 -	0.99	1.00	1.00
15 -	0.80	1.00	0.89
16 -	0.99	1.00	1.00
17 -	1.00	0.97	0.98
18 -	1.00	0.98	0.99
19 -	0.48	0.83	0.61
20 -	0.64	0.40	0.49
21 -	0.87	0.96	0.91
22 -	1.00	0.93	0.96
23 -	0.94	1.00	0.97
24 -	0.99	0.98	0.98
25 -	0.98	0.97	0.97
26 -	0.99	1.00	0.99
27 -	0.84	0.82	0.83
28 -	1.00	1.00	1.00
29 -	1.00	1.00	1.00

```
import numpy as np
import matplotlib.pyplot as plt
from sklearn.metrics import roc curve
def plot cumulative gains(model, test loader, device):
    Plot a Cumulative Gains Chart.
    Aras:
        model (torch.nn.Module): The trained model.
        test loader (torch.utils.data.DataLoader): DataLoader for the
test dataset.
        device (torch.device): Device to perform testing on (CPU or
GPU).
    model.eval()
    all labels = []
    all probabilities = []
    # Collect true labels and model probabilities
    with torch.no grad():
        for images, labels in tgdm(test loader, desc="Calculating
Cumulative Gains Data"):
            images, labels = images.to(device), labels.to(device)
            outputs = model(images)
            probabilities = torch.nn.functional.softmax(outputs,
dim=1)
            all probabilities.extend(probabilities.cpu().numpy())
            all labels.extend(labels.cpu().numpy())
    all labels = np.array(all labels)
    all probabilities = np.array(all probabilities)
    # Assuming binary classification (adjust for multi-class if
necessary)
    positive class = 1 # Define the positive class
    probabilities_positive = all_probabilities[:, positive_class]
    # Sort probabilities and labels together
    sorted indices = np.argsort(probabilities positive)[::-1]
    sorted labels = all labels[sorted indices]
    # Calculate cumulative gains
    total positives = np.sum(sorted labels == positive class)
    cumulative positives = np.cumsum(sorted labels == positive class)
    percentage of data = np.arange(\frac{1}{1}, len(sorted labels) + \frac{1}{1}) /
len(sorted labels)
    percentage of positives = cumulative positives / total positives
    # Plot cumulative gains
```

```
plt.figure(figsize=(8, 6))
    plt.plot(percentage of data, percentage of positives,
label="Model", color="blue")
    plt.plot([0, 1], [0, 1], 'k--', label="Random", color="gray")
    plt.title("Cumulative Gains Chart", fontsize=16)
    plt.xlabel("Percentage of Data", fontsize=12)
    plt.ylabel("Percentage of Positives", fontsize=12)
    plt.legend()
    plt.grid()
    plt.tight layout()
    plt.show()
plot cumulative gains(model, test loader, device)
Calculating Cumulative Gains Data: 100%
[02:16<00:00, 92.73it/s]
<ipython-input-27-25354360c501>:47: UserWarning: color is redundantly
defined by the 'color' keyword argument and the fmt string "k--" (->
color='k'). The keyword argument will take precedence.
  plt.plot([0, 1], [0, 1], 'k--', label="Random", color="gray")
```





```
import numpy as np
import matplotlib.pyplot as plt
def plot lift curve(model, test loader, device):
    Plot a Lift Curve.
   Args:
        model (torch.nn.Module): The trained model.
        test loader (torch.utils.data.DataLoader): DataLoader for the
test dataset.
        device (torch.device): Device to perform testing on (CPU or
GPU).
    0.00
    model.eval()
    all labels = []
    all probabilities = []
    # Collect true labels and model probabilities
    with torch.no grad():
        for images, labels in tqdm(test loader, desc="Calculating Lift
Curve Data"):
            images, labels = images.to(device), labels.to(device)
            outputs = model(images)
            probabilities = torch.nn.functional.softmax(outputs,
dim=1)
            all_probabilities.extend(probabilities.cpu().numpy())
            all labels.extend(labels.cpu().numpy())
    all labels = np.array(all labels)
    all probabilities = np.array(all probabilities)
    # Assuming binary classification (adjust for multi-class if
necessary)
    positive class = 1 # Define the positive class
    probabilities positive = all probabilities[:, positive class]
    # Sort probabilities and labels together
    sorted indices = np.argsort(probabilities positive)[::-1]
    sorted labels = all labels[sorted indices]
    # Calculate cumulative gains
    total positives = np.sum(sorted labels == positive class)
    cumulative positives = np.cumsum(sorted labels == positive class)
    percentage of data = np.arange(1, len(sorted labels) + 1) /
len(sorted labels)
    lift = cumulative positives / (percentage of data *
total positives)
    # Plot lift curve
```

