

## GAN-BASED SYNTHETIC BRAIN MR IMAGE GENERATION

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### ABSTRACT

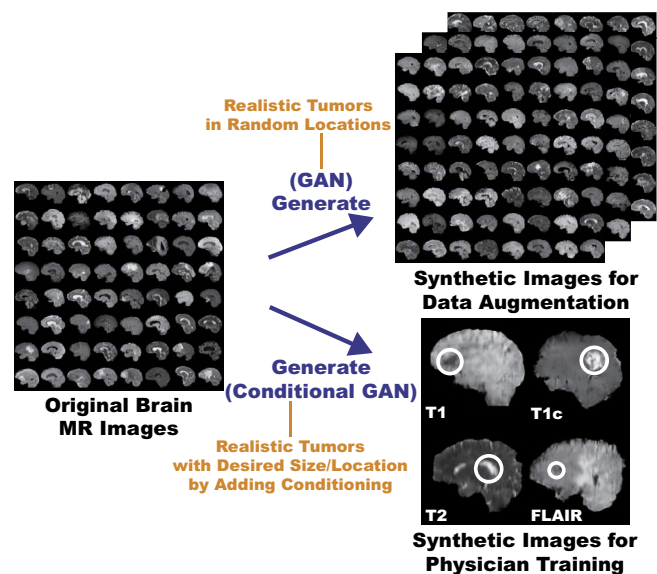
In medical imaging, it remains a challenging and valuable goal how to generate realistic medical images completely different from the original ones; the obtained synthetic images would improve diagnostic reliability, allowing for data augmentation in computer-assisted diagnosis as well as physician training. In this paper, we focus on generating synthetic multi-sequence brain Magnetic Resonance (MR) images using Generative Adversarial Networks (GANs). This involves difficulties mainly due to low contrast MR images, strong consistency in brain anatomy, and intra-sequence variability. Our novel realistic medical image generation approach shows that GANs can generate  $128 \times 128$  brain MR images avoiding artifacts. In our preliminary validation, even an expert physician was unable to accurately distinguish the synthetic images from the real samples in the Visual Turing Test.

**Index Terms**— Generative Adversarial Networks, Synthetic Medical Image Generation, Brain MRI, Data Augmentation, Physician Training, Visual Turing Test

### 1. INTRODUCTION

Along with classic methods [1], Convolutional Neural Networks (CNNs) have recently revolutionized medical image analysis [2], including brain Magnetic Resonance Imaging (MRI) segmentation [3]. However, CNN training demands extensive medical data that are laborious to obtain [4]. To overcome this issue, data augmentation techniques via reconstructing original images are common for better performance, such as geometry and intensity transformations [5, 6].

However, those reconstructed images intrinsically resemble the original ones, leading to limited performance improvement in terms of generalization abilities; thus, generating re-



**Fig. 1.** Potential applications of the proposed GAN-based synthetic brain MR image generation: (1) data augmentation for better diagnostic accuracy by generating random realistic images giving insights in classification; (2) physician training for better understanding various diseases to prevent misdiagnosis by generating desired realistic pathological images.

alistic (similar to the real image distribution) but completely new images is essential. In this context, Generative Adversarial Network (GAN)-based data augmentation has excellently performed in general computer vision tasks. It attributes to GAN's good generalization ability from matching the generated distribution from noise variables to the real one with a sharp value function. Especially, Shrivastava *et al.* (SimGAN) outperformed the state-of-the-art with a relative 21% improvement in eye-gaze estimation [7].

So, how can we generate realistic medical images completely different from the original samples? Our aim is to gen-

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erate synthetic multi-sequence brain MR images using GANs, which is essential in medical imaging to increase diagnostic reliability, such as via data augmentation in computer-assisted diagnosis as well as physician training and teaching (Fig. 1) [8]. However, this is extremely challenging—MR images are characterized by low contrast, strong visual consistency in brain anatomy, and intra-sequence variability. Our novel GAN-based approach for medical data augmentation adopts Deep Convolutional GAN (DCGAN) [9] and Wasserstein GAN (WGAN) [10] to generate realistic images, and an expert physician validates them via the Visual Turing Test [11].

**Research Questions.** We mainly address two questions:

- **GAN Selection:** Which GAN architecture is well-suited for realistic medical image generation?
- **Medical Data Augmentation:** How can we handle MR images with specific intra-sequence variability?

**Contributions.** Our main contributions are as follows:

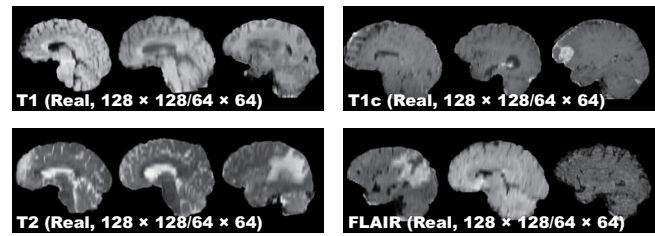
- **MR Image Generation:** This research shows that WGAN can generate realistic multi-sequence brain MR images, possibly leading to valuable clinical applications: data augmentation and physician training.
- **Medical Image Generation:** This research provides how to exploit medical images with intrinsic intra-sequence variability towards GAN-based data augmentation for medical imaging.

## 2. GENERATIVE ADVERSARIAL NETWORKS

Since the breakthrough paper by Goodfellow *et al.* in 2014 [12], GANs have shown promising results for image generation in general computer vision [13]. GANs generate highly realistic images, without a well-defined objective function associated with difficult training accompanying oscillations and mode collapse—i.e., a common failure case where the generator learns with extremely low variety. Whereas Variational Autoencoders (VAEs) [14], the other most used deep generative models, have an objective likelihood function to optimize, and could so generate blurred samples because of the injected noise and imperfect reconstruction [15].

Therefore, many medical imaging researchers have begun to use GANs recently, such as in image super-resolution [16], anomaly detection [17], and estimating CT images from the corresponding MR images [18]. As GANs allow adding conditioning on the class labels and images, they often use such conditional GANs to produce desired images, while it makes learning robust latent spaces difficult.

Differently from a very recent work of GANs for biological image synthesis (fluorescence microscopy) [19], to the best of our knowledge, this is the first GAN-based realistic brain tumor MR image generation approach aimed at data augmentation and physician training. Instead of reconstructing real brain MR images themselves with respect to geometry/intensity, a completely different approach—generating



**Fig. 2.** Example real MR images used for training the GANs: the resized sagittal multi-sequence brain MRI scans of patients with HGG on the BRATS 2016 training dataset [20].

novel realistic images using GANs—may become a clinical breakthrough.

## 3. MATERIALS AND METHODS

Towards clinical applications utilizing realistic brain MR images, we generate synthetic brain MR images from the original samples using GANs. Here, we compare the most used two GANs, DCGAN and WGAN, to find a well-suited GAN between them for medical image generation—it must avoid mode collapse and generate realistic MR images with high resolution.

### 3.1. The BRATS 2016 Dataset

This paper exploits a dataset of multi-sequence brain MR images to train GANs with sufficient data and resolution, which was originally produced for the Multimodal Brain Tumor Image Segmentation Benchmark (BRATS) challenge [20]. In particular, the BRATS 2016 training dataset contains 220 High-Grade Glioma (HGG) and 54 Low-Grade Glioma (LGG) cases, with T1-weighted (T1), contrast enhanced T1-weighted (T1c), T2-weighted (T2), and Fluid Attenuation Inversion Recovery (FLAIR) sequences—they were skull stripped and resampled to isotropic  $1\text{mm} \times 1\text{mm} \times 1\text{mm}$  resolution with image dimension  $240 \times 240 \times 155$ ; among the different sectional planes, we use the sagittal multi-sequence scans of patients with HGG to show that our GANs can generate a complete view of the whole brain anatomy (allowing for visual consistency among the different brain lobes), including also severe tumors for clinical purpose.

### 3.2. Proposed GAN-based Image Generation Approach

#### 3.2.1. Pre-processing

We select the slices from #80 to #149 among the whole 240 slices to omit initial/final slices, since they convey a negligible amount of useful information and could affect the training. The images are resized to both  $64 \times 64$  and  $128 \times 128$  from  $240 \times 155$  for better GAN training (DCGAN architecture results in stable training on  $64 \times 64$  [9], and so  $128 \times 128$  is reasonably a high-resolution). Fig. 2 shows some real MR images used for training; each sequence contains 15,400 images with 220 patients  $\times$  70 slices (61,600 in total).

### 3.2.2. GAN-based MR Image Generation

DCGAN and WGAN generate six types of images as follows:

- T1 sequence ( $128 \times 128$ ) from the real T1;
- T1c sequence ( $128 \times 128$ ) from the real T1c;
- T2 sequence ( $128 \times 128$ ) from the real T2;
- FLAIR sequence ( $128 \times 128$ ) from the real FLAIR;
- Concat sequence ( $128 \times 128$ ) from concatenating the real T1, T1c, T2, and FLAIR (i.e., feeding the model with samples from all the MRI sequences);
- Concat sequence ( $64 \times 64$ ) from concatenating the real T1, T1c, T2, and FLAIR.

Concat sequence refers to a new ensemble sequence for an alternative data augmentation, containing the features of all four sequences. We also generate  $64 \times 64$  Concat images to compare the generation performance in terms of image size.

**DCGAN.** DCGAN [9] is a standard GAN [12] with a convolutional architecture for unsupervised learning; this generative model uses up-convolutions interleaved with ReLU non-linearity and batch-normalization.

Let  $p_{\text{data}}$  be a generating distribution over data  $\mathbf{x}$ . The generator  $G(\mathbf{z}; \theta_g)$  is a mapping to data space that takes a prior on input noise variables  $p_z(\mathbf{z})$ , where  $G$  is a neural network with parameters  $\theta_g$ . Similarly, the discriminator  $D(\mathbf{x}; \theta_d)$  is a neural network with parameters  $\theta_d$  that takes either real data or synthetic data and outputs a single scalar probability that  $\mathbf{x}$  came from the real data. The discriminator  $D$  maximizes the probability of classifying both training examples and samples from  $G$  correctly while the generator  $G$  minimizes the likelihood; it is formulated as a minimax two-player game with value function  $V(G, D)$ :

$$\min_G \max_D V(D, G) = \mathbb{E}_{\mathbf{x} \sim p_{\text{data}}(\mathbf{x})} [\log D(\mathbf{x})] + \mathbb{E}_{\mathbf{z} \sim p_z(\mathbf{z})} [\log(1 - D(G(\mathbf{z})))] \quad (1)$$

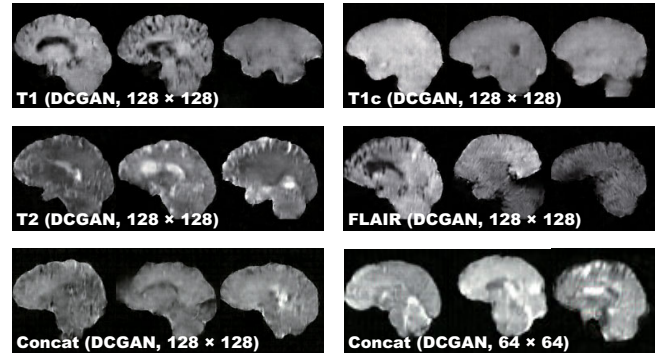
This can be reformulated as the minimization of the Jensen-Shannon (JS) divergence between the distribution  $p_{\text{data}}$  and another distribution  $p_g$  derived from  $p_z$  and  $G$ .

**DCGAN Implementation Details.** We use the same DCGAN architecture [9] with no tanh in the generator, ELU as the discriminator, all filters of size  $4 \times 4$ , and a half channel size for DCGAN training. A batch size of 64 and Adam optimizer with  $2.0 \times 10^{-4}$  learning rate were implemented.

**WGAN.** WGAN [10] is an alternative to traditional GAN training, as the JS divergence is limited, such as when it is discontinuous; this novel GAN achieves stable learning with less mode collapse by replacing it to the Earth Mover (EM) distance (a.k.a. the Wasserstein-1 metrics):

$$W(p_g, p_r) = \inf_{p \in \Pi(p_g, p_r)} \mathbb{E}_{(\mathbf{x}, \mathbf{x}') \sim p} \|\mathbf{x} - \mathbf{x}'\|, \quad (2)$$

where  $\Pi(p_g, p_r)$  is the set of all joint distributions  $p$  whose marginals are  $p_g$  and  $p_r$ , respectively. In other words,  $p$



**Fig. 3.** Example synthetic MR images yielded by DCGAN.

implies how much mass must be transported from one distribution to another. This distance intuitively indicates the cost of the optimal transport plan.

**WGAN Implementation Details.** We use the same DCGAN architecture [9] for WGAN training. A batch size of 64 and Root Mean Square Propagation (RMSprop) optimizer with  $5.0 \times 10^{-5}$  learning rate were implemented.

### 3.3. Clinical Validation Using the Visual Turing Test

To quantitatively evaluate how realistic the synthetic images are, an expert physician was asked to constantly classify a random selection of 50 real/50 synthetic MR images as real or synthetic shown in a random order for each GAN/sequence, without previous training stages revealing which is real/synthetic; Concat images were classified together with real T1, T1c, T2, and FLAIR images in equal proportion. The so-called Visual Turing Test [11] uses binary questions to probe a human ability to identify attributes and relationships in images. For these motivations, it is commonly used to evaluate GAN-generated images, such as for SimGAN [7]. This applies also to medical images in clinical environments [21], wherein physicians' expertise is critical.

## 4. RESULTS

This section shows how DCGAN and WGAN generate synthetic brain MR images. The results include instances of synthetic images and their quantitative evaluation of the realism by an expert physician. The training took about 2 (1) hours to train each  $128 \times 128$  ( $64 \times 64$ ) sequence on an Nvidia GeForce GTX 980 GPU, increasingly learning realistic features.

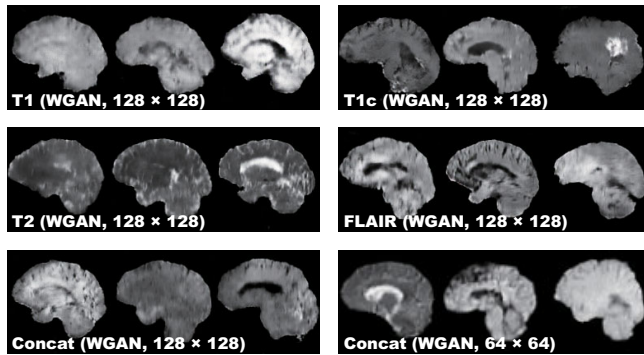
### 4.1. MR Images Generated by GANs

**DCGAN.** Fig. 3 illustrates examples of synthetic images by DCGAN. The images look similar to the real samples. Concat images combine appearances and patterns from all the four sequences used in training. Since DCGAN's value function could be unstable, it often generates hyper-intense T1-like images analogous to mode collapse for  $64 \times 64$  Concat images, while sharing the same hyper-parameters with  $128 \times 128$ .



**Table 1.** Visual Turing Test results by a physician for classifying real vs synthetic images. It should be noted that proximity to 50% of accuracy indicates superior performance (chance = 50%).

	Accuracy (%)	Real Selected as Real	Real as Synt	Synt as Real	Synt as Synt
T1 (DCGAN, $128 \times 128$ )	70	26	24	6	44
T1c (DCGAN, $128 \times 128$ )	71	24	26	3	47
T2 (DCGAN, $128 \times 128$ )	64	22	28	8	42
FLAIR (DCGAN, $128 \times 128$ )	54	12	38	8	42
Concat (DCGAN, $128 \times 128$ )	77	34	16	7	43
Concat (DCGAN, $64 \times 64$ )	54	13	37	9	41
T1 (WGAN, $128 \times 128$ )	64	20	30	6	44
T1c (WGAN, $128 \times 128$ )	55	13	37	8	42
T2 (WGAN, $128 \times 128$ )	58	19	31	11	39
FLAIR (WGAN, $128 \times 128$ )	62	16	34	4	46
Concat (WGAN, $128 \times 128$ )	66	31	19	15	35
Concat (WGAN, $64 \times 64$ )	53	18	32	15	35



**Fig. 4.** Example synthetic MR images yielded by WGAN.

**WGAN.** Fig. 4 shows the example output of WGAN in each sequence. Outperforming remarkably DCGAN, WGAN successfully captures the sequence-specific texture and the appearance of the tumors while maintaining the realism of the original brain MR images. As expected,  $128 \times 128$  Concat images tend to have more messy and unrealistic artifacts than  $64 \times 64$  Concat ones, especially around the boundaries of the brain, due to the introduction of unexpected intensity patterns.

#### 4.2. Visual Turing Test Results

Table 1 shows the confusion matrix concerning the Visual Turing Test. Even the expert physician found classifying real and synthetic images challenging, especially in lower resolution due to their less detailed appearances unfamiliar in clinical routine, even for highly hyper-intense  $64 \times 64$  Concat images by DCGAN; distinguishing Concat images was easier compared to the case of T1, T1c, T2, and FLAIR images because the physician often felt odd from the artificial sequence. WGAN succeeded to deceive the physician significantly better than DCGAN for all the MRI sequences except FLAIR images (62% to 54%).

#### 5. CONCLUSION

Our preliminary results show that GANs, especially WGAN, can generate  $128 \times 128$  realistic multi-sequence brain MR images that even an expert physician is unable to accurately distinguish from the real, leading to valuable clinical applications, such as data augmentation and physician training. This attributes to WGAN's good generalization ability with a sharp value function. In this context, DCGAN might be unsuitable due to both the inferior realism and mode collapse in terms of intensity. We only use the slices of interest in training to obtain desired MR images and generate both original/Concat sequence images for data augmentation in medical imaging.

This study confirms the synthetic image quality by the human expert evaluation, but a more objective computational evaluation for GANs should also follow, such as Classifier Two-Sample Tests (C2ST) [22], which assesses whether two samples are drawn from the same distribution. Currently this work uses sagittal MR images alone, so we will generate coronal and transverse images in the near future. As this research uniformly selects middle slices in pre-processing, better data generation demands developing a classifier to only select brain MRI slices with/without tumors.

Towards data augmentation, while realistic images give more insights on geometry/intensity transformations in classification, more realistic images do not always assure better data augmentation, so we have to find suitable image resolutions and sequences; that is why we generate both high-resolution images and Concat images, yet they looked more unrealistic for the physician. For physician training, generating desired realistic tumors by adding conditioning requires exploring extensively the latent spaces of GANs.

Overall, our novel GAN-based realistic brain MR image generation approach sheds light on diagnostic and prognostic medical applications; future studies on these applications are needed to confirm our encouraging results.

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