

Adversarial Correction Networks for Brain MRI Segmentation in 2D Manner

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1 Introduction

The development of deep learning techniques has largely improved the state-of-the-art segmentation methods [2, 3, 8]. Recently, fully convolutional networks (FCN) [3] provides a great choice for semantic image segmentation [2–4, 6–9]. Thus, we propose a FCN-like network to automatically segment the brain MRI. Due to a lot of issues in medical image field (small data set size, 3D format and so on), we propose several mechanisms (such as using a transmission module, residual learning module and intermedia-supervision modules) to solve the training problems of the network. Most importantly, we provide an adversarial correction network [1, 5] to evaluate the quality of segmentation network and thus we can improve the training for difficult cases. Dropout is also involved to relieve overfitting. We use generalized dice loss hybridized by softmax loss as the objective function.

This work is done in a 2D manner due to time limit. Personally, it is worth to run the 3D version if time allowed.

2 Experiments

We use MRBrain2018 dataset (see the description in the below paragraph) for training in a leave-one-out manner. And the patch size we extracted for training is 3x192x192. We are planning to evaluate our proposed approach on the test dataset of brain MRI segmentation.

The results on training validation likes (show case 7):

dice1= 0.893537644759 dice2= 0.965670723041 dice3 = 0.914421178163 dice4 = 0.99999997917 dice5 = 0.881257114503 dice6 = 0.983529680882 dice7 = 0.93848676381 dice8 = 0.99472058969 dice9 = 0.0 dice10 = 0.0

And the results on validation validation (show case 1):

dice1 = 0.846024607653 dice2 = 0.837783064583 dice3 = 0.890493967206 dice4 = 0.68446979516 dice5 = 0.827850313027 dice6 = 0.942282799052 dice7 = 0.904149703592 dice8 = 0.754513481823 dice9 = 0.0 dice10 = 0.0

“Image data used in this challenge were acquired on a 3T scanner at the UMC Utrecht (the Netherlands). For each of the 30 subjects, fully annotated multi-sequence (T1-weighted, T1-weighted inversion recovery and T2-FLAIR) scans are available. The 30 subjects include patients with diabetes, dementia and Alzheimers, and matched controls (with increased cardiovascular risk) with varying degrees of atrophy and white matter lesions (age > 50).”

References

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