

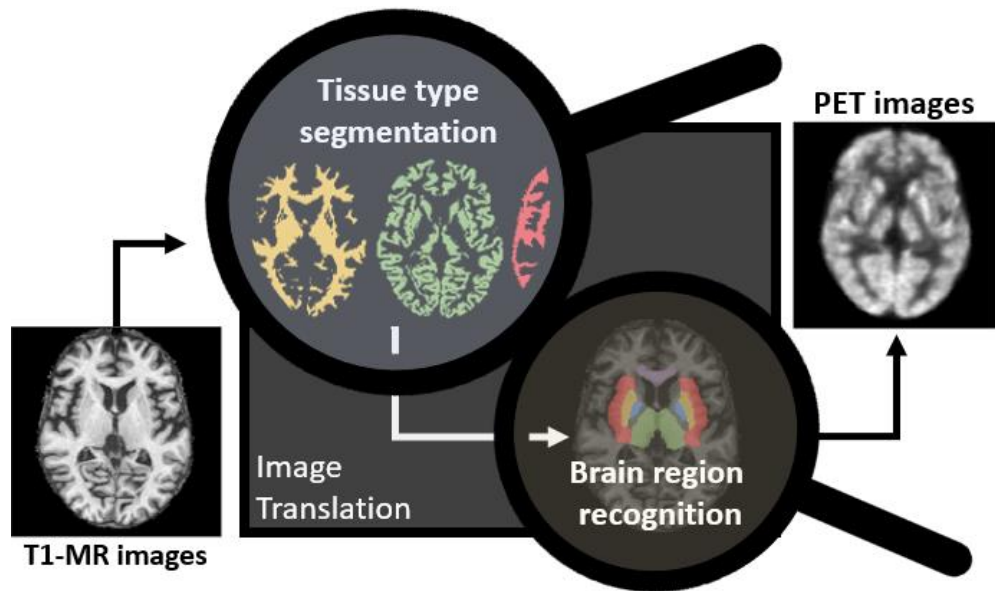
Demystifying T1-MRI to FDG¹⁸-PET Image Translation via Representational Similarity

Chia-Hsiang Kao, Yong-Sheng Chen, Li-Fen Chen, and Wei-Chen Chiu
National Yang Ming Chiao Tung University, Taiwan



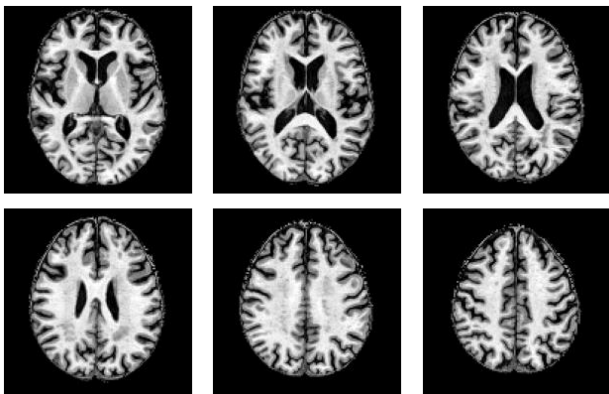
Take home message

We use representational similarity to verify our proposed hypotheses that the translation from T1-MR to PET images comprises the recognition of brain tissue types and brain regions.

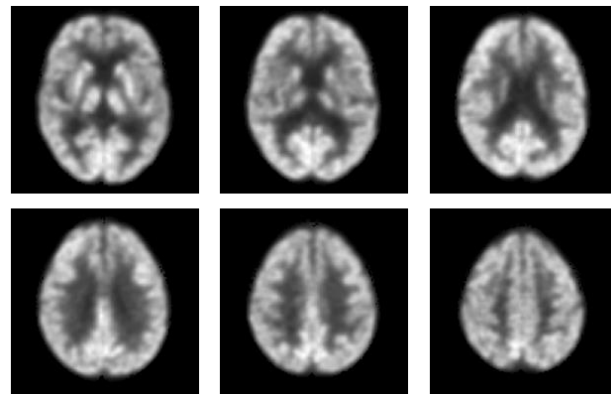


Introduction There is a growing research trend in using AI for cross-modality medical image translation.

**T1-weighted Magnetic resonance imaging
(T1-MRI)**



**FDG¹⁸-positron emission tomography
(PET)**



Principle

MRI uses strong magnetic fields to identify the anatomical structure of the brain.

PET measures the glucose uptake ability of tissue to quantify the metabolic process.

Drawback

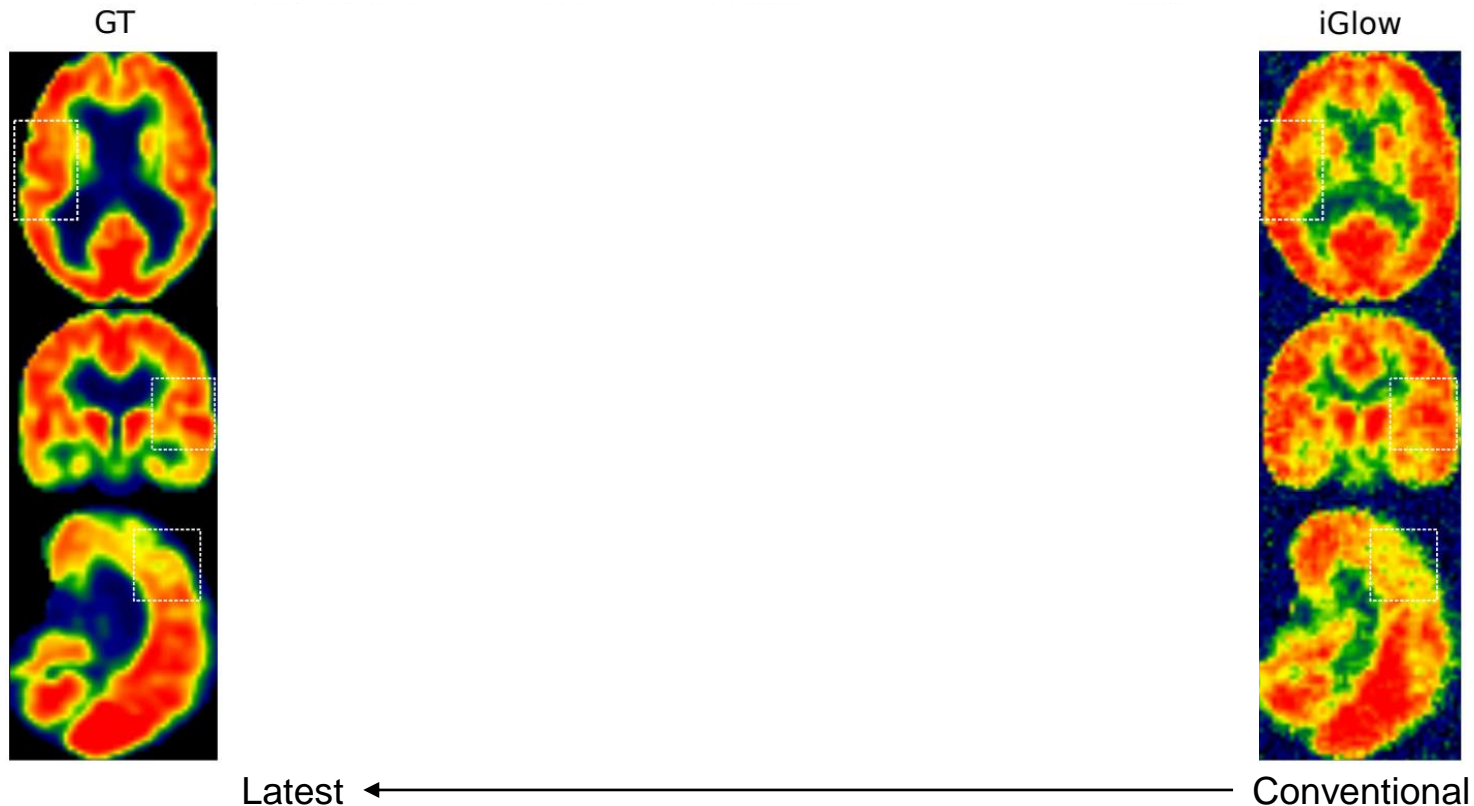
time consuming, high cost

injection of radioactive tracer, high cost

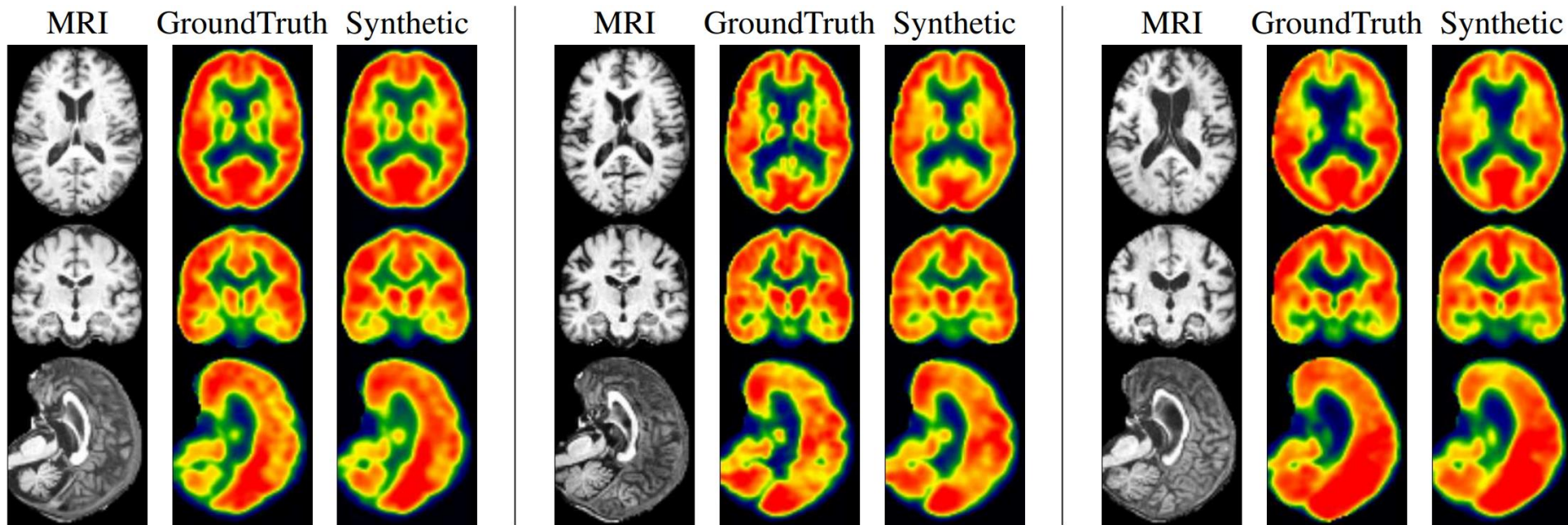
DL-based image synthesis/translation

Nie, D. (2017), Pan, Y. (2018),
Sikka, A. (2018), Sun, H. (2019), Lan, H. (2020)

Introduction There is a growing research trend in using AI for cross-modality medical image translation.



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Synthetic images are meaningful for subjects in both extremes of disease spectrum.

Left: CN. Middle: MCI. Right: AD. The generated PET images show consistency of hypometabolism (less red, more yellow) with the ground truth image.

Introduction However, there are several major concerns when it comes to clinical applications.

- Robustness to distributional biases (population, organs, diseases).



- Robustness to systemic biases (scanning devices and protocols, preprocessing techniques).



SIEMENS

PHILIPS

- Robustness to out-of-distribution cases, noise and adversarial perturbations.

Introduction Even more, deep learning models remain black-box for most AI scientists.

Analysis | [Open Access](#) | [Published: 23 August 2021](#)

Common pitfalls and mitigating bias in machine learning for medicine

[Michael Roberts](#) , [Derek Driggs](#), [Angelica I. Aviles-Rivero](#), [Christian](#), [Zhongzhao Teng](#), [Effrossyni Gkrani](#)


[Schönlieb](#)

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Peeking into a black box, the fairness and generalizability of a MIMIC-III benchmark


[Eliane Röösli](#), [Selen Bozkurt](#) & [Tina Hernandez-Boussard](#) 

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Mitigating bias in machine learning for medicine


[Kerstin N. Vokinger](#) , [Stefan Feuerriegel](#) & [Aaron S. Kesselheim](#)

[Communications Medicine](#) **1**, Article number: 25 (2021) | [Cite this article](#)

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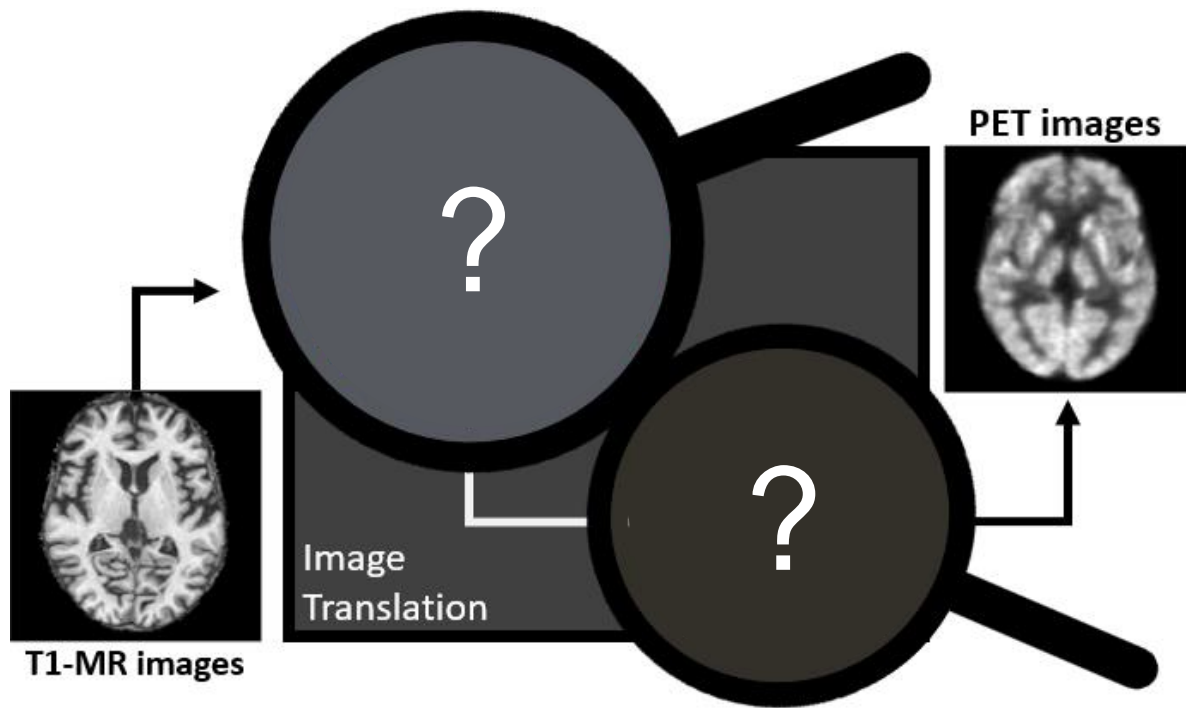
Underdiagnosis bias of artificial intelligence algorithms applied to chest radiographs in under-served patient populations

[Laleh Seyyed-Kalantari](#) , [Haoran Zhang](#), [Matthew B. A. McDermott](#), [Irene Y. Chen](#) & [Marzyeh Ghassemi](#)

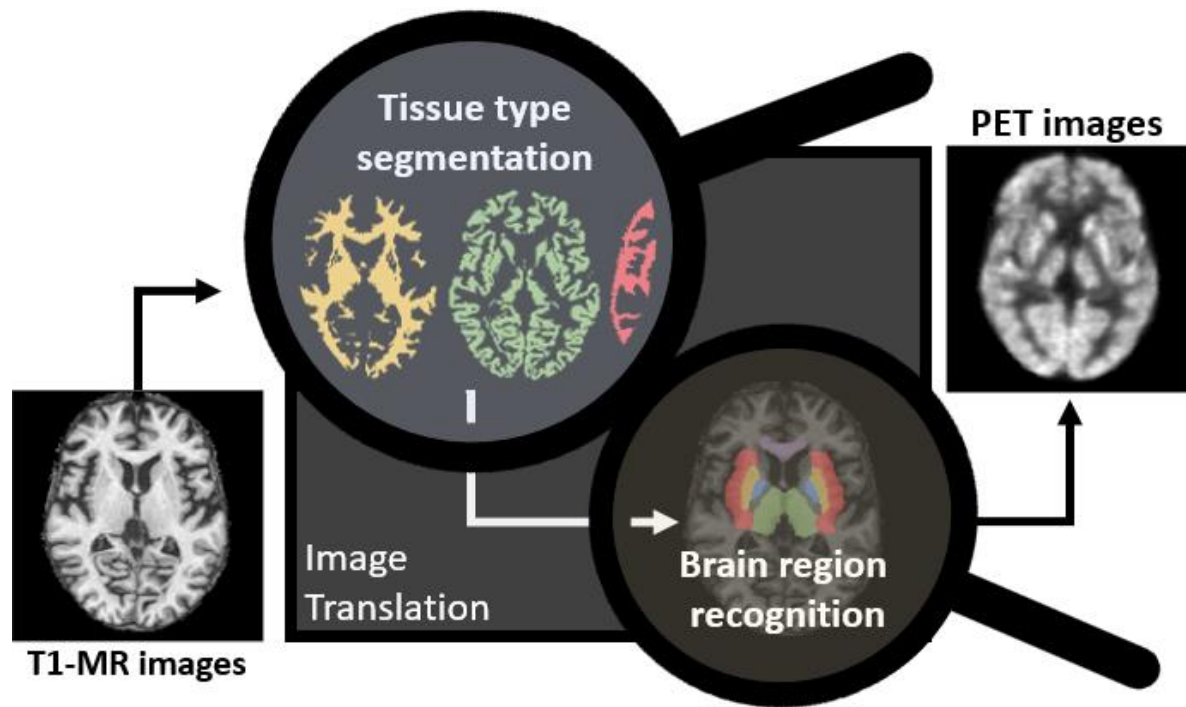
[Nature Medicine](#) **27**, 2176–2182 (2021) | [Cite this article](#)

6594 Accesses | **2** Citations | **58** Altmetric | [Metrics](#)

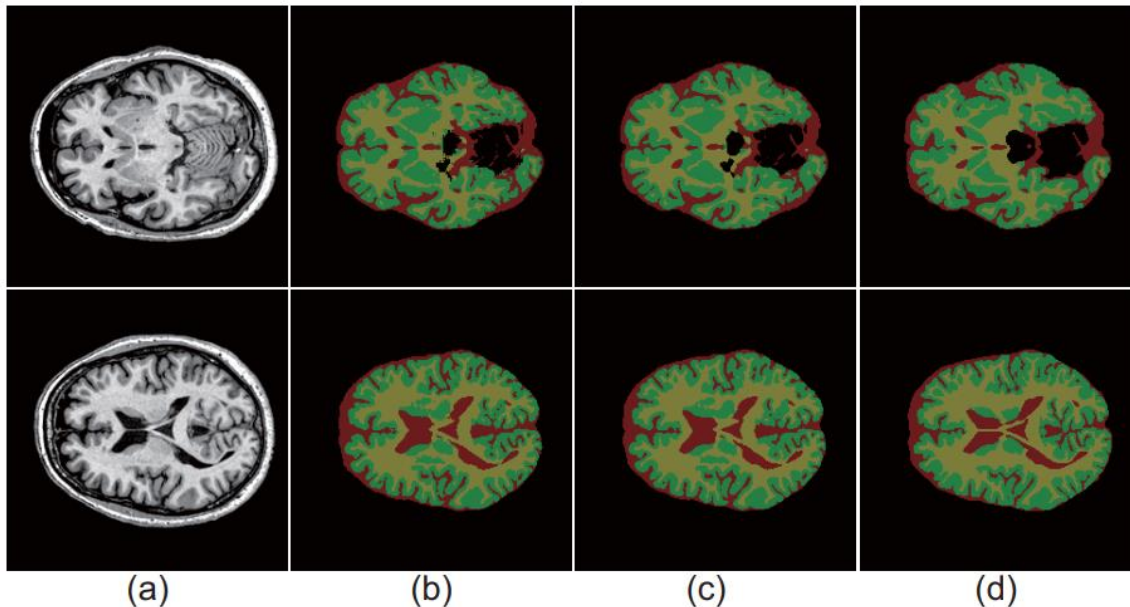
Motivation How do deep learning models translate T1-MRI images to FDG-PET images?



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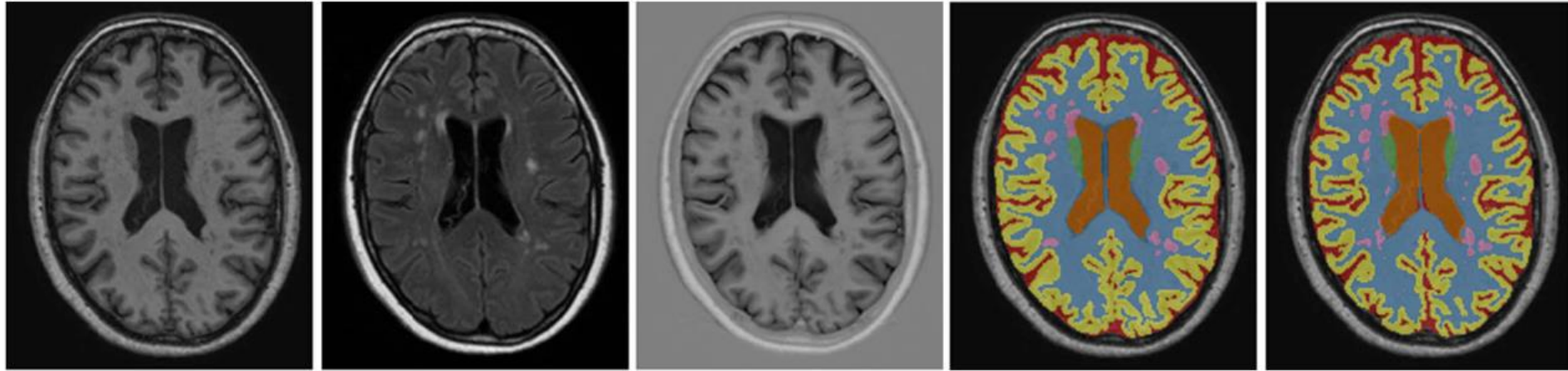


Literature Deep learning has been deployed for brain tissue segmentation and brain region identification.



The example results of validation data (yellow, green, and red colors represent the WM, GM, and CSF, respectively): (a) original MR images, (b) results of VoxResNet, (c) results of Autocontext VoxResNet, (d) ground truth label.

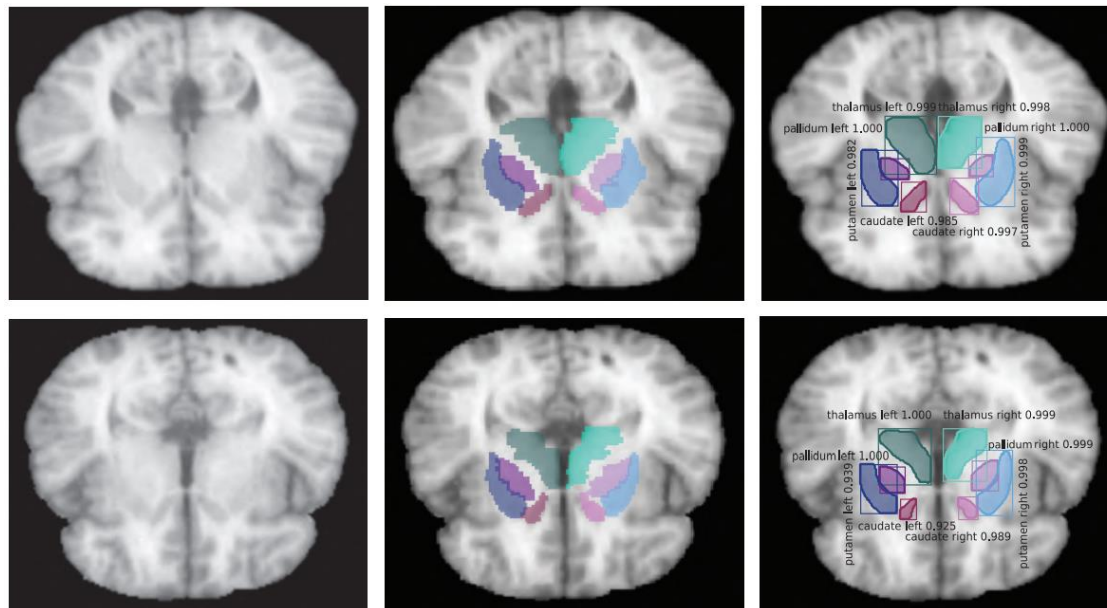
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Example segmentation for one of the test images from the MRBrainS13 challenge, trained using the 5 training images available within MRBrainS13.

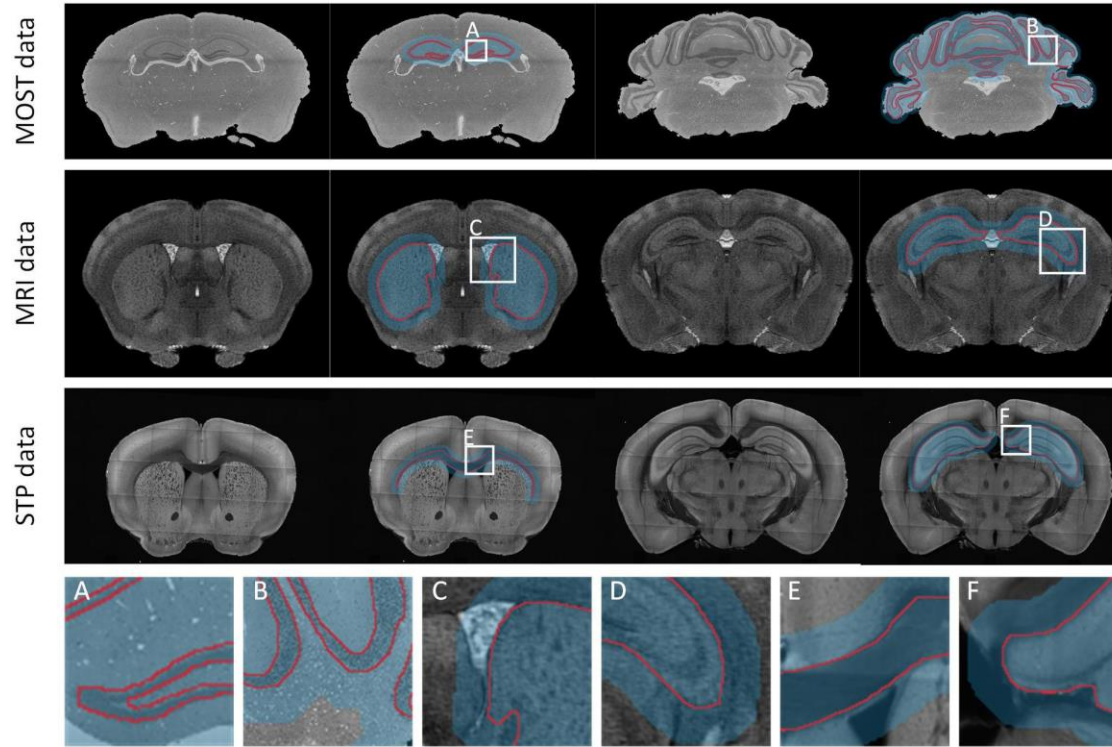
From left to right: T1-weighted image, T2-weighted FLAIR image, T1-weighted IR image, **reference segmentation** and **automatic segmentation**.

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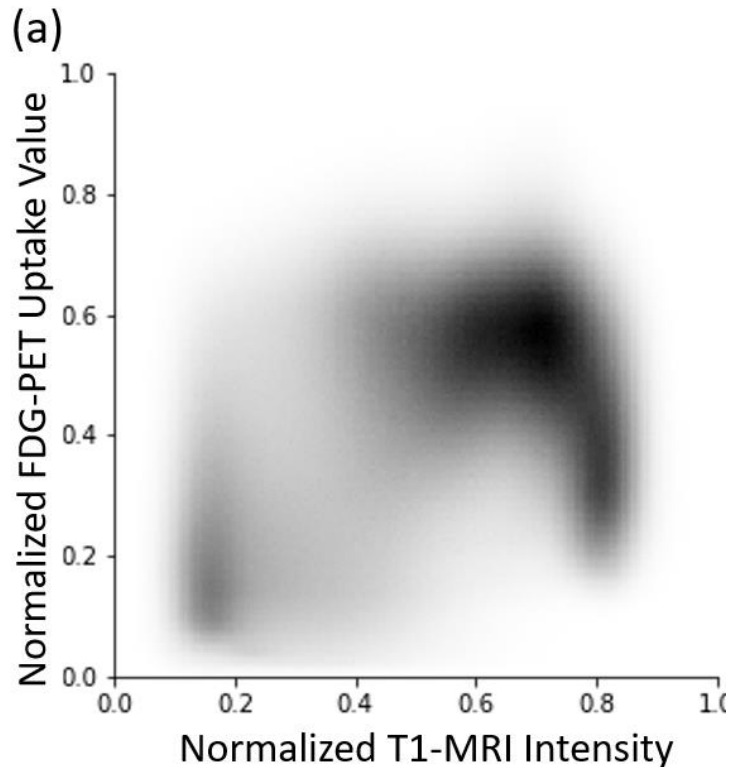
Left: Randomly selected human brain MR scans are shown. Middle: Ground-truth masks of the corresponding MR scans in (a). Right: Outputs of our proposed deep learning models.

Literature Deep learning has been deployed for brain tissue segmentation and brain region identification.

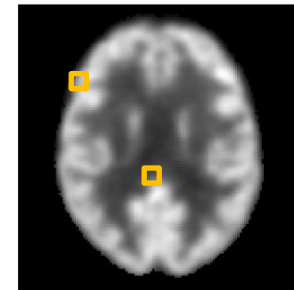


Tan, C., Guan, Y., Feng, Z., Ni, H., Zhang, Z., Wang, Z., ... & Li, A. (2020). DeepBrainSeg: Automated brain region segmentation for micro-optical images with a convolutional neural network. *Frontiers in neuroscience*, 14, 179.

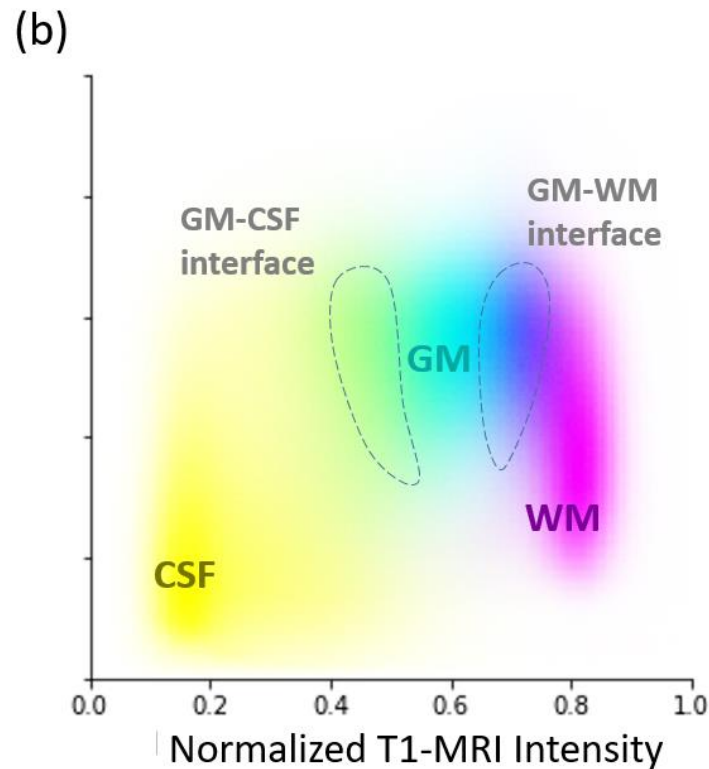
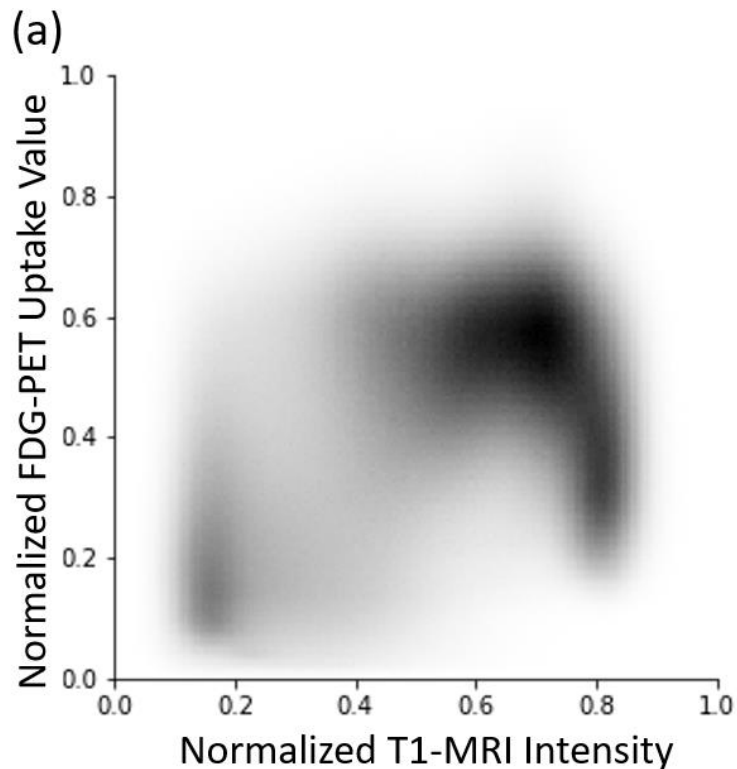
Hypothesis Deep learning models learn to segment brain tissue types.



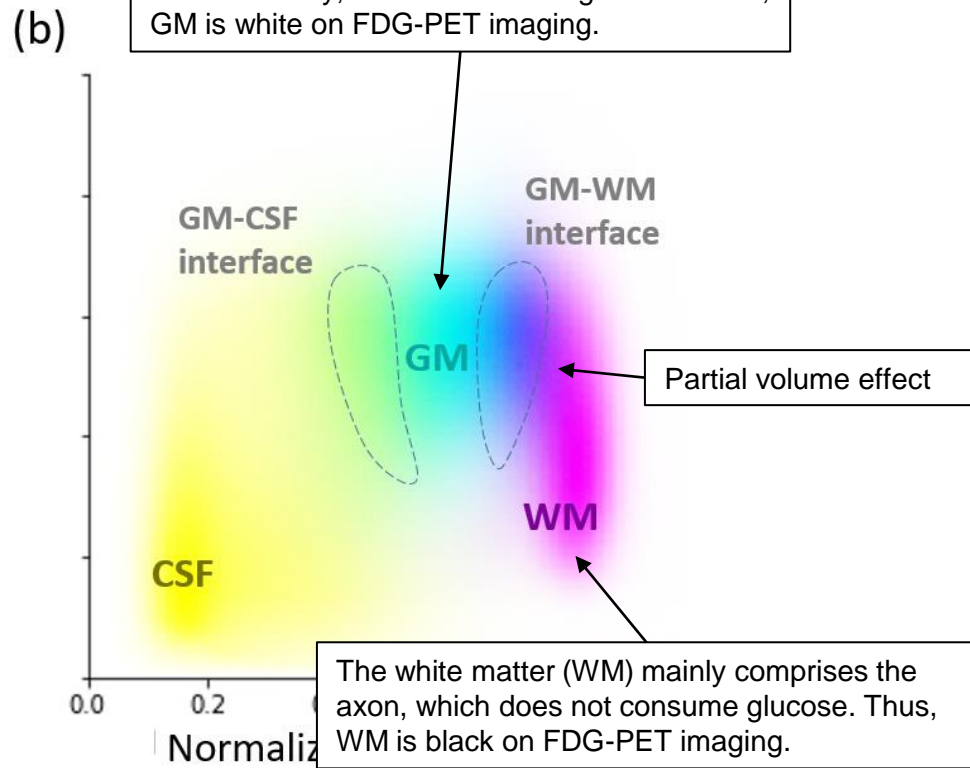
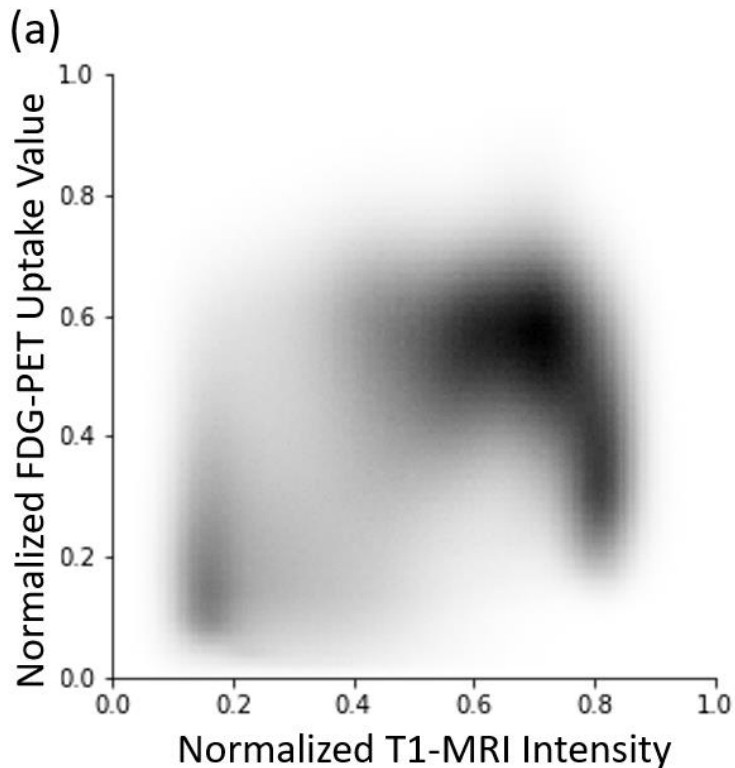
1. Align brain images from both T1-MRI and FDG-PET.
2. Use the value from T1-MRI as x and that from FDG-PET as y to obtain (x, y) pairs.



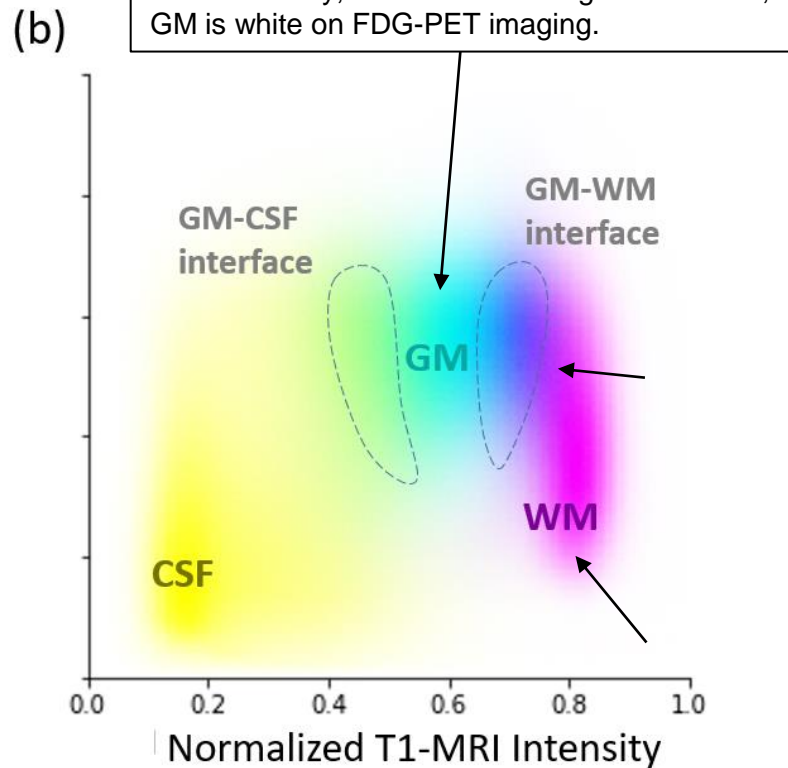
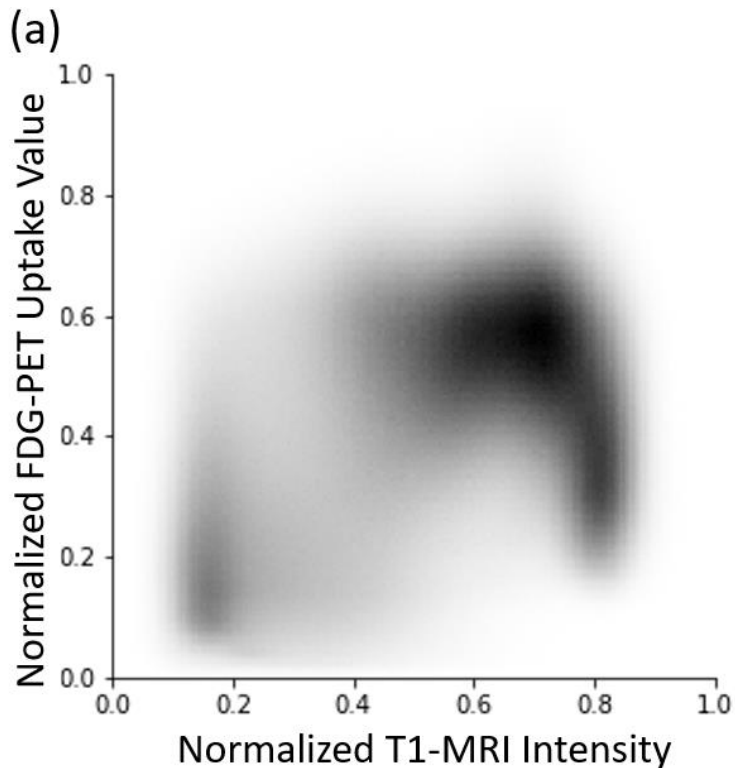
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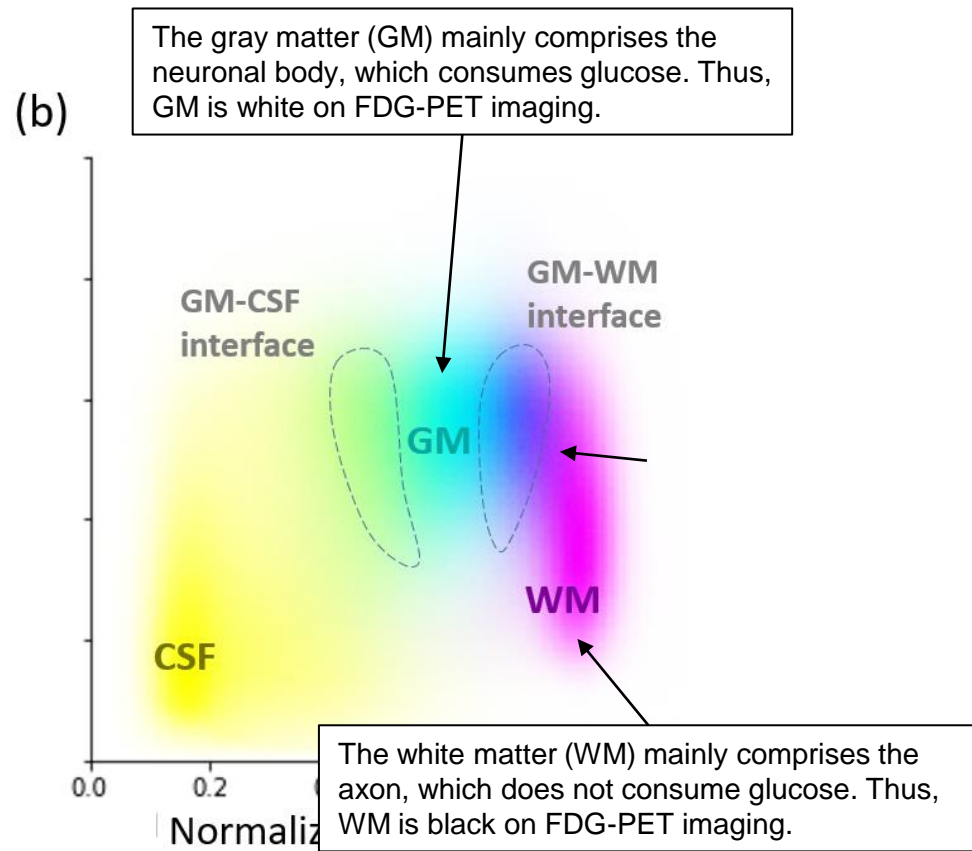
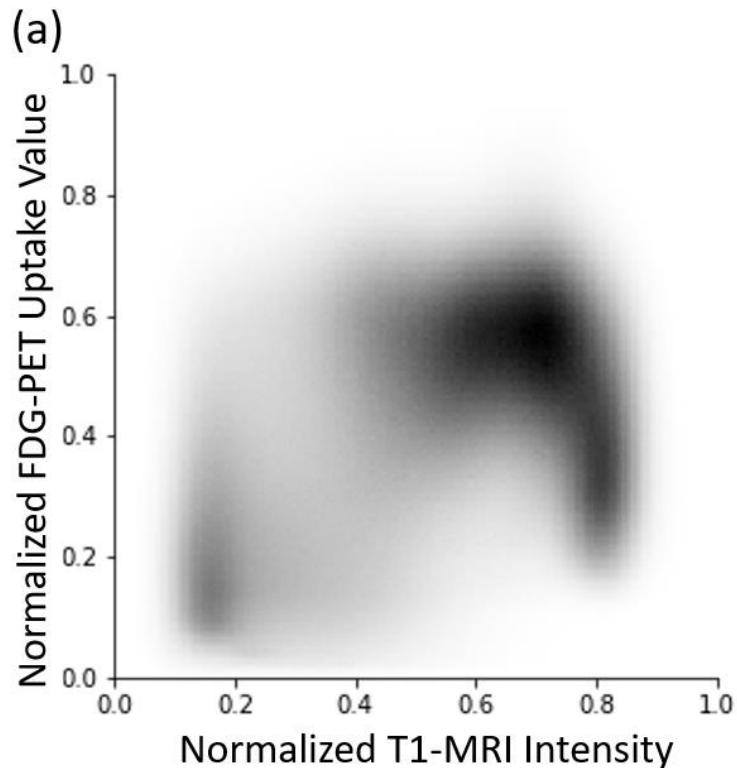
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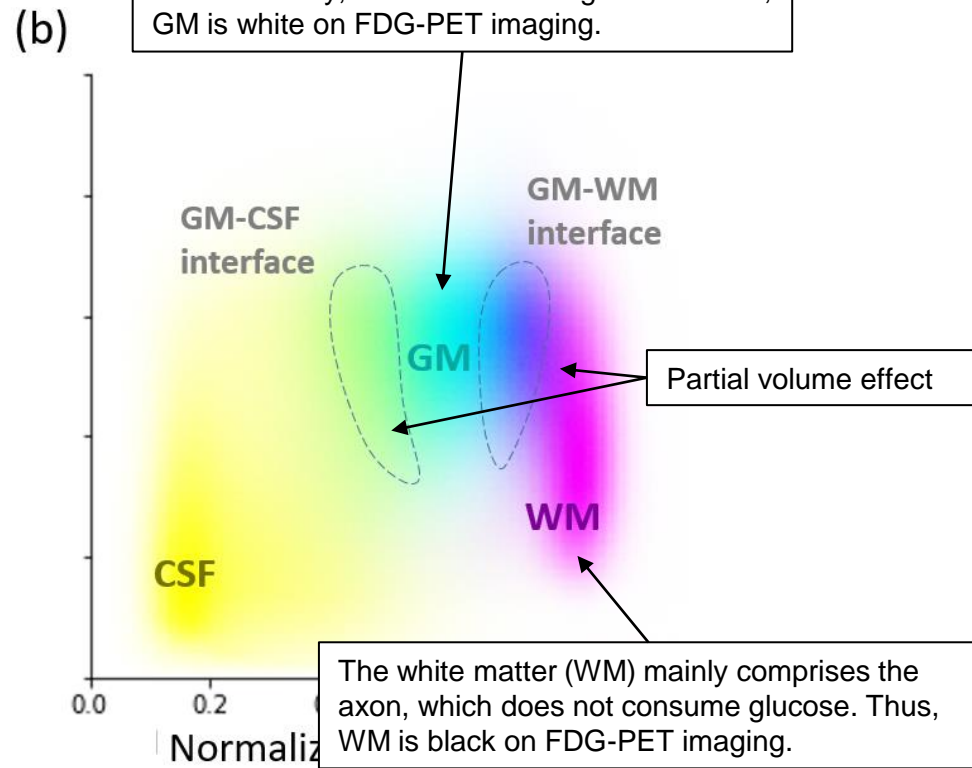
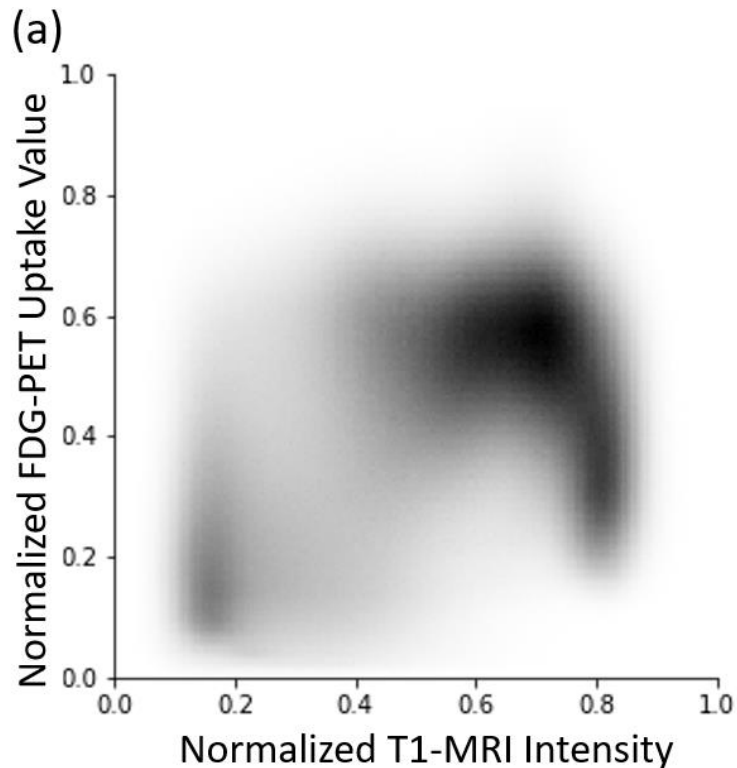
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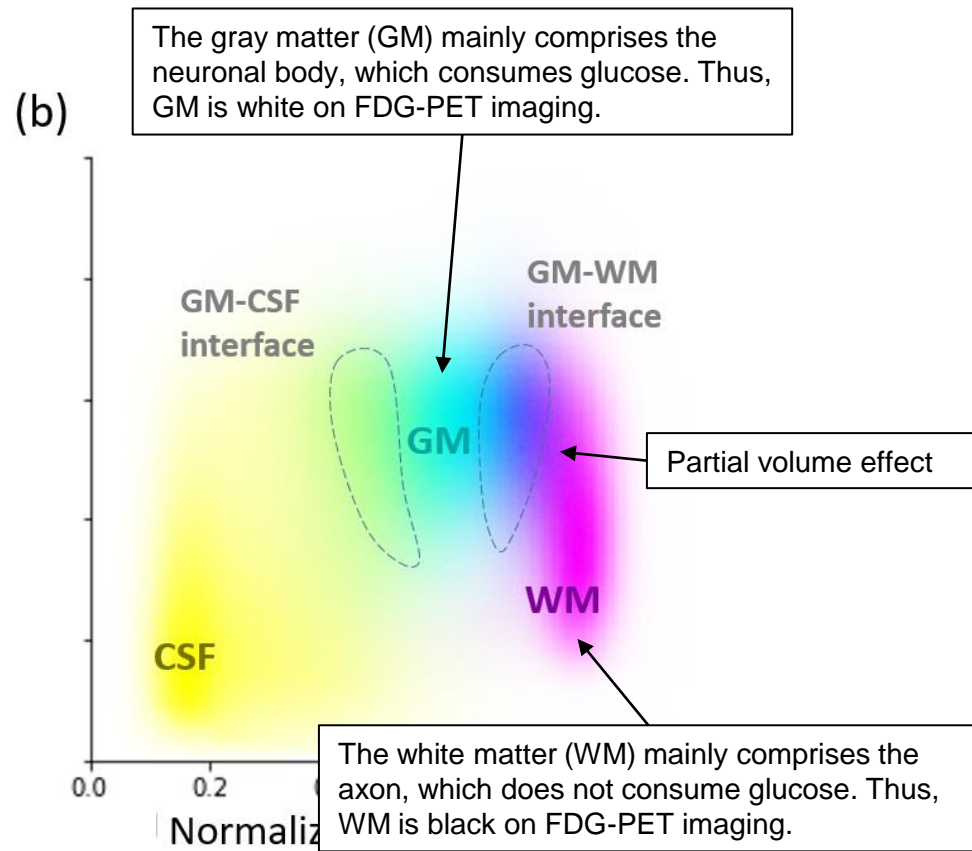
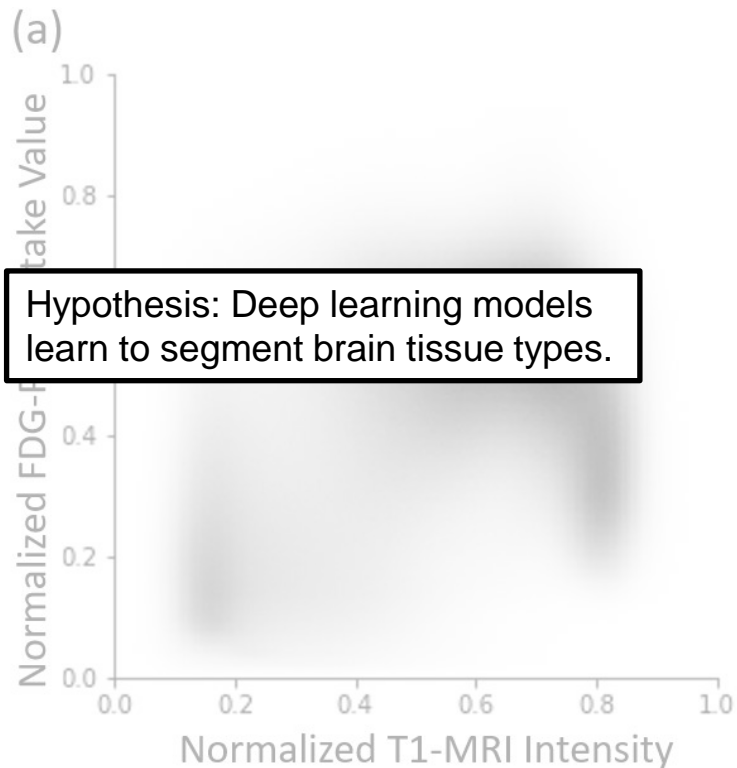
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Method

Dataset

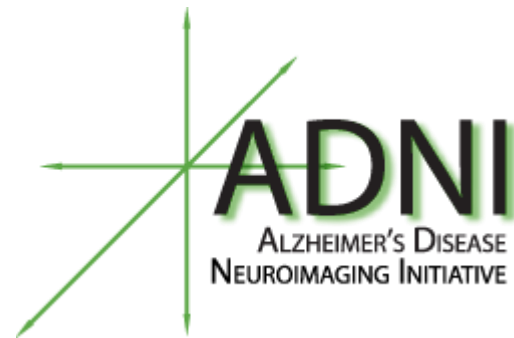
- Obtain **1300+ MRI-PET paired data** from Alzheimer's Disease Neuroimaging Initiative (ADNI), composed of brain images from cognitive normal people and patients with mild cognitive impairment or Alzheimer's disease. (International, multi-center, multi-scanning protocols, standard preprocessing procedures)
- Preprocess data according to the standard method with SPM software.

Deep learning model

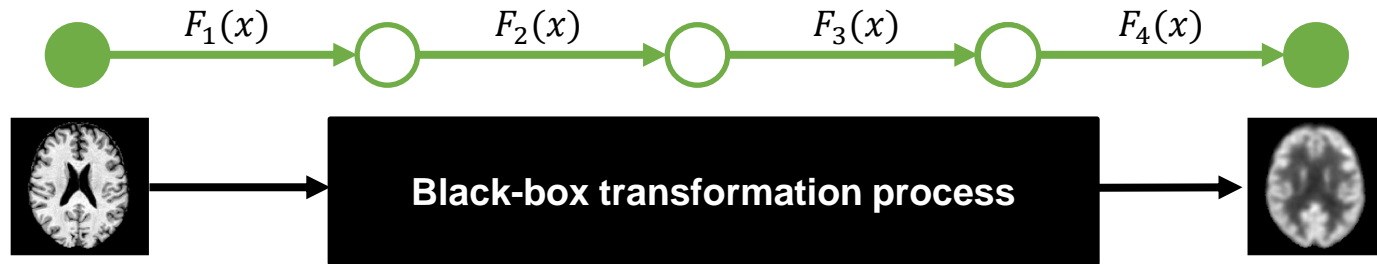
- Adopt a widely-used 2D Unet model for MRI-to-PET image translation, which comprises several nonlinear transformation functions $\{F_1, \dots, F_n\}$.

Goal

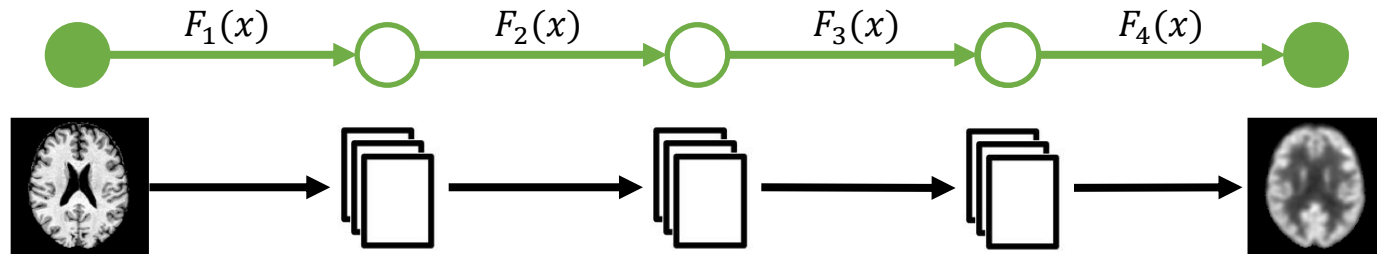
- Measure and quantify the information of “brain tissue type” along the data transformation process.



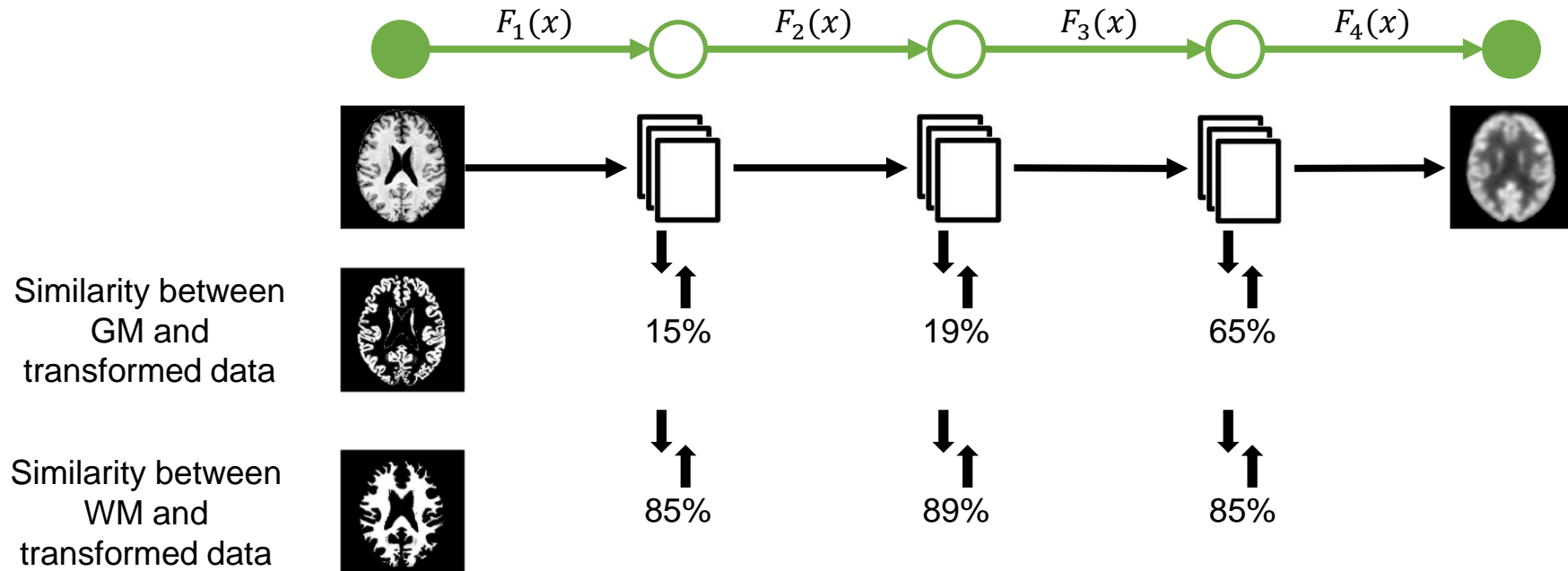
Method



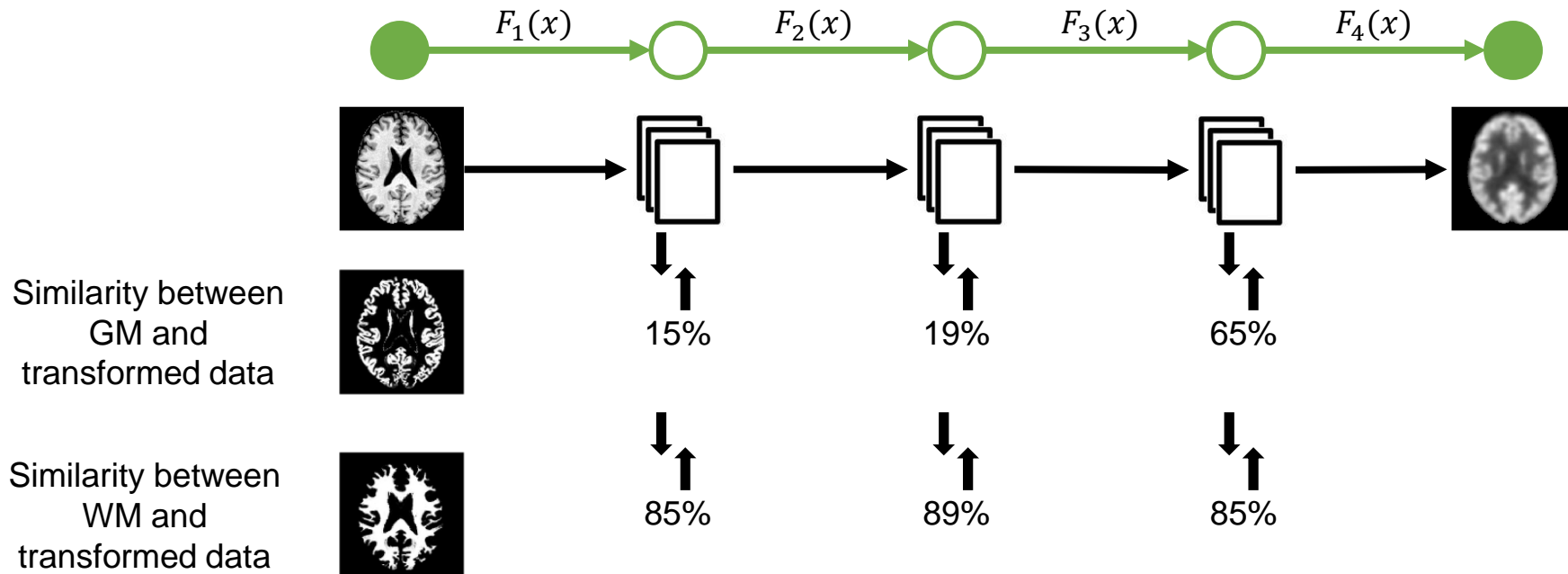
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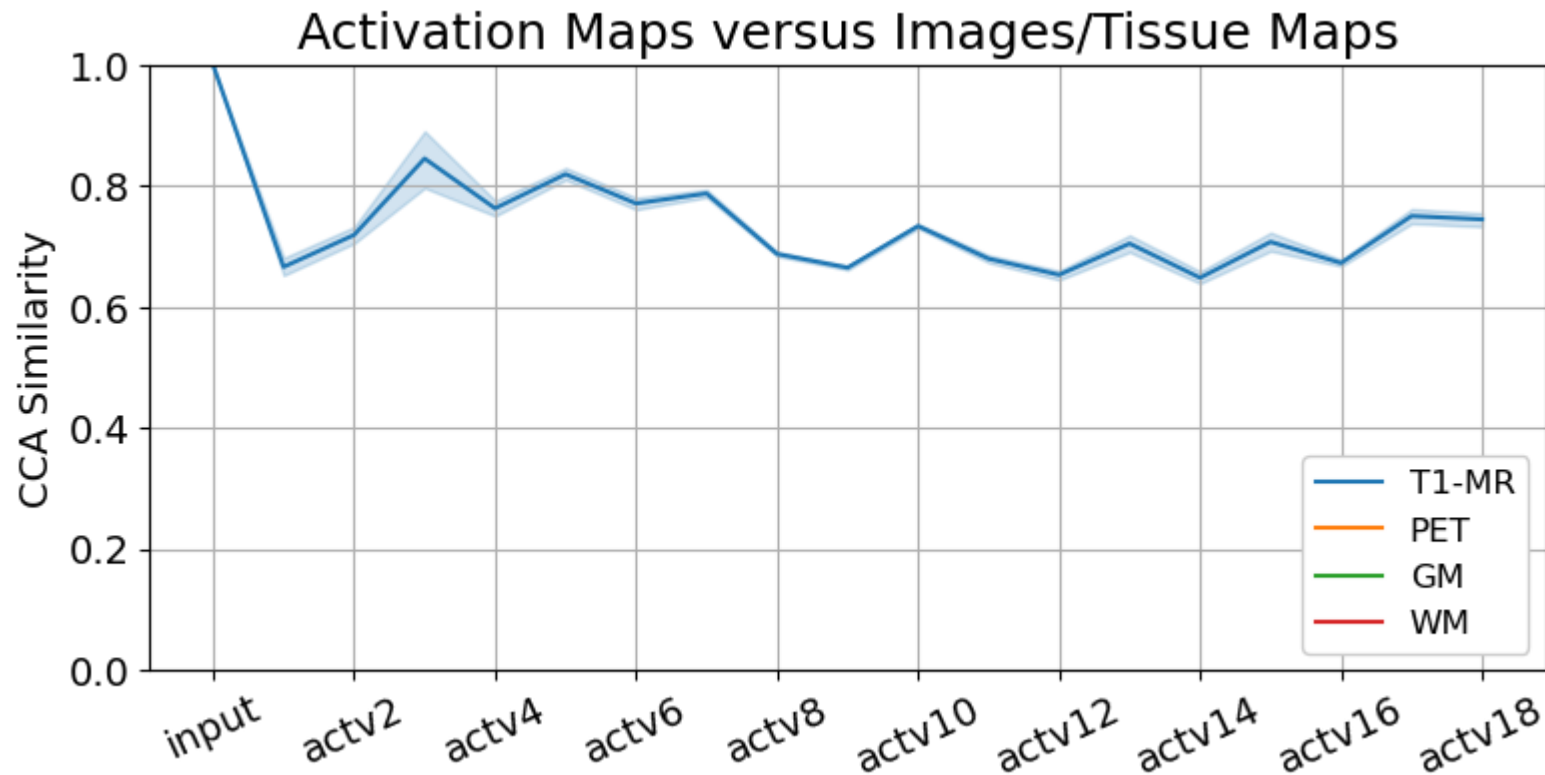


Method

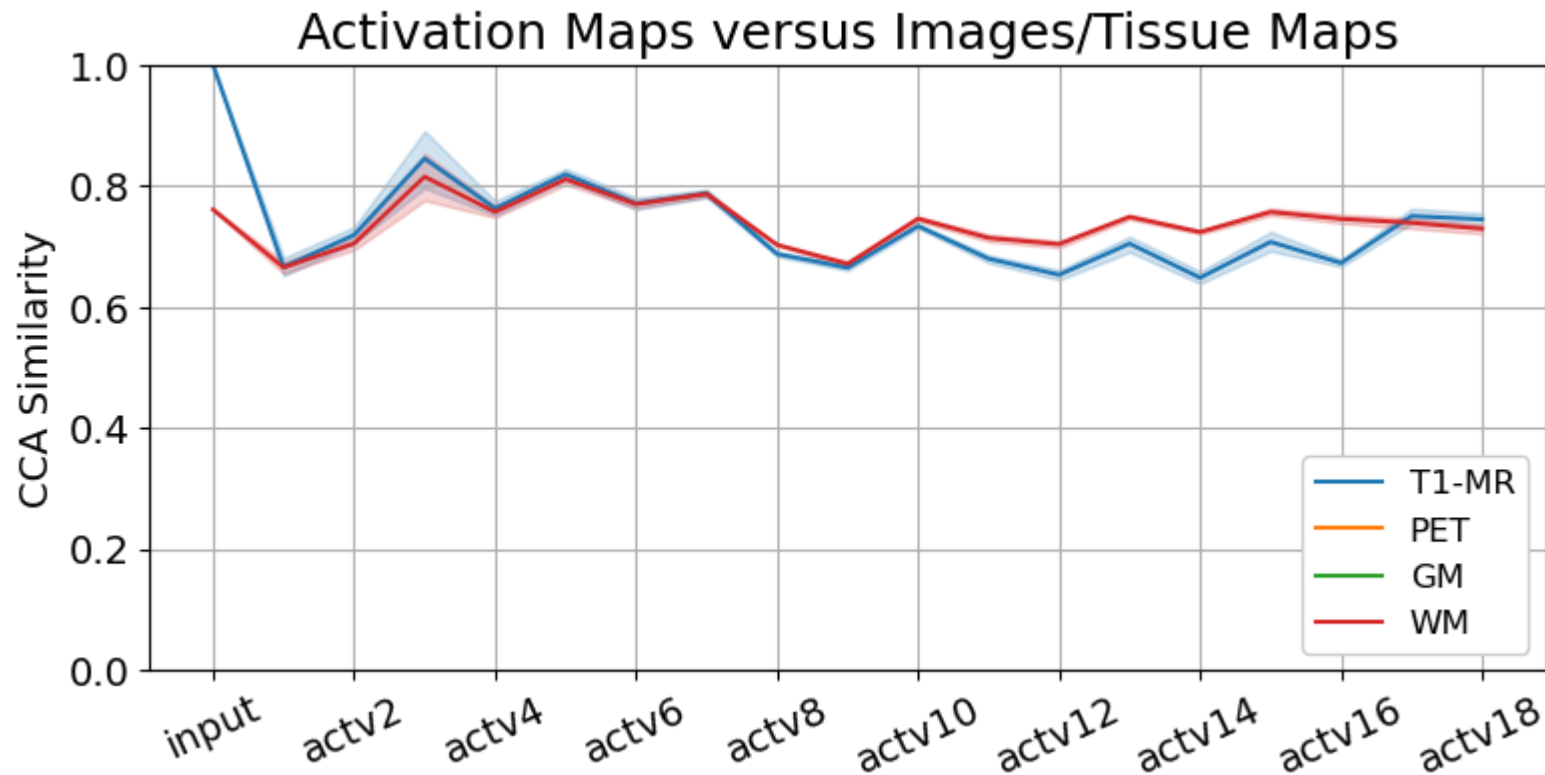


We can also compute the similarity between MRI (or PET) and the transformed data.

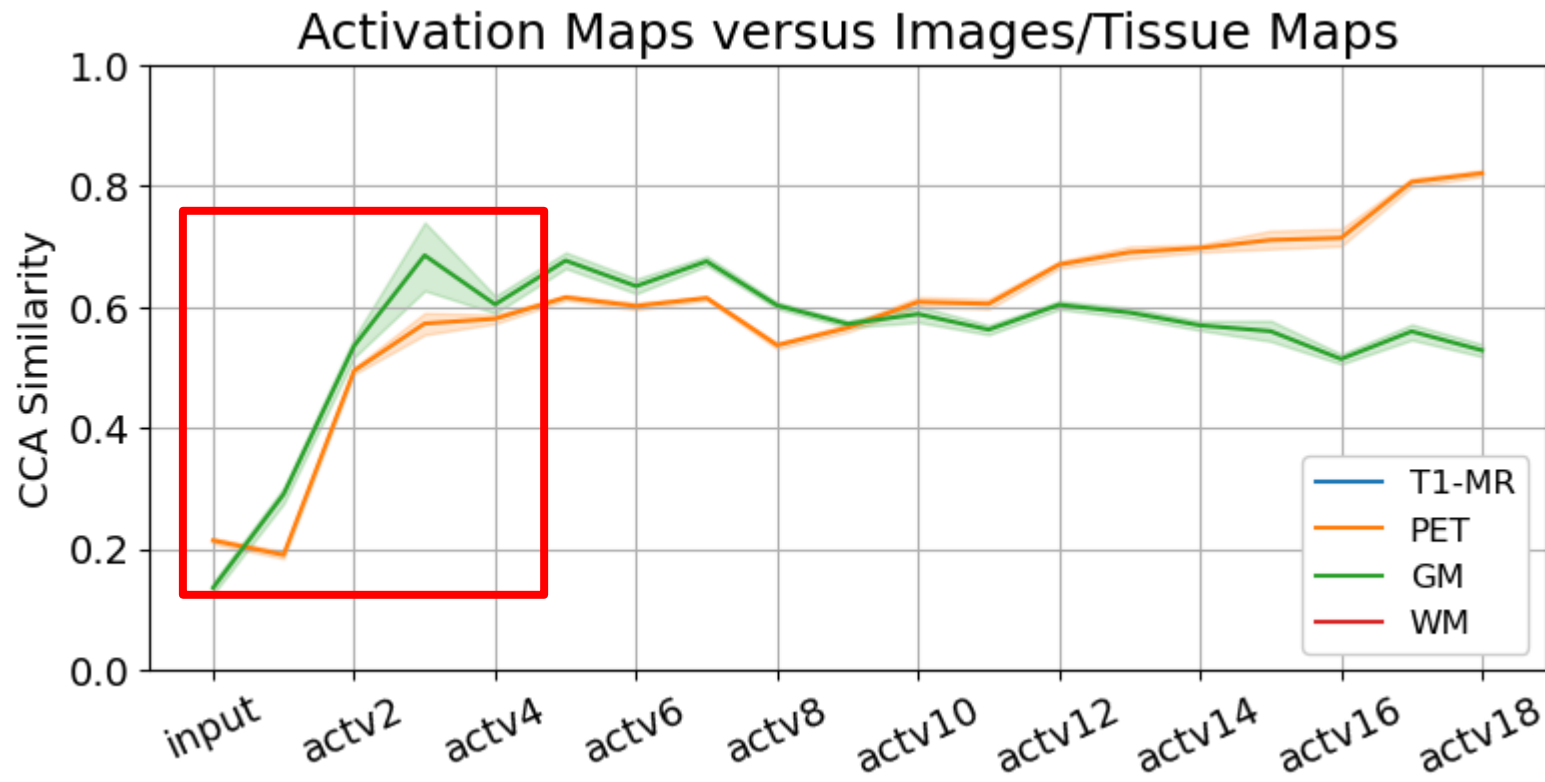
Results Brain tissues are segmented in the early encoding stage of image translation.



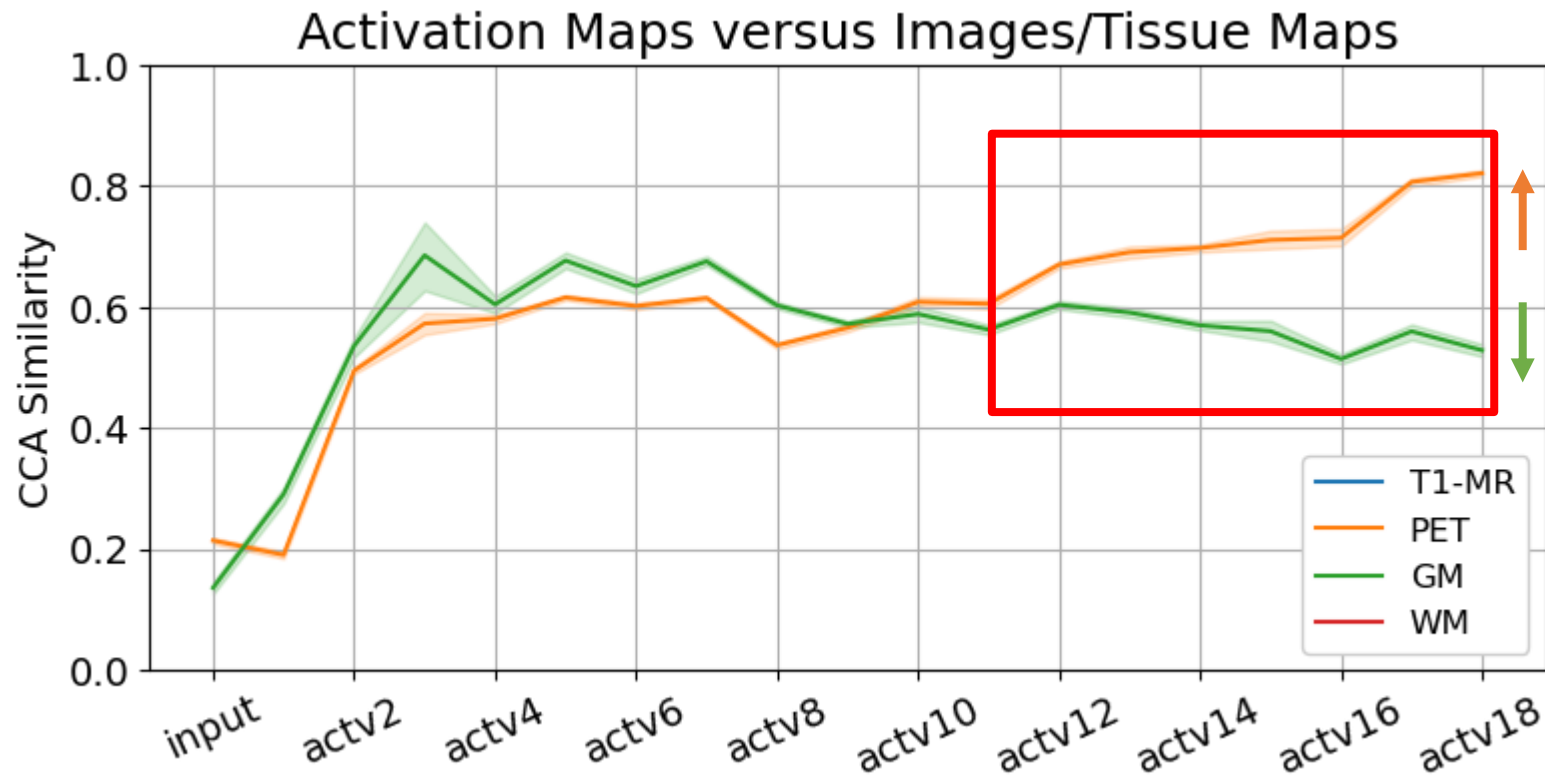
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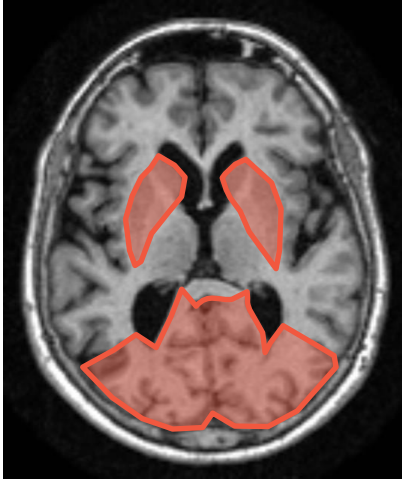


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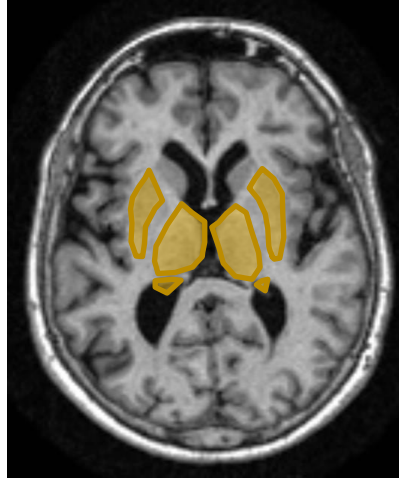


Hypothesis Deep learning models learn to segment brain regions.

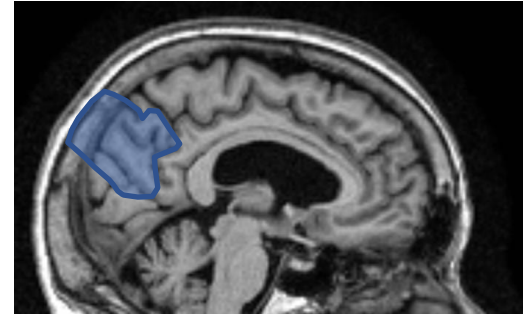
Normal higher uptake regions



Least altered regions during aging

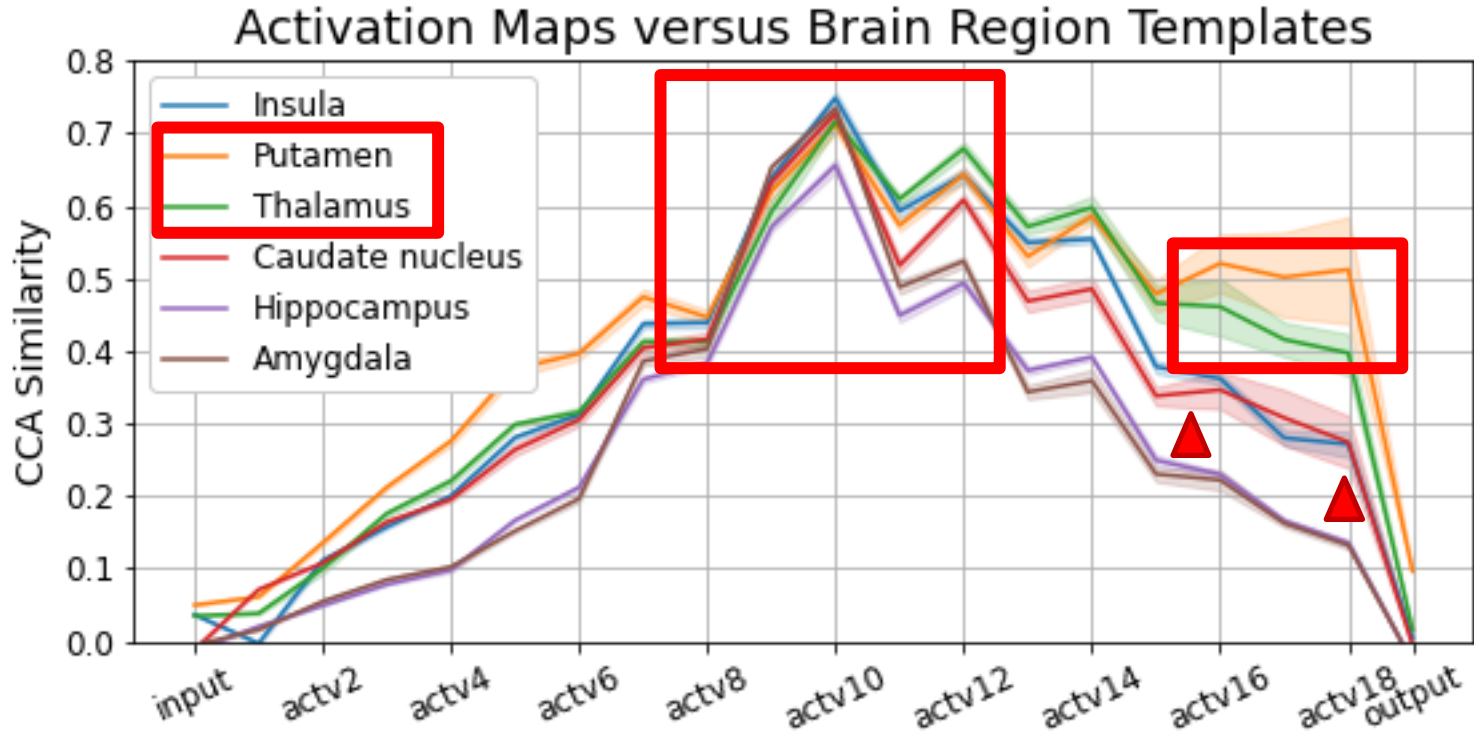


Regions of hypometabolism in AD



Brain: normal variations and benign findings in fluorodeoxyglucose-PET/computed tomography imaging. (PET Clin. 2014)
Voxel-based mapping of brain gray matter volume and glucose metabolism profiles in normal aging. (Neurobiology of Aging. 2007)
Brain PET in the diagnosis of Alzheimer's disease. (Clin Nucl Med. 2014)

Results Brain regions are identified in the decoding stage of translation.



Limitation

Dataset – organs, diseases, modalities (e.g., different radioactive tracers), machines.

Models – various deep learning models.

Measurement of similarity – curse of dimensionality.

Take home message

We use representational similarity to verify our proposed hypotheses that the translation from T1-MR to PET images comprises the recognition of brain tissue types and brain regions.

