



Retrieval-Augmented Few-shot Skull Stripping: Implementation and Evaluation

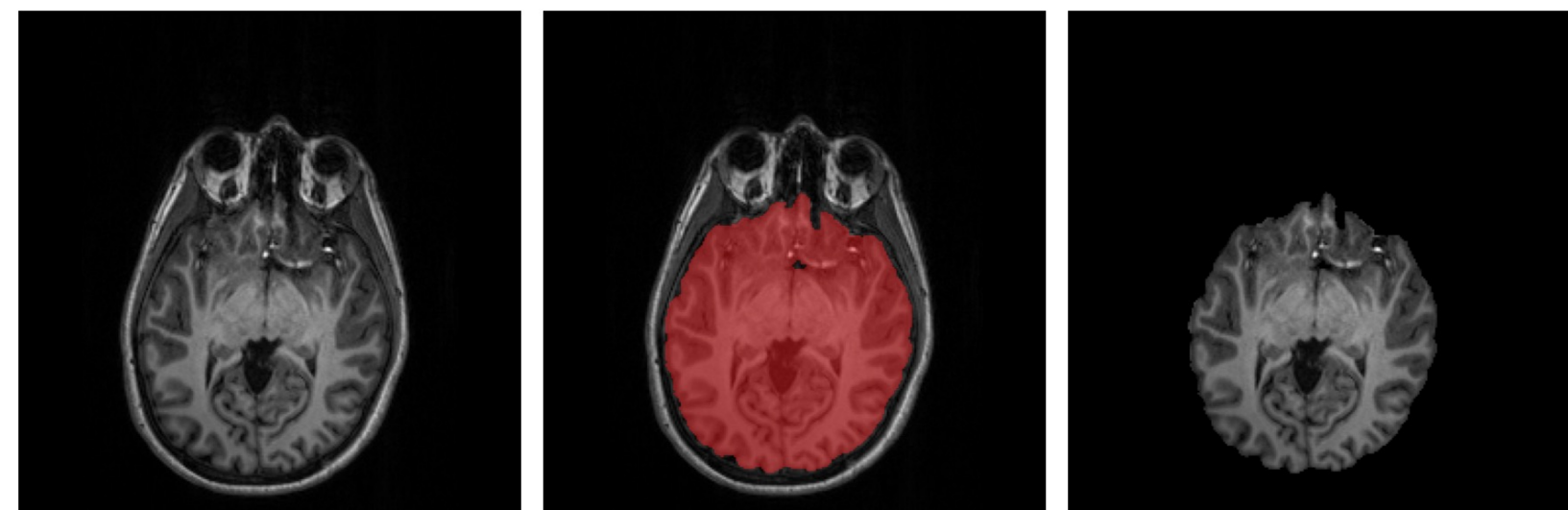
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GitHub

INTRODUCTION

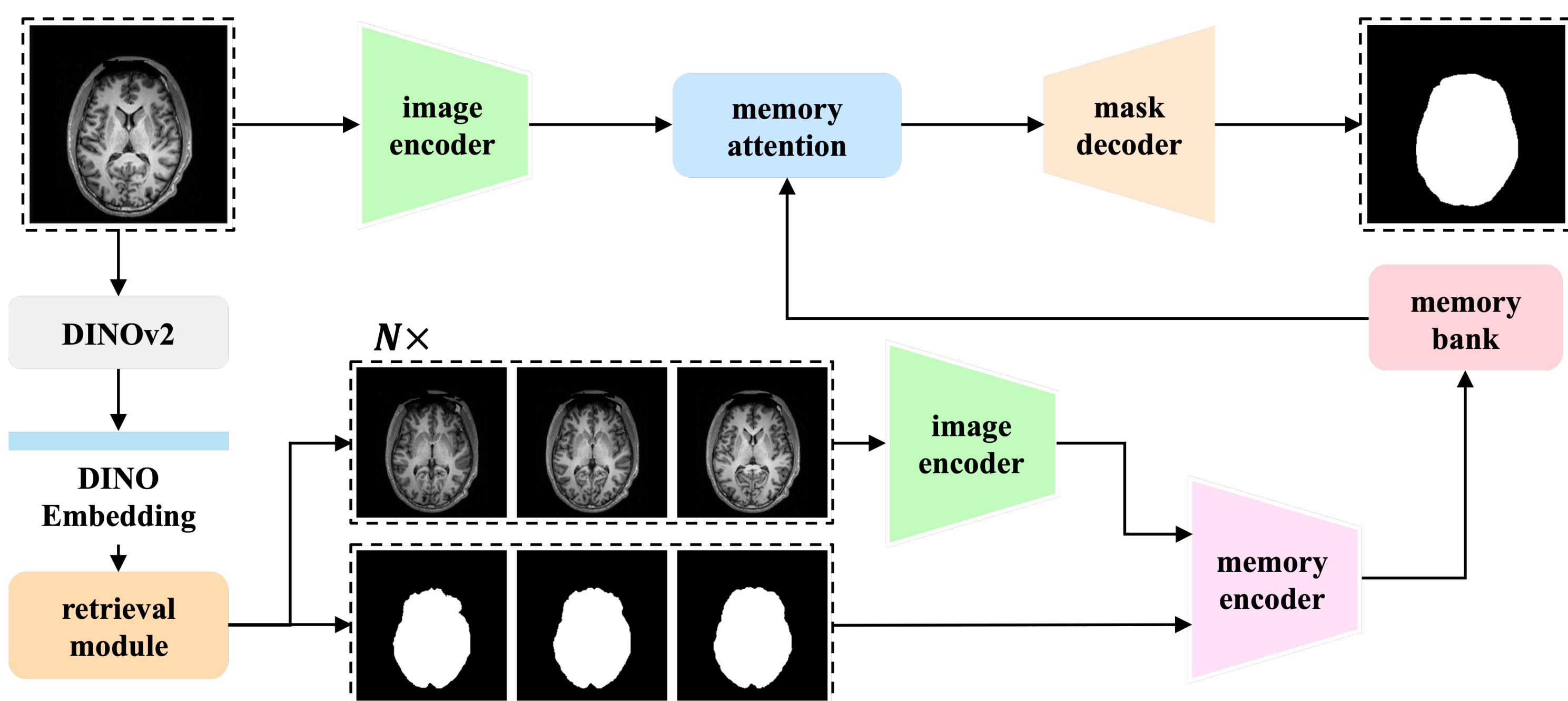
- Skull Stripping is a crucial preprocessing step in brain image analysis, as it removes non-brain tissues (e.g., skull, scalp) from MRI scans. This process improves the accuracy of downstream tasks such as brain tissue segmentation, registration, and disease diagnosis.



- Recent visual foundation models such as DINOv2 [2] and SAM 2 [3] show strong zero-shot generalization across various domain shifts, including the medical images.
- Furthermore, recent study[1] has demonstrated that Retrieval-Augmentation (RA) significantly enhances performance in medical image segmentation tasks.
- In this work, we implement the Retrieval-Augmented Few-shot Medical Image Segmentation (RAFS) framework and evaluate its effectiveness on the skull stripping task. Note that, this method is “training-free approach”, which means it does not require any additional training.
- All codes are available at https://github.com/kdh-yu/BME_Capstone1.

METHOD

- Pipeline Overview



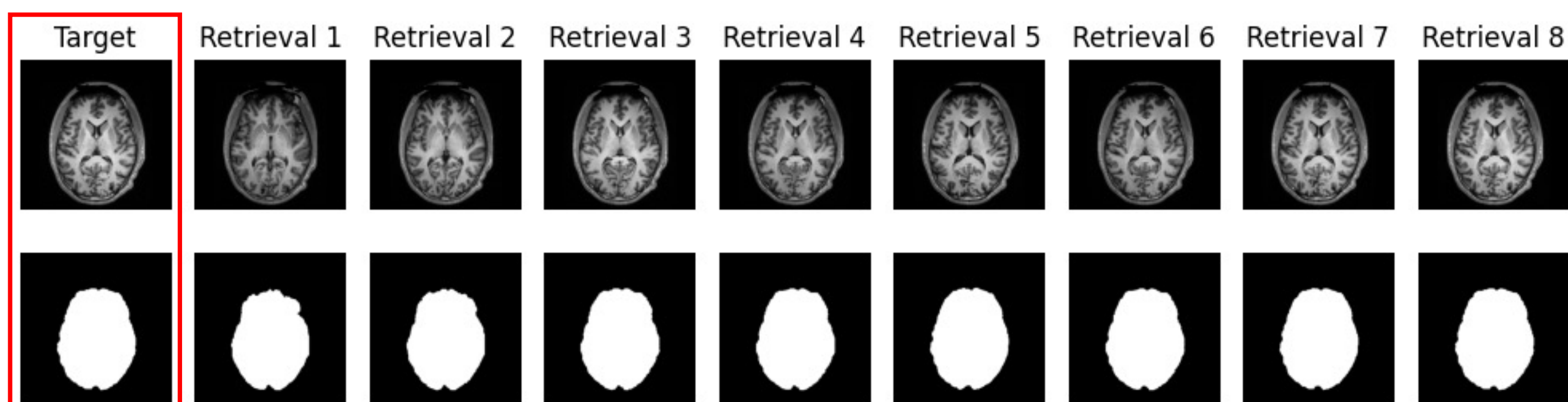
- First Stage: FAISS Index Construction

- DINOv2 Vision Transformer extracts normalized [CLS] embedding from brain MRI scans.
- 784 axial slices from 3 annotated volumetric data are added to a FAISS index



- Second Stage: Retrieval-Augmented Segmentation

- Retrieval: The FAISS Index is queried to retrieve N similar images, which are then treated as “Past Frames”.
- Segmentation: SAM 2 Video Segmentation is performed to predict the current image’s brain mask.



RESULTS

- Implementation Details

- FAISS Index: Configured with Euclidean Distance (L2)
 - Vectors are normalized during both constructing and querying
 - Model retrieves 8 slices to predict current mask. (N=8)
- SAM 2: Hiera-Large Model
- DINOv2: ViT-S/14 with registers

- Dataset

- NFBS Skull Stripped Repository [4]
 - Anonymized T1w MRI Dataset
 - Includes 125 volumetric data with size (256, 256, 192)

- Quantitative Analysis (Dice Score)

	Slice-wise	Volume-wise
RAFS (ours)	0.8969	0.9594
U-Net (3 Subj)	0.9250	0.9747
U-Net	0.9677	0.9918
Swin U-Net (3 Subj)	0.8279	0.9204
Swin U-Net	0.9671	0.9881

- Our implemented method demonstrates performance on par with fully-supervised U-Net and Swin U-Net
- Notably, it also excels when applied to volumetric data, suggesting it’s a promising solution for practical skull stripping tasks.

CONCLUSION

- Our work demonstrates that the Retrieval-Augmented Few-shot Medical Image Segmentation (RAFS) framework achieves robust performance, not only on 2D slices but also on 3D volumetric data.
- One key advantage of RAFS is its training-free nature, eliminating the need for extensive fine-tuning on new datasets, which makes RAFS a highly efficient and adaptable solution for skull stripping, with the potential to be expanded for diverse tasks such as tissue segmentation.

LIMITATIONS

- One limitation is its reduced ability to produce highly intricate or geometrically complex masks. We hypothesize this stems from two main factors:
 - A potential loss of fine-grained detail when integrating retrieved images and masks into memory
 - Performance bottlenecks exacerbated by significant domain shift



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