

NeuroScope: CycleGAN Domain Adaptation for Multi-Site Brain MRI Harmonization

(A) Model Architecture

ARCHITECTURE

Generator:

- 6 Residual Blocks
- 64 base filters ($\rightarrow 256$)
- Instance Normalization
- Reflection Padding
- Tanh output

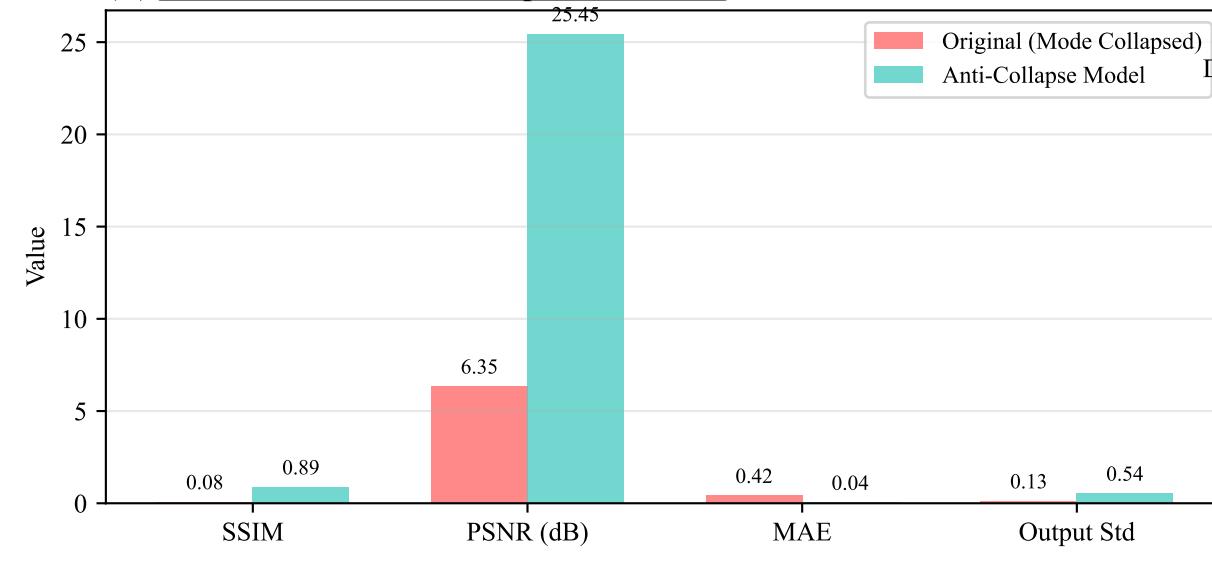
Discriminator:

- PatchGAN (70x70)
- Spectral Normalization
- LeakyReLU (0.2)

Input/Output:

- 4 channels (T1, T1-Gd, T2, FLAIR)
- 256 x 256 resolution

(C) Performance Metrics Comparison



(B) Training Configuration

TRAINING CONFIGURATION

Anti-Mode-Collapse:

- TTUR: $G_lr=2e-4$, $D_lr=1e-4$
- Label Smoothing (0.9/0.1)
- Spectral Normalization
- Replay Buffer (50 samples)
- Gradient Clipping (1.0)

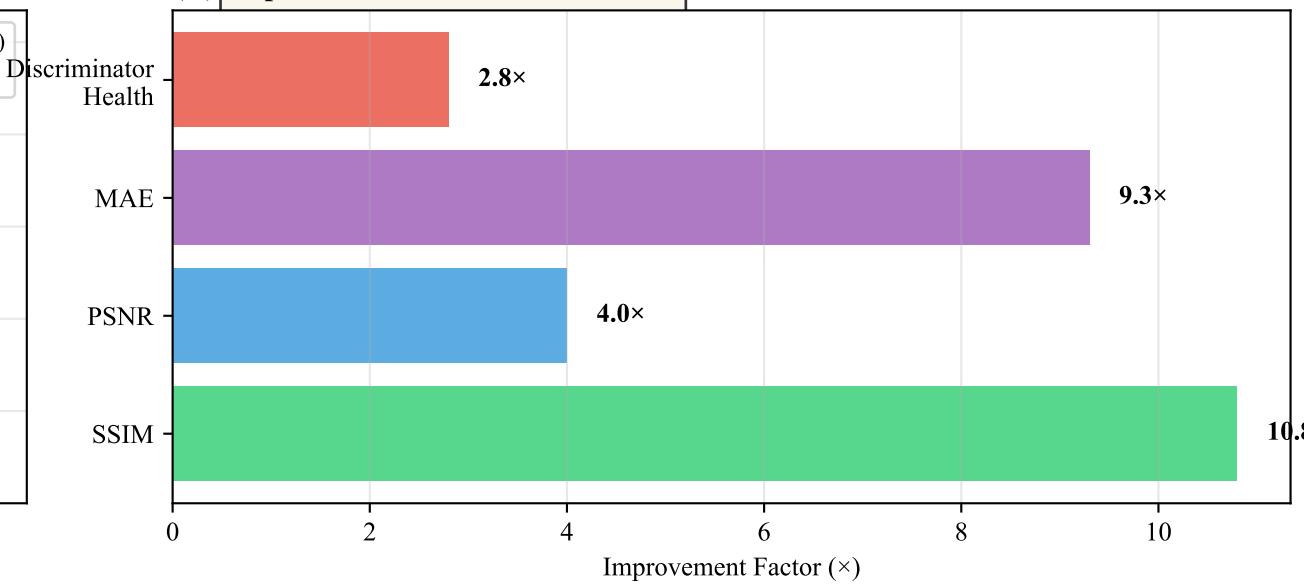
Loss Weights:

- $\lambda_{cycle} = 10.0$
- $\lambda_{identity} = 5.0$
- $\lambda_{GAN} = 1.0$

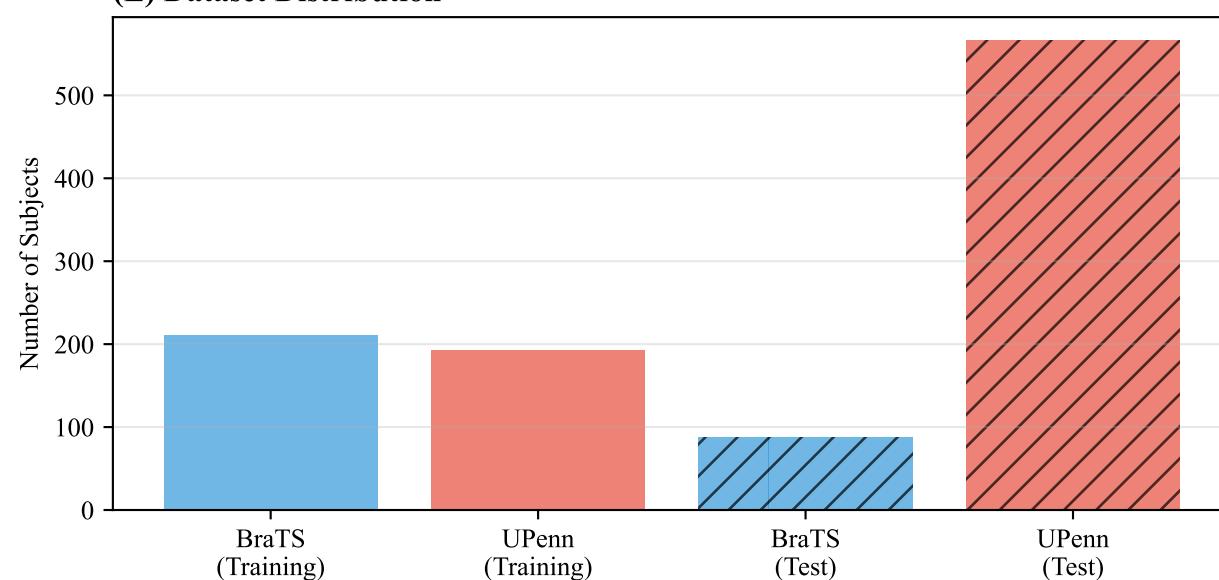
Training:

- 30 epochs
- Batch size: 4

(D) Improvement Over Baseline



(E) Dataset Distribution



(F) Results Summary

FINAL RESULTS SUMMARY

Evaluation on 654 test subjects:

Cycle Consistency:

- SSIM: 0.886 ± 0.016
- PSNR: 25.45 ± 1.37 dB
- MAE: 0.0448 ± 0.0098

Mode Collapse: ✓ RESOLVED

- Output diversity: 0.541 (healthy)
- $D_A: 0.134$, $D_B: 0.142$ (balanced)

Improvement over original:

- SSIM: 10.8 \times better
- PSNR: +19.1 dB
- Training: 40% faster (30 vs 50 epochs)