

master-thesis copy

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Getting Started We will be using TensorFlow and Keras for data augmentation and matplotlib for displaying the images.

```
[1]: from glob import glob
import pandas as pd
import cv2
from scripts.visualization import Visualization
import matplotlib.pyplot as plt
import numpy as np
```

Reading Data

```
[2]: # Path to all data
data_dir = './lgg-mri-segmentation/kaggle_3m'

# img size
IMG_SIZE = 512
```

```
[3]: images_paths = []
masks_paths = glob(f'{data_dir}/*/*_mask*')

for i in masks_paths:
    images_paths.append(i.replace('_mask', ''))

df = pd.DataFrame(data= {'images_paths': images_paths, 'masks_paths':
    ↪masks_paths})
df.head()
```

```
[3]:                                     images_paths
0  ./lgg-mri-segmentation/kaggle_3m/TCGA_CS_6667_... \
1  ./lgg-mri-segmentation/kaggle_3m/TCGA_CS_6667_...
2  ./lgg-mri-segmentation/kaggle_3m/TCGA_CS_6667_...
3  ./lgg-mri-segmentation/kaggle_3m/TCGA_CS_6667_...
4  ./lgg-mri-segmentation/kaggle_3m/TCGA_CS_6667_...

                                     masks_paths
0  ./lgg-mri-segmentation/kaggle_3m/TCGA_CS_6667_...
1  ./lgg-mri-segmentation/kaggle_3m/TCGA_CS_6667_...
```

```

2 ./lgg-mri-segmentation/kaggle_3m/TCGA_CS_6667_...
3 ./lgg-mri-segmentation/kaggle_3m/TCGA_CS_6667_...
4 ./lgg-mri-segmentation/kaggle_3m/TCGA_CS_6667_...

```

```

[4]: def pos_neg_diagnosis(masks_paths):
      value = np.max(cv2.imread(masks_paths))
      if value > 0 :
          return 1
      else:
          return 0

df['label'] = df['masks_paths'].apply(lambda x: pos_neg_diagnosis(x))
df

```

```

[4]:                                     images_paths
0      ./lgg-mri-segmentation/kaggle_3m/TCGA_CS_6667_... \
1      ./lgg-mri-segmentation/kaggle_3m/TCGA_CS_6667_...
2      ./lgg-mri-segmentation/kaggle_3m/TCGA_CS_6667_...
3      ./lgg-mri-segmentation/kaggle_3m/TCGA_CS_6667_...
4      ./lgg-mri-segmentation/kaggle_3m/TCGA_CS_6667_...
...
3924    ./lgg-mri-segmentation/kaggle_3m/TCGA_FG_A60K_...
3925    ./lgg-mri-segmentation/kaggle_3m/TCGA_FG_A60K_...
3926    ./lgg-mri-segmentation/kaggle_3m/TCGA_FG_A60K_...
3927    ./lgg-mri-segmentation/kaggle_3m/TCGA_FG_A60K_...
3928    ./lgg-mri-segmentation/kaggle_3m/TCGA_FG_A60K_...

                                     masks_paths  label
0      ./lgg-mri-segmentation/kaggle_3m/TCGA_CS_6667_...      0
1      ./lgg-mri-segmentation/kaggle_3m/TCGA_CS_6667_...      0
2      ./lgg-mri-segmentation/kaggle_3m/TCGA_CS_6667_...      0
3      ./lgg-mri-segmentation/kaggle_3m/TCGA_CS_6667_...      0
4      ./lgg-mri-segmentation/kaggle_3m/TCGA_CS_6667_...      0
...
3924    ./lgg-mri-segmentation/kaggle_3m/TCGA_FG_A60K_...      0
3925    ./lgg-mri-segmentation/kaggle_3m/TCGA_FG_A60K_...      0
3926    ./lgg-mri-segmentation/kaggle_3m/TCGA_FG_A60K_...      0
3927    ./lgg-mri-segmentation/kaggle_3m/TCGA_FG_A60K_...      0
3928    ./lgg-mri-segmentation/kaggle_3m/TCGA_FG_A60K_...      0

[3929 rows x 3 columns]

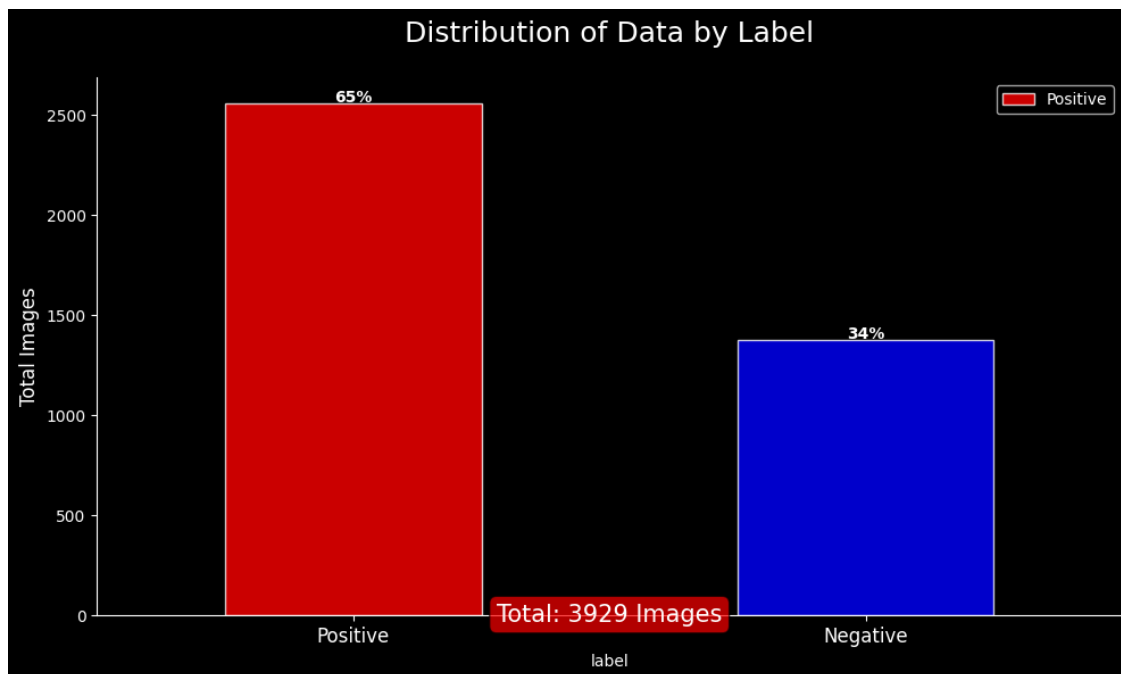
```

Data Distribution

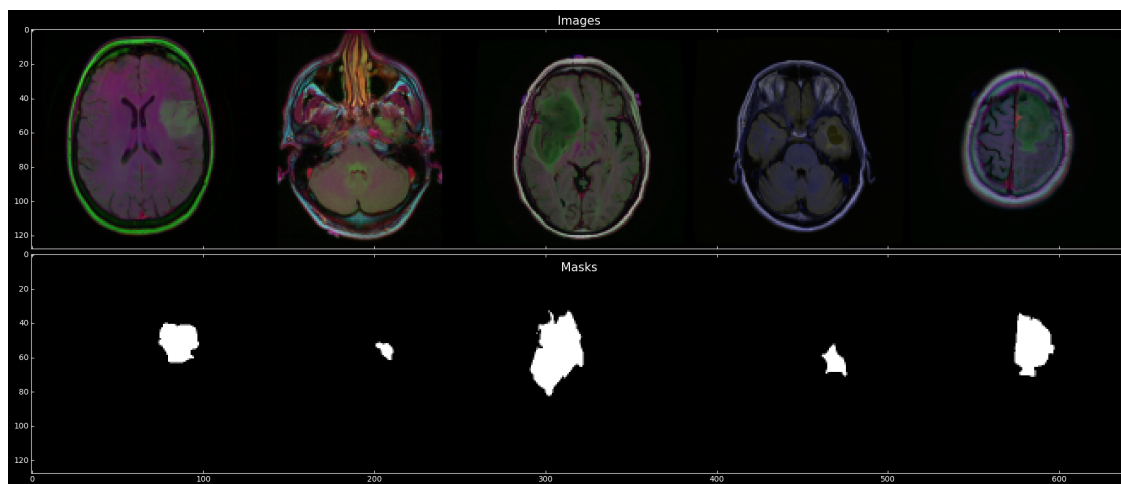
```

[5]: visualization = Visualization(df)
      visualization.plot_distribution_grouped_by_label()

```



```
[6]: visualization = Visualization(df)
      visualization.plot_images_and_masks()
```



Data Loading In the code below, we have loaded 80% training, 10% validation, and a 10% test set with labels and metadata.

```
[7]: mask_df = df[df['label'] == 1]
      mask_df.shape
```

[7]: (1373, 3)

```
[8]: from sklearn.model_selection import train_test_split
# Split df into train_df and val_df
train_df, val_df = train_test_split(mask_df, stratify=mask_df.label,
    ↪test_size=0.1)
train_df = train_df.reset_index(drop=True)
val_df = val_df.reset_index(drop=True)

# Split train_df into train_df and test_df
val_df, test_df = train_test_split(val_df, stratify=val_df.label, test_size=0.3)
val_df = val_df.reset_index(drop=True)
test_df = test_df.reset_index(drop=True)

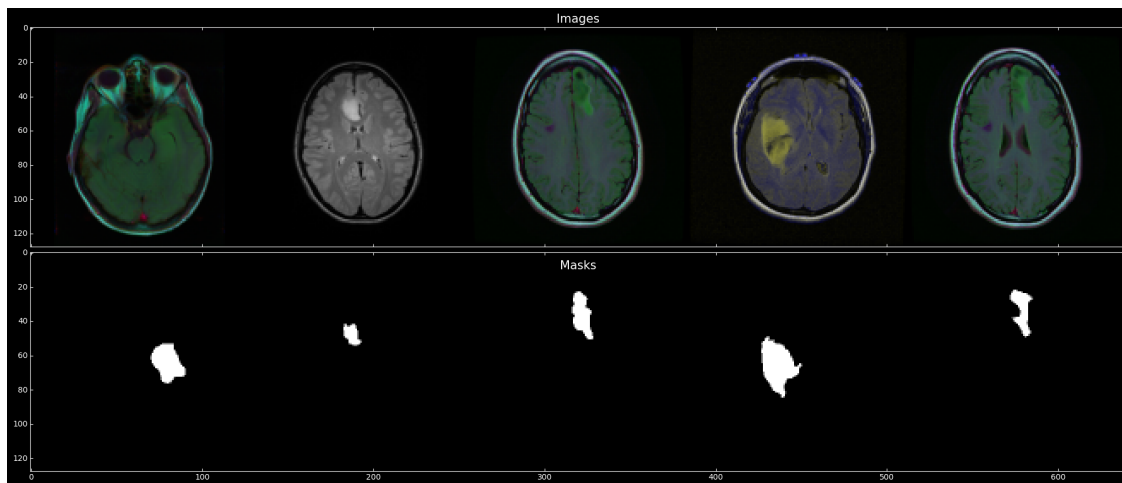
# train_df = train_df[:1000]
print(f"Train: {train_df.shape} \nVal: {val_df.shape} \nTest: {test_df.shape}")
```

Train: (1235, 3)

Val: (96, 3)

Test: (42, 3)

```
[9]: visualization = Visualization(train_df)
visualization.plot_images_and_masks()
```



GAN

```
[10]: train_df
```

```
[10]:                                     images_paths
0      ./lgg-mri-segmentation/kaggle_3m/TCGA_HT_7686_... \
1      ./lgg-mri-segmentation/kaggle_3m/TCGA_DU_A5TR_...
```

```

2      ./lgg-mri-segmentation/kaggle_3m/TCGA_DU_7299_...
3      ./lgg-mri-segmentation/kaggle_3m/TCGA_DU_7014_...
4      ./lgg-mri-segmentation/kaggle_3m/TCGA_HT_7605_...
...
1230    ./lgg-mri-segmentation/kaggle_3m/TCGA_DU_A5TW_...
1231    ./lgg-mri-segmentation/kaggle_3m/TCGA_HT_8107_...
1232    ./lgg-mri-segmentation/kaggle_3m/TCGA_HT_7608_...
1233    ./lgg-mri-segmentation/kaggle_3m/TCGA_FG_7637_...
1234    ./lgg-mri-segmentation/kaggle_3m/TCGA_DU_7014_...

                                masks_paths  label
0      ./lgg-mri-segmentation/kaggle_3m/TCGA_HT_7686_...      1
1      ./lgg-mri-segmentation/kaggle_3m/TCGA_DU_A5TR_...      1
2      ./lgg-mri-segmentation/kaggle_3m/TCGA_DU_7299_...      1
3      ./lgg-mri-segmentation/kaggle_3m/TCGA_DU_7014_...      1
4      ./lgg-mri-segmentation/kaggle_3m/TCGA_HT_7605_...      1
...
1230    ./lgg-mri-segmentation/kaggle_3m/TCGA_DU_A5TW_...      1
1231    ./lgg-mri-segmentation/kaggle_3m/TCGA_HT_8107_...      1
1232    ./lgg-mri-segmentation/kaggle_3m/TCGA_HT_7608_...      1
1233    ./lgg-mri-segmentation/kaggle_3m/TCGA_FG_7637_...      1
1234    ./lgg-mri-segmentation/kaggle_3m/TCGA_DU_7014_...      1

```

[1235 rows x 3 columns]

```

[11]: from scripts.brain_mri_dataset import BrainMriDataset
      from torch.utils.data import DataLoader
      IMG_SIZE = 64
      BATCH_SIZE = 26

      # train
      train_dataset = BrainMriDataset(df=train_df, img_size=IMG_SIZE)
      train_dataloader = DataLoader(train_dataset, batch_size=BATCH_SIZE,
      ↪num_workers=4, shuffle=True)

      # val
      val_dataset = BrainMriDataset(df=val_df, img_size=IMG_SIZE)
      val_dataloader = DataLoader(val_dataset, batch_size=BATCH_SIZE, num_workers=4,
      ↪shuffle=True)

      #test
      test_dataset = BrainMriDataset(df=test_df, img_size=IMG_SIZE)
      test_dataloader = DataLoader(test_dataset, batch_size=BATCH_SIZE,
      ↪num_workers=4, shuffle=True)

```

```

[12]: images, masks, labels = next(iter(train_dataloader))

```

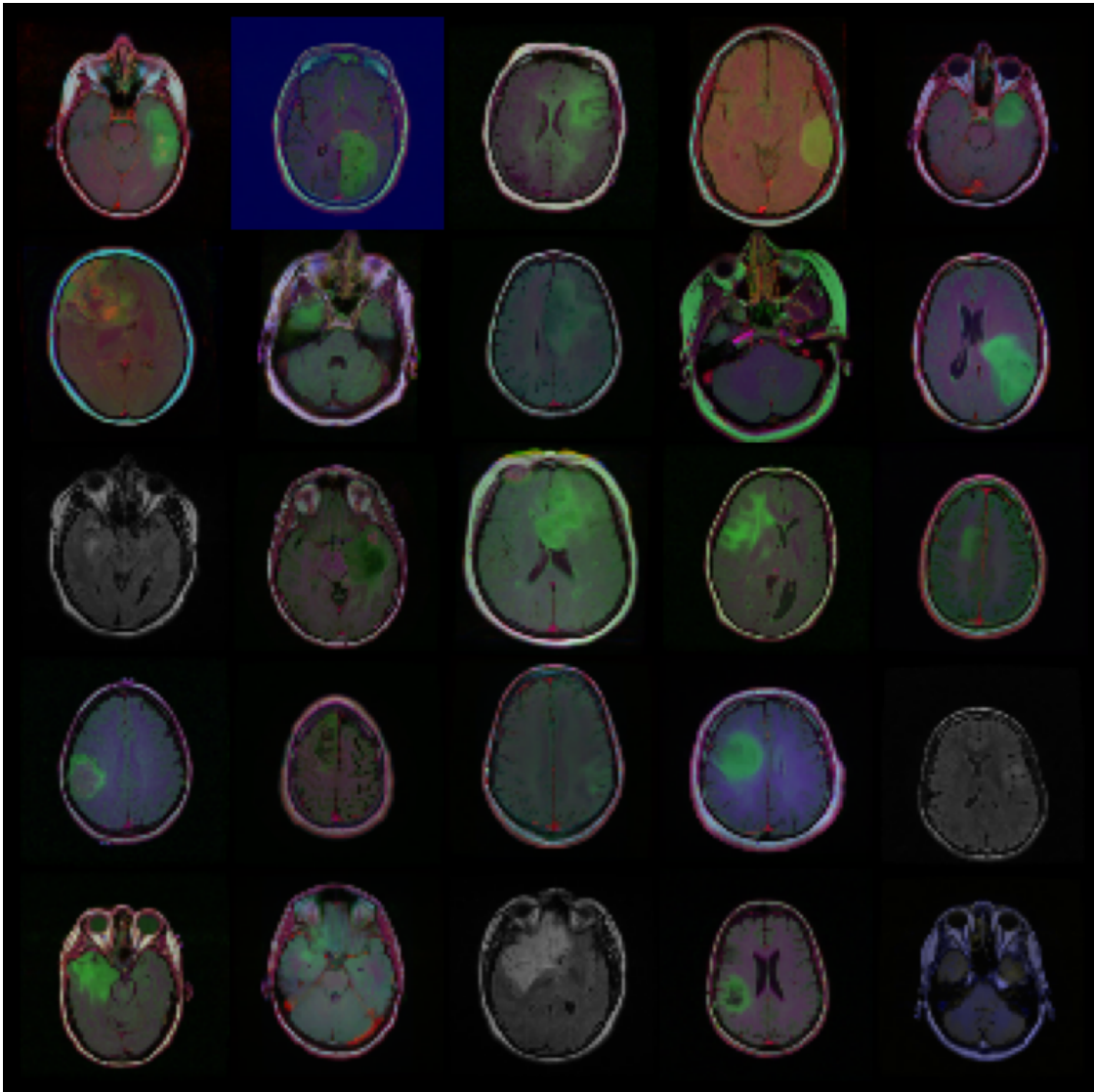
```
[13]: print(images.shape, masks.shape)
```

```
torch.Size([26, 64, 64, 3]) torch.Size([26, 64, 64, 3])
```

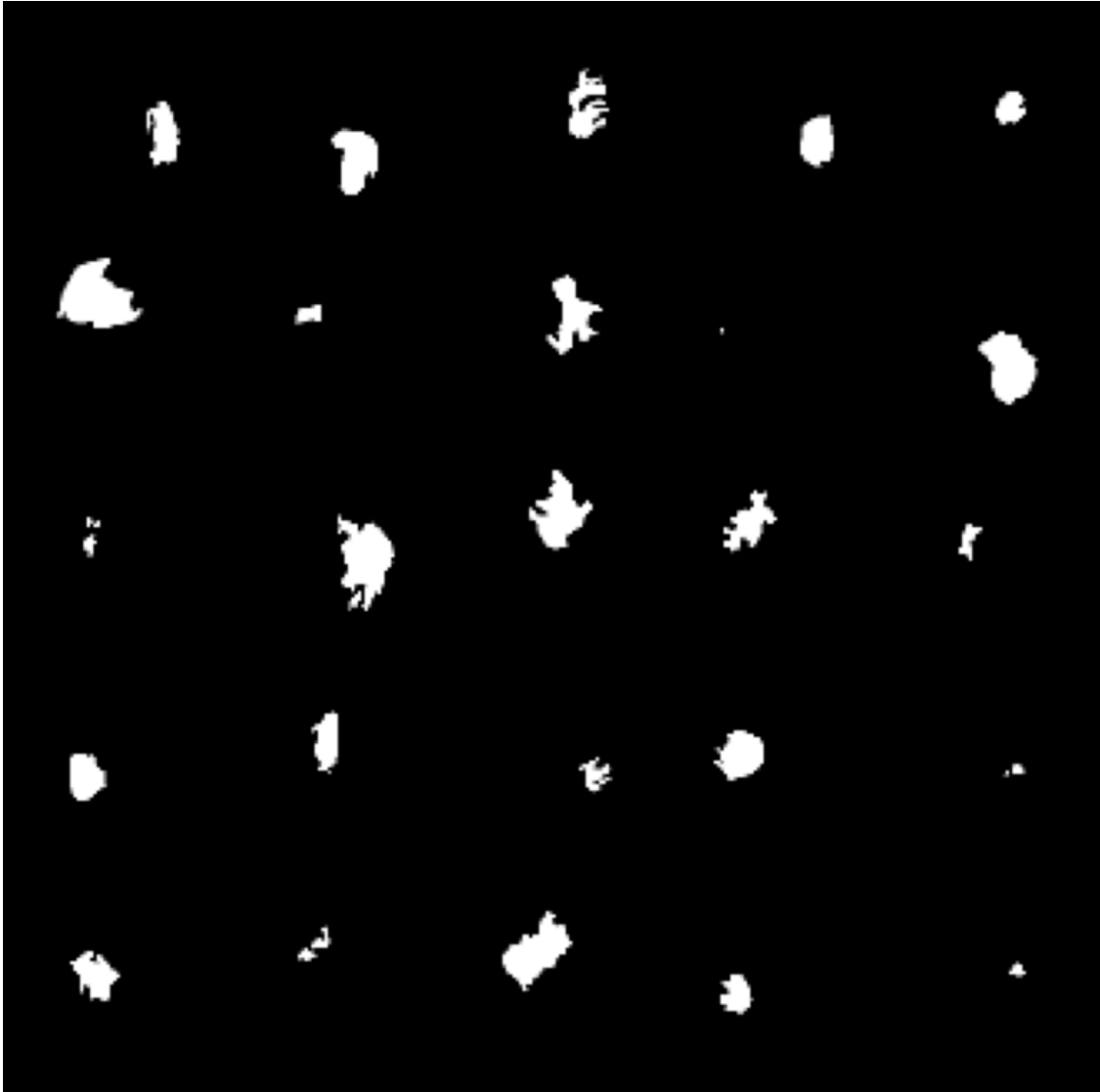
```
[14]: print(images[0].shape, masks[0].shape)
```

```
torch.Size([64, 64, 3]) torch.Size([64, 64, 3])
```

```
[15]: visualization = Visualization(train_df)  
visualization.plot_images(images)
```



```
[16]: visualization.plot_images(masks)
```



```
[17]: from scripts.gan import SimpleGAN

# Instantiate and train the GAN
gan = SimpleGAN(img_size=IMG_SIZE)
```

Metal device set to: Apple M1 Pro
Model: "Discriminator"

Layer (type)	Output Shape	Param #
Discriminator-Hidden-Layer-1 (Conv2D)	(None, 32, 32, 64)	3136

Discriminator-Hidden-Layer-Activation-1 (LeakyReLU)	(None, 32, 32, 64)	0
Discriminator-Hidden-Layer-2 (Conv2D)	(None, 16, 16, 128)	131200
Discriminator-Hidden-Layer-Activation-2 (LeakyReLU)	(None, 16, 16, 128)	0
Discriminator-Hidden-Layer-3 (Conv2D)	(None, 8, 8, 256)	524544
Discriminator-Hidden-Layer-Activation-3 (LeakyReLU)	(None, 8, 8, 256)	0
Discriminator-Flatten-Layer (Flatten)	(None, 16384)	0
Discriminator-Flatten-Layer-Dropout (Dropout)	(None, 16384)	0
Discriminator-Output-Layer (Dense)	(None, 1)	16385

```
=====
Total params: 675,265
Trainable params: 0
Non-trainable params: 675,265
```

```
-----
Model: "Generator"
```

Layer (type)	Output Shape	Param #
Generator-Hidden-Layer-1 (Dense)	(None, 8192)	827392
Generator-Hidden-Layer-Reshape-1 (Reshape)	(None, 8, 8, 128)	0
Generator-Hidden-Layer-2 (Conv2DTranspose)	(None, 16, 16, 128)	262272
Generator-Hidden-Layer-Activation-2 (ReLU)	(None, 16, 16, 128)	0
Generator-Hidden-Layer-3 (Conv2DTranspose)	(None, 32, 32, 256)	524544

Generator-Hidden-Layer-Activation-3 (ReLU)	(None, 32, 32, 256)	0
Generator-Hidden-Layer-4 (Conv2DTranspose)	(None, 64, 64, 512)	2097664
Generator-Hidden-Layer-Activation-4 (ReLU)	(None, 64, 64, 512)	0
Generator-Output-Layer (Conv2D)	(None, 64, 64, 3)	38403

```

=====
Total params: 3,750,275
Trainable params: 3,750,275
Non-trainable params: 0

```

```

-----
WARNING:tensorflow:Error in loading the saved optimizer state. As a result, your
model is starting with a freshly initialized optimizer.
WARNING:tensorflow:No training configuration found in the save file, so the
model was *not* compiled. Compile it manually.
Models loaded.

```

```
[18]: gan.train(images, epochs=5000, batch_size=128)
```

Models are already loaded. Training skipped.

```
[19]: # Generate synthetic images
num_images = 16
generated_images = gan.generate_images(num_images)
```

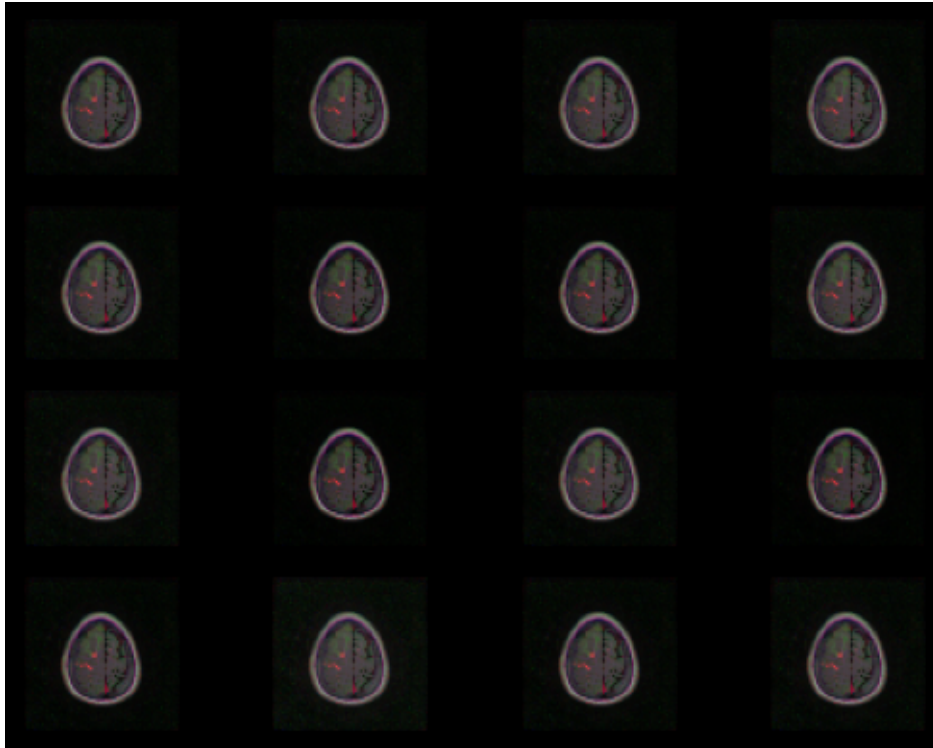
```
1/1 [=====] - 0s 79ms/step
```

```
2023-09-05 09:05:16.054403: W
tensorflow/tsl/platform/profile_utils/cpu_utils.cc:128] Failed to get CPU
frequency: 0 Hz
```

```
[20]: def display_images(images):
    fig, axs = plt.subplots(4,4)
    count = 0
    for i in range(4):
        for j in range(4):
            axs[i, j].imshow((images[count] * 0.5) + 0.5)
            axs[i, j].axis('off')
            count += 1
    plt.show()
```

```
[21]: # Display a few samples from the dataset
```

```
[22]: # Display the generated images
display_images(generated_images)
```



```
[23]: print(generated_images.shape)
```

```
(16, 64, 64, 3)
```

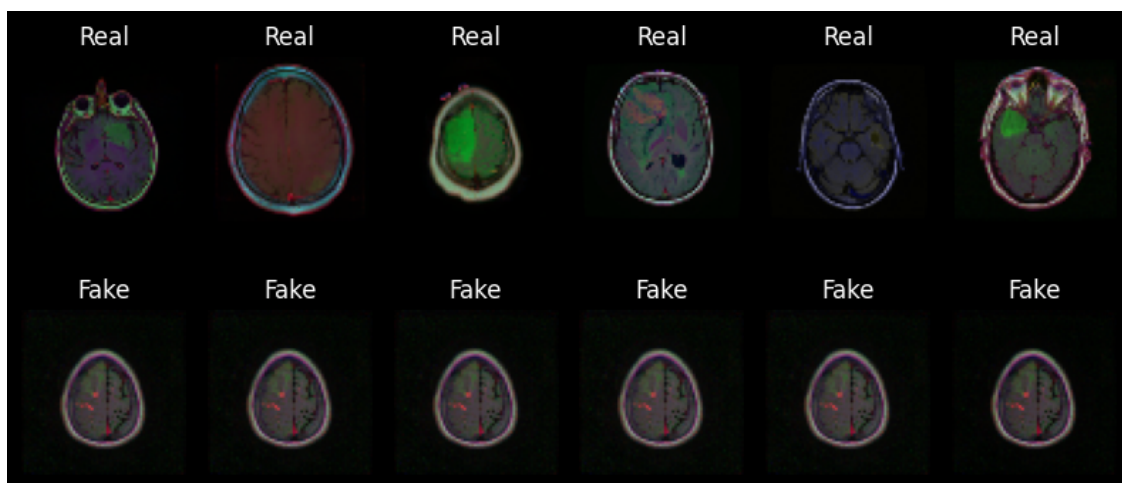
```
[24]: test_images, test_masks, test_labels = next(iter(test_dataloader))
```

1 Model evaluation

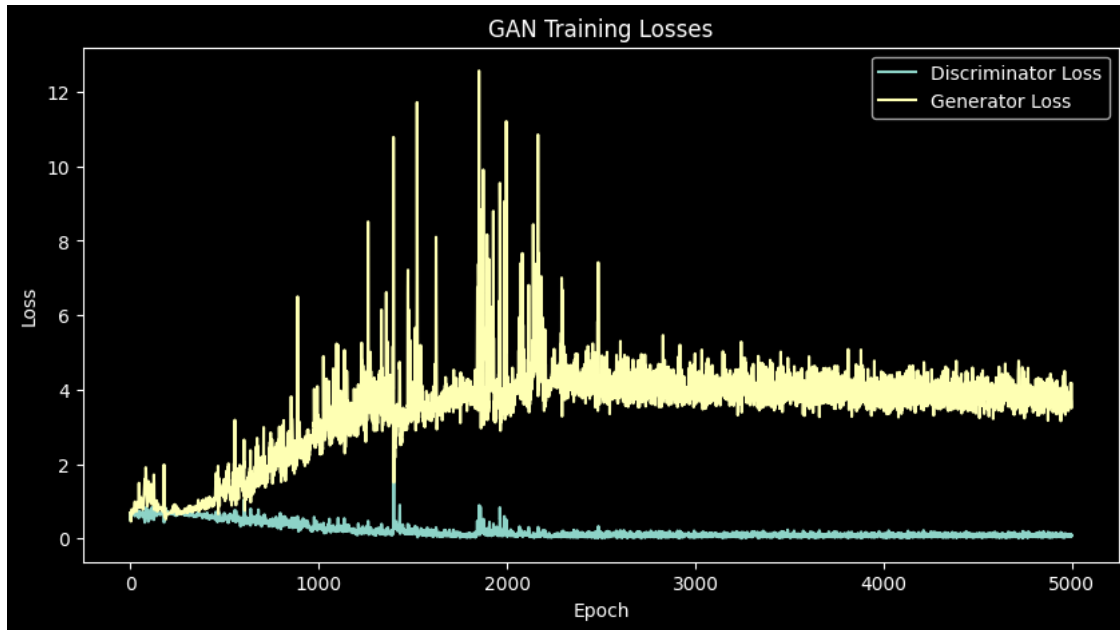
```
[25]: gan.plot_real_vs_fake(test_images, 1000, 4)
```



```
[26]: gan.plot_real_vs_fake(test_images, 1000, 6)
```



```
[27]: gan.plot_losses()
```



2 Brain Cancer Classifier

The fundamental idea underlying this classifier involves a two-step training process. Initially, the model is trained using authentic data, followed by a subsequent training round where both authentic and synthetic data are used. This approach aims to assess whether the classifier's performance exhibits improvement after incorporating synthetic data alongside genuine data.

```
[28]: from scripts.brain_cancer_classifier import BrainCancerClassifier
```

```
[29]: classifier = BrainCancerClassifier()
```

Based on real images

```
[30]: all_data = df
```

```
[31]: # Split df into train_df and val_df
all_train_df, all_val_df = train_test_split(all_data, stratify=all_data.label,
    ↪ test_size=0.1)
all_train_df = all_train_df.reset_index(drop=True)
all_val_df = all_val_df.reset_index(drop=True)

# Split train_df into train_df and test_df
all_val_df, all_test_df = train_test_split(all_val_df, stratify=all_val_df.
    ↪ label, test_size=0.3)
all_val_df = all_val_df.reset_index(drop=True)
all_test_df = all_test_df.reset_index(drop=True)
```

```
print(f"Train: {all_train_df.shape} \nVal: {all_val_df.shape} \nTest: \n{all_test_df.shape}")
```

Train: (3536, 3)

Val: (275, 3)

Test: (118, 3)

```
[32]: IMG_SIZE = 64
      BATCH_SIZE = 26

      # train
      all_train_dataset = BrainMriDataset(df=all_train_df, img_size=IMG_SIZE)
      all_train_dataloader = DataLoader(all_train_dataset, batch_size=BATCH_SIZE,
      ↪num_workers=4, shuffle=True)

      # val
      all_val_dataset = BrainMriDataset(df=all_val_df, img_size=IMG_SIZE)
      all_val_dataloader = DataLoader(all_val_dataset, batch_size=BATCH_SIZE,
      ↪num_workers=4, shuffle=True)

      #test
      all_test_dataset = BrainMriDataset(df=all_test_df, img_size=IMG_SIZE)
      all_test_dataloader = DataLoader(all_test_dataset, batch_size=BATCH_SIZE,
      ↪num_workers=4, shuffle=True)
```

```
[33]: all_train_images, all_train_masks, all_train_labels =
      ↪next(iter(all_train_dataloader))
```

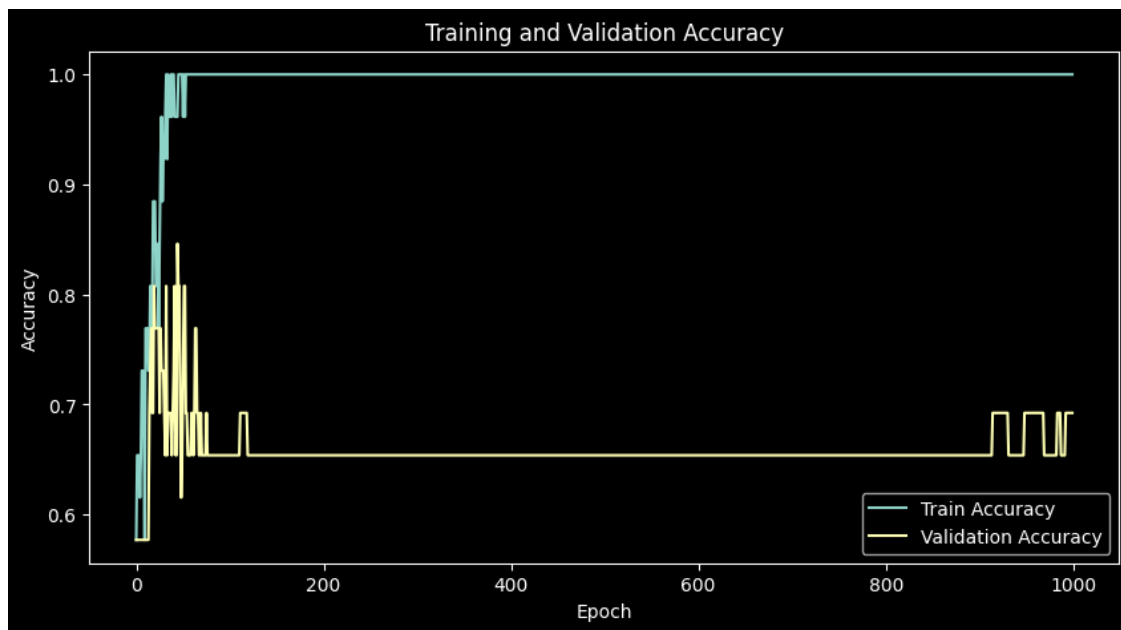
```
[34]: all_val_images, all_val_masks, all_val_labels = next(iter(all_val_dataloader))
```

```
[35]: classifier.train(all_train_images, all_train_labels, all_val_images,
      ↪all_val_labels)
```

Model saved.

```
[36]: all_test_images, all_test_masks, all_test_labels =
      ↪next(iter(all_test_dataloader))
```

```
[37]: classifier.plot_training_history()
```



```
[38]: classifier.evaluate(all_test_images, all_test_labels)
```

```
1/1 [=====] - 0s 66ms/step - loss: 3.9830 - accuracy: 0.6538
```

```
Test Loss: 3.982983350753784
```

```
Test Accuracy: 0.6538462042808533
```

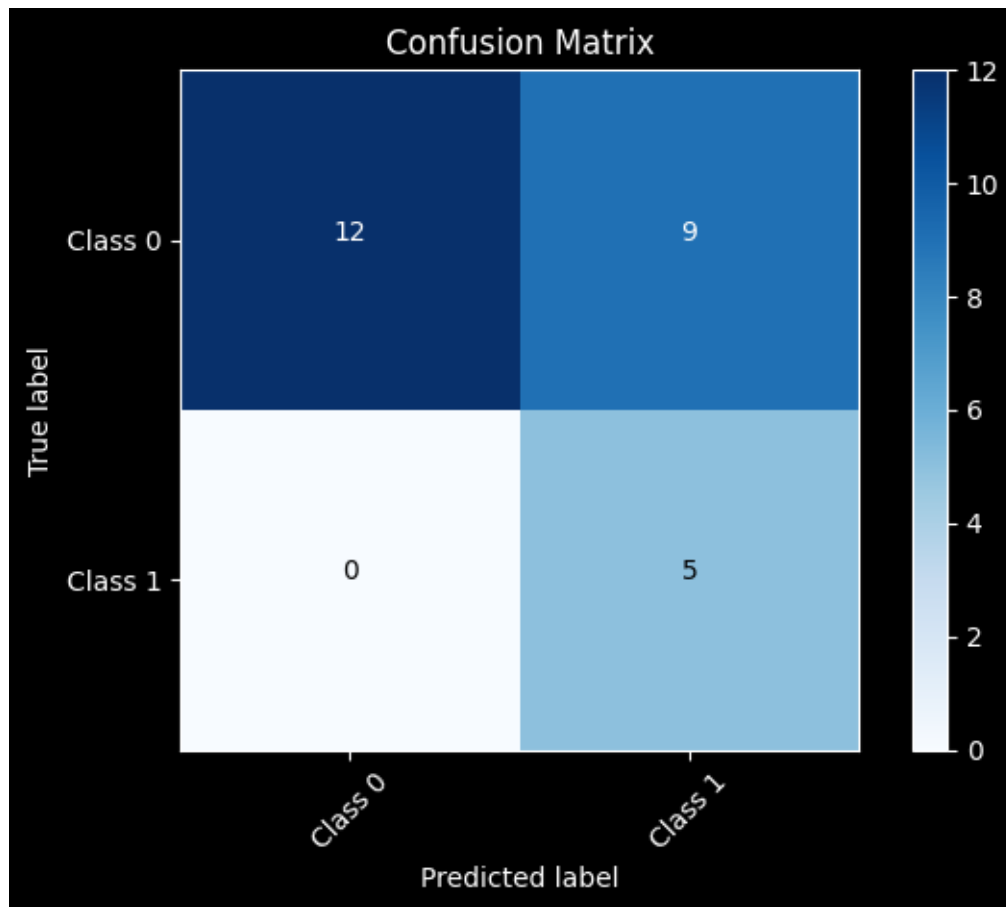
```
1/1 [=====] - 0s 51ms/step
```

```
Confusion Matrix:
```

```
[[12  9]
 [ 0  5]]
```

```
Classification Report:
```

	precision	recall	f1-score	support
0	1.00	0.57	0.73	21
1	0.36	1.00	0.53	5
accuracy			0.65	26
macro avg	0.68	0.79	0.63	26
weighted avg	0.88	0.65	0.69	26



```
[39]: from scripts.brain_mri_dataset import CombinedBrainMriDataset
```

```
[40]: IMG_SIZE = 64
      BATCH_SIZE = 26

      # Instantiate the CombinedBrainMriDataset
      combined_dataset = CombinedBrainMriDataset(original_df=all_train_df,
      ↪ generated_images=generated_images, img_size=IMG_SIZE)

      # Create a DataLoader for the combined dataset
      combined_dataloader = DataLoader(combined_dataset, batch_size=BATCH_SIZE,
      ↪ shuffle=True)
```

Based on fake images

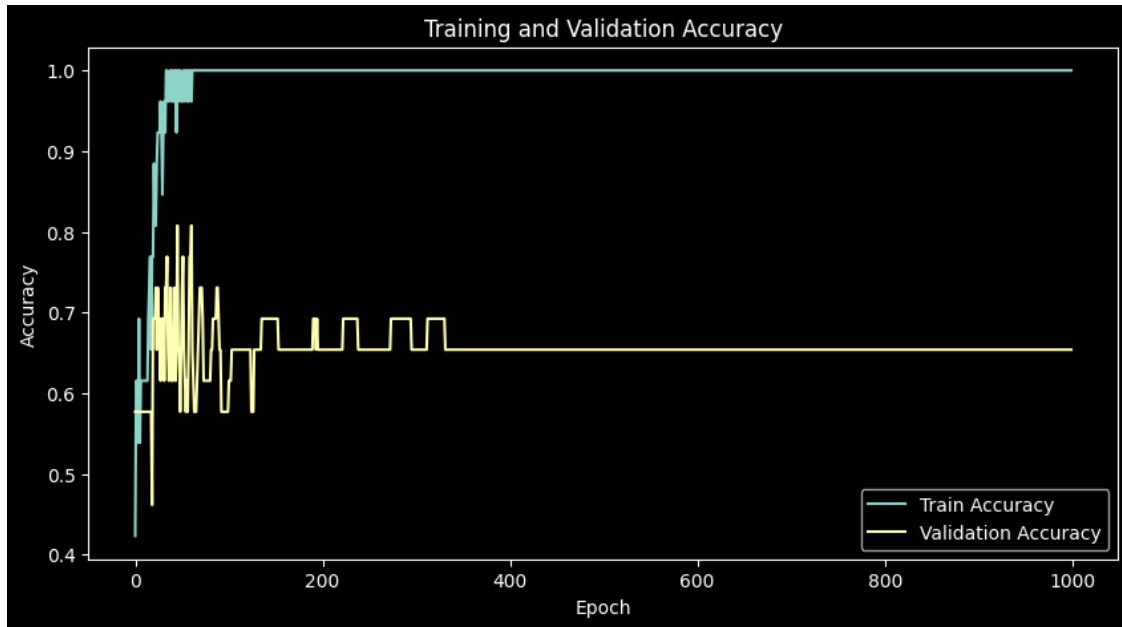
```
[41]: balanced_train_images, balanced_train_masks, balanced_train_labels =
      ↪ next(iter(combined_dataloader))
```

```
[42]: classifier_with_fake_data = BrainCancerClassifier()
```

```
[43]: classifier_with_fake_data.train(balanced_train_images, balanced_train_labels,
    ↪all_val_images, all_val_labels)
```

Model saved.

```
[44]: classifier_with_fake_data.plot_training_history()
```



```
[45]: classifier_with_fake_data.evaluate(all_test_images, all_test_labels)
```

```
1/1 [=====] - 0s 20ms/step - loss: 4.0785 - accuracy: 0.6923
```

```
Test Loss: 4.078546524047852
```

```
Test Accuracy: 0.692307710647583
```

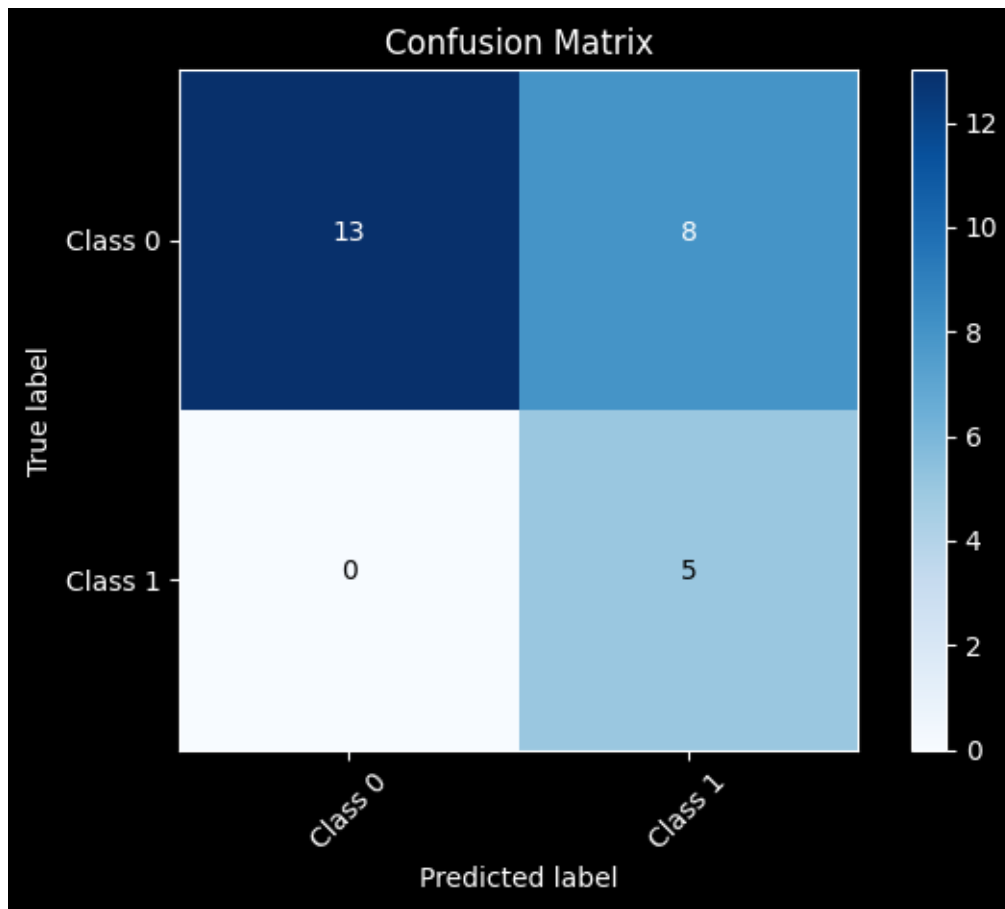
```
1/1 [=====] - 0s 52ms/step
```

```
Confusion Matrix:
```

```
[[13  8]
 [ 0  5]]
```

```
Classification Report:
```

	precision	recall	f1-score	support
0	1.00	0.62	0.76	21
1	0.38	1.00	0.56	5
accuracy			0.69	26
macro avg	0.69	0.81	0.66	26
weighted avg	0.88	0.69	0.72	26



[]: