

Clustering-based Unsupervised Brain MRI Segmentation

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INTRODUCTION

Background

- Semantic and lesion/tumor segmentation of brain magnetic resonance imaging (MRI) has many uses in research and clinical applications.
- Annotation of brain MRI is traditionally done by radiologists with specialized skills.
- Supervised methods are not applicable due to difficulty in collecting large, high-quality labels.

Previous Work

- Generative models trained purely on healthy brain images (AnoVAEGAN). [1]
- K-means with feature vectors from 3D CNNs to segment lung cancer in pathology images. [2]
- Unsupervised encoder learning for natural image segmentation (IIC, MoCo). [3, 4]

Data

- Human Connectome Project (HCP) public database. [5]
- Data of 99 subjects were processed and used.

Oracle and baseline

- Oracle data comes from FreeSurfer segmentation results, labels ranging from 0 to 14175. [6]
- 114 labels that are actually presented in the 99 subjects are reclassified into 5 categories: Brain Mask, Deep Grey Matter, Grey Matter, White Matter, and Cerebrospinal fluid (CSF). [7]
- Baseline comes from the mean-shift algorithm. [8]

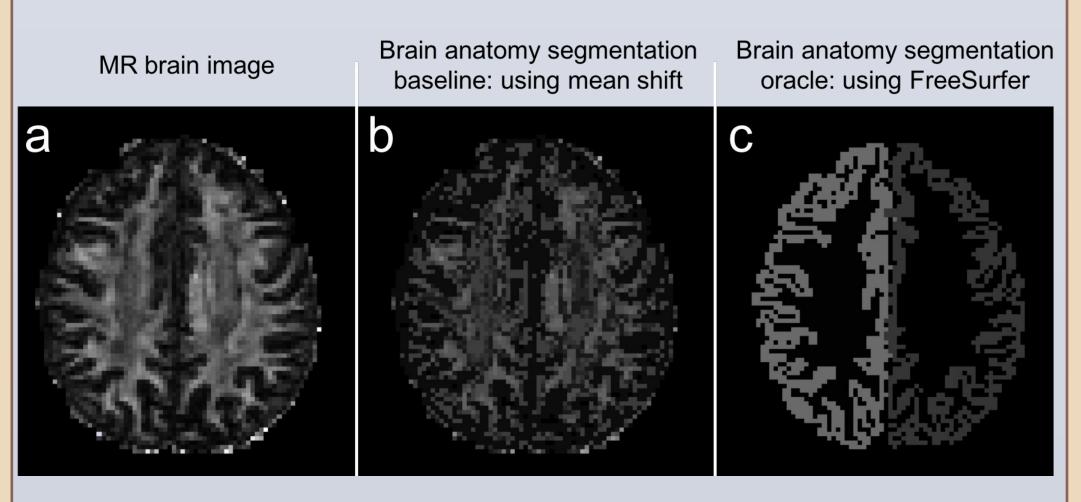


Figure 1. Example of a brain MR image (a), and segmentations results using the mean shift algorithm as baseline (b) and the FreeSurfer as oracle (c).

METHODS

Data preprocessing

- 1. Image normalization
- 2. Artifact correction
- 3. Coregistration between different contrasts
- 4. MRI physical model fitting (DTI) [9]

(64 contrasts-> 4 contrasts as below)

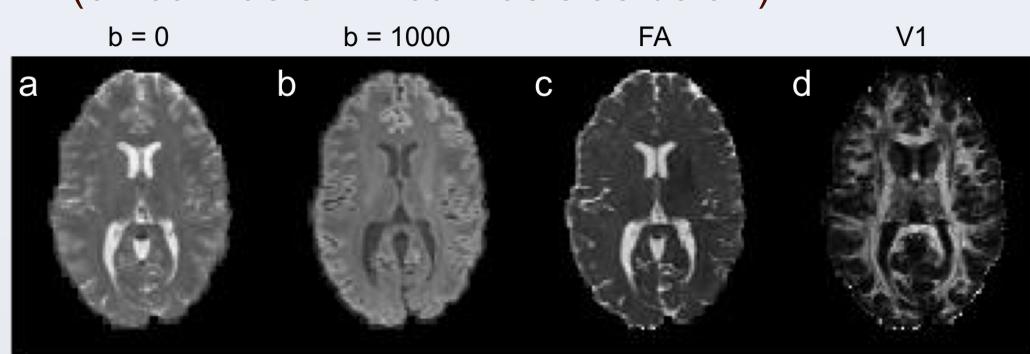
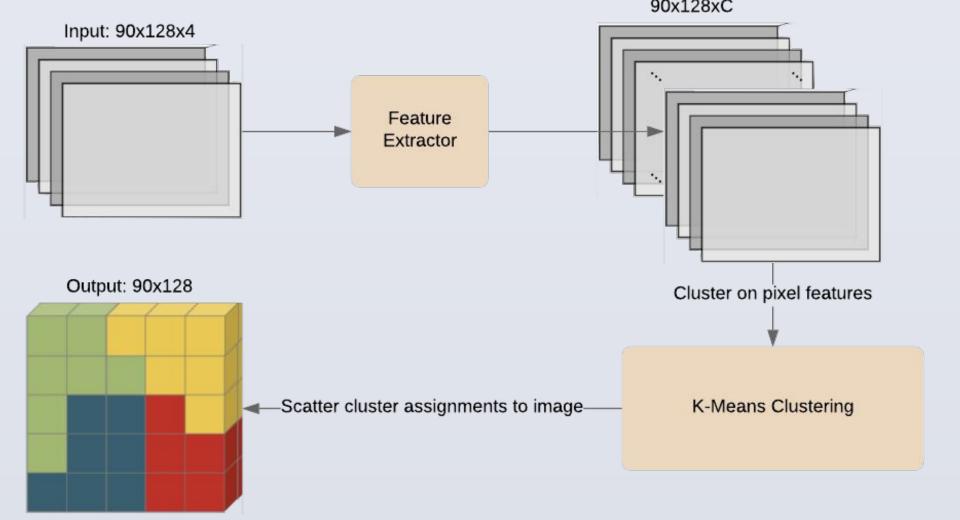
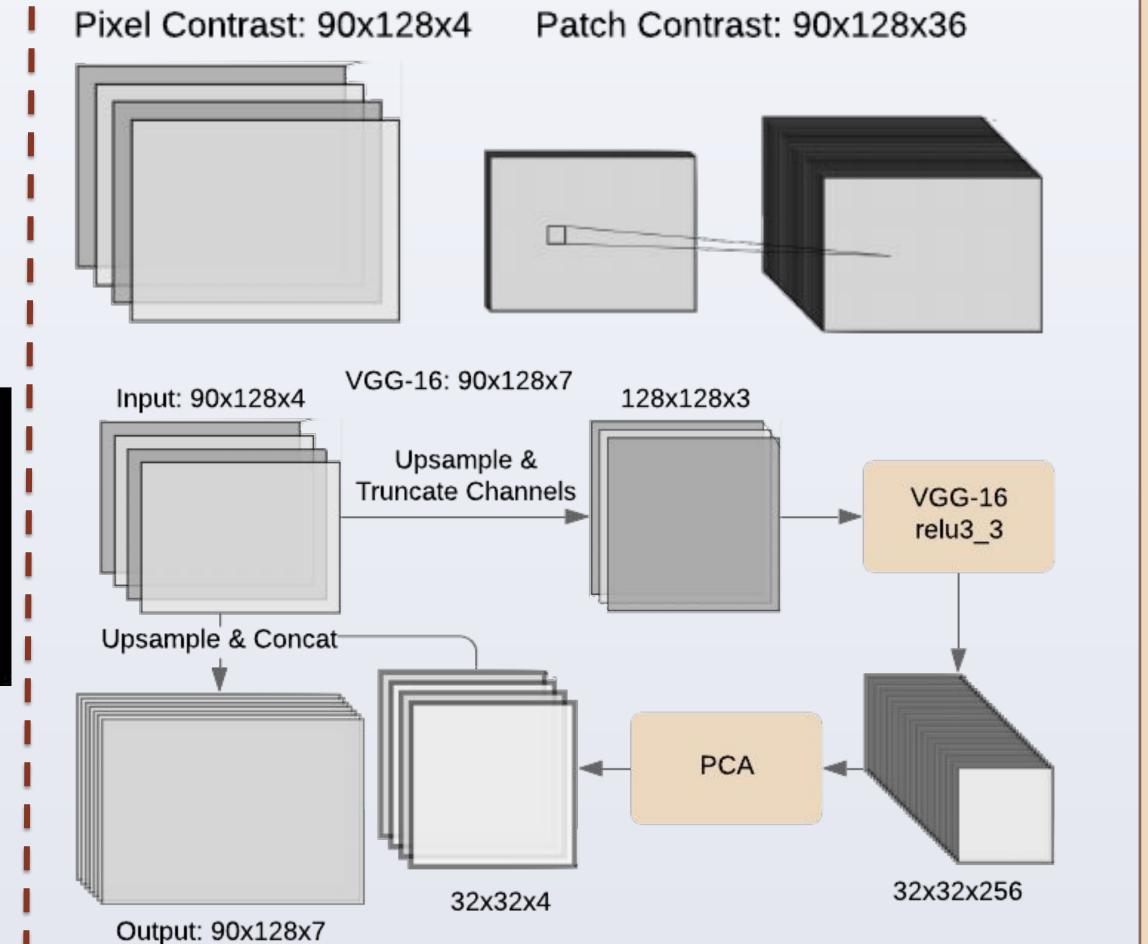


Figure 2. One representative example of preprocessed images from the HCP dataset (a: averaged b=0 image, b: averaged b=1000 image, c/d: FA /V1 maps derived from all b=1000 images).

Segmentation pipeline



Feature Extraction



Evaluation

Segmentation label matching with the Hungarian method:

- 1. Cost matrix calculation between labels
- 2. Linear assignment to find a permutation [10] Segmentation Accuracy Evaluation:
 - 1. Pixel accuracy: (TP + TN)/(TP + TN + FP + FN)
 - 2. Sørensen-Dice coefficient (Dice score)

RESULTS

Different patch sizes for feature extraction

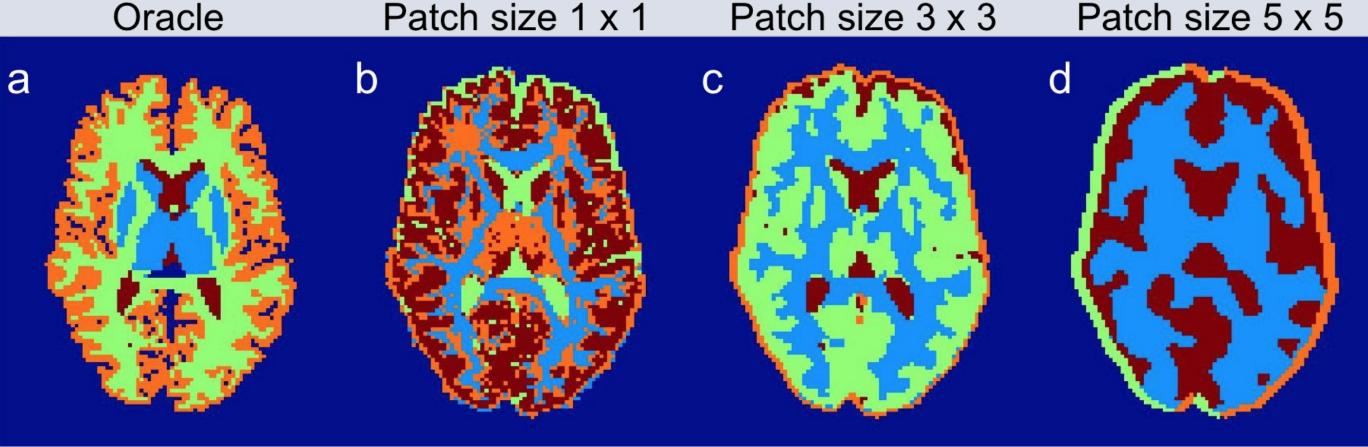
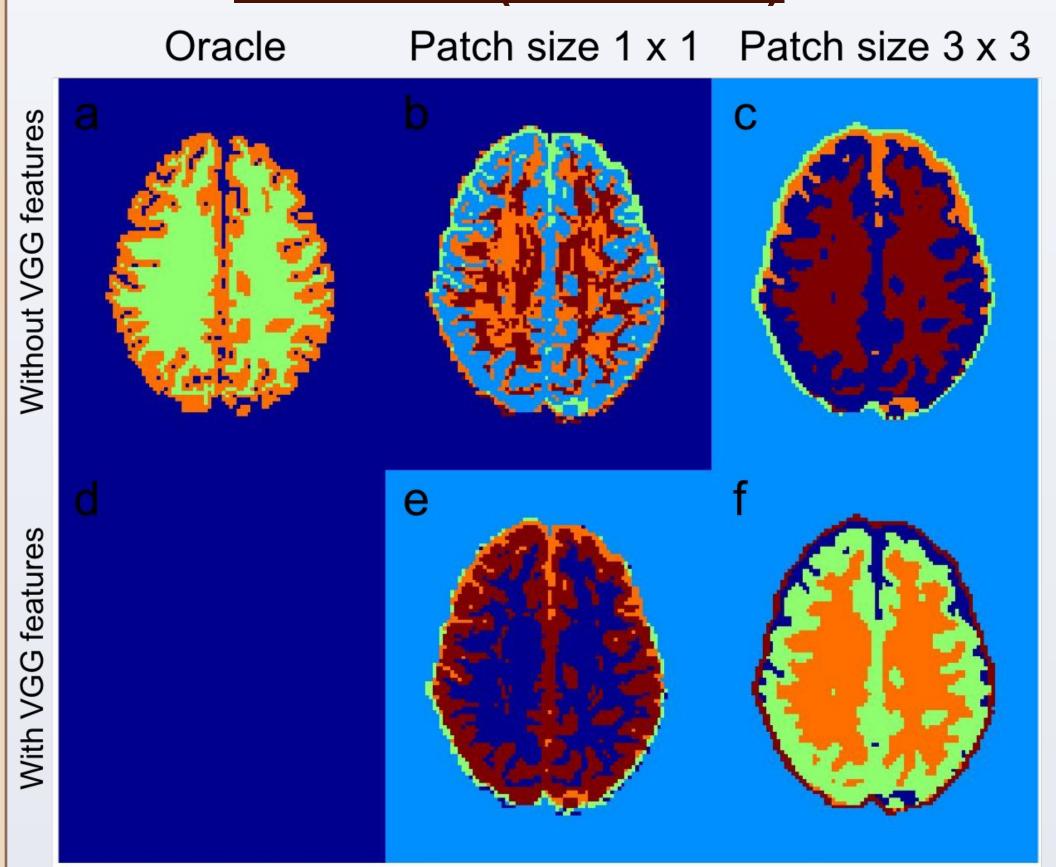


Figure 3. One representative segmentation results using FreeSurfer (a), and K-means with different patch sizes (b-d). For the choice of the patch sizes, there is a trade-off between the noise level and the details. The result is too noisy with patch size 1 (b), and too many details are lost with patch size 5 (d).

	Grey matter	White matter	Deep grey matter	CSF	Brain mask	Pixel accuracy
Patch size = 1 x 1	0.78 ± 0.09	0.79 ± 0.04	0.05 ± 0.08	0.22 ± 0.26	0.94 ± 0.01	0.84 ± 0.04
Patch size = 3 x 3	0.80 ± 0.10	0.75 ± 0.09	0.03 ± 0.06	0.20 ± 0.28	0.94 ± 0.01	0.84 ± 0.04
Patch size = 5 x 5	0.74 ± 0.15	0.67 ± 0.14	0.01 ± 0.04	0.16 ± 0.27	0.93 ± 0.01	0.81 ± 0.05
Patch size = 3 x 3 with features from pretrained VGG	0.80 ± 0.10	0.74 ± 0.11	0.02 ± 0.05	0.20 ± 0.28	0.92 ± 0.03	0.83 ± 0.05

Table 1. Dice scores of five different classes and pixel accuracy with different segmentation settings.

RESULTS (continued)



ANALYSIS

- Increasing the patch size trades off detail in segmentation in favor of being more robust against noise.
- Including features from pretrained VGG network reduces misclassification. [11]
- The proposed method works well in classifying grey and white matter, and poorly in classifying CSF and deep grey matter.

FUTURE WORK

- Explore more feature extraction methods
- Leverage other quantitative metrics for evaluation

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