

ClinicalFMamba: Advancing Clinical Assessment using Mamba-based Multimodal Neuroimaging Fusion

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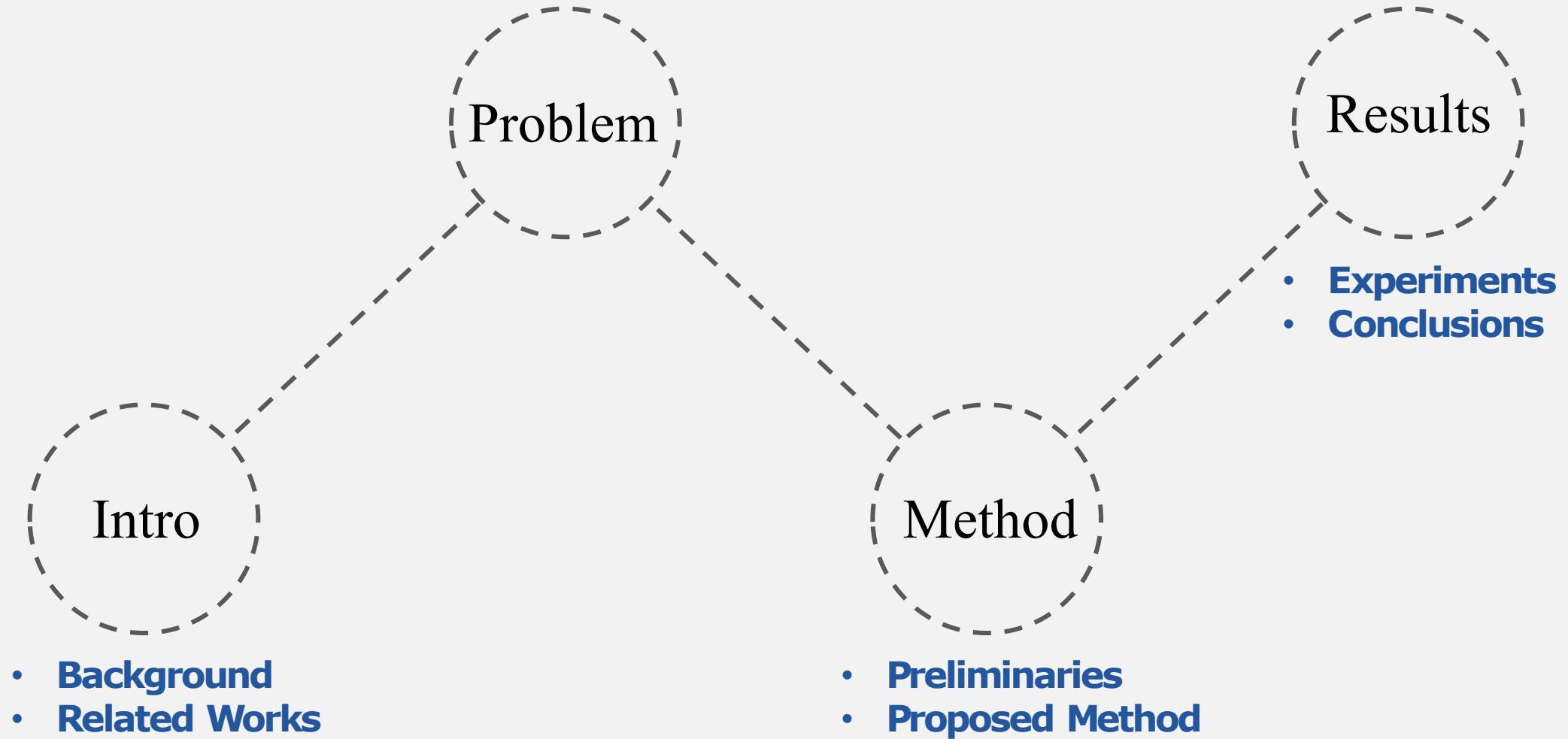
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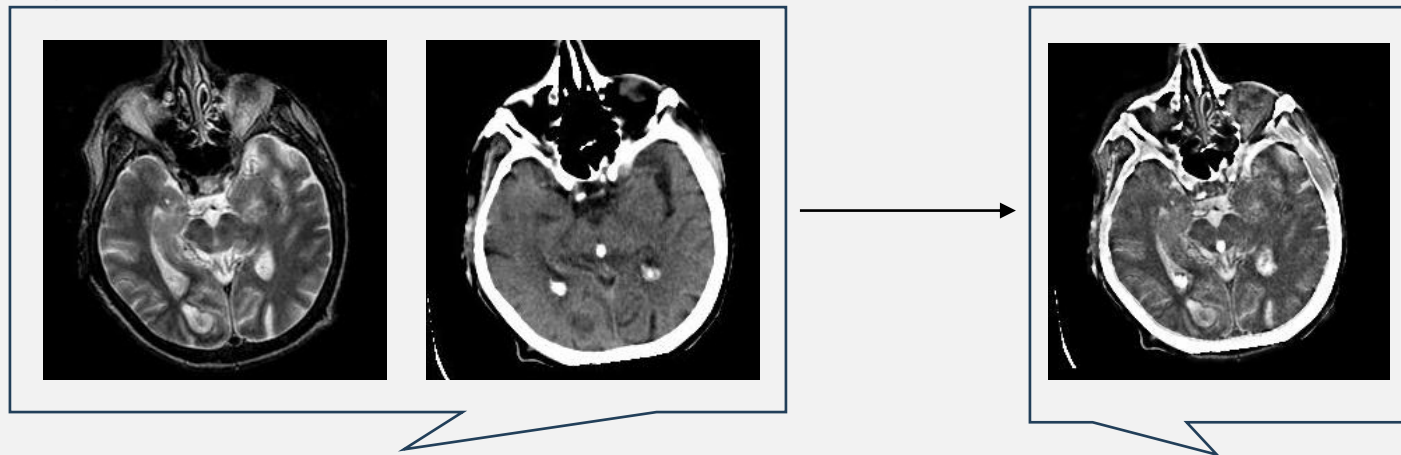
Outline



Intro

Background

- Multimodal medical image fusion plays an increasingly prominent role in clinical diagnosis.
- It aims to aggregate complementary information from different image modalities to produce higher-quality fused images (e.g., anatomical and functional images)
- Due to hardware constraints and current physical imaging principles, individual modalities can only capture specific aspects of tissue characteristics, leading to incomplete diagnostic information.



Related Works

- CNN-based:
 - MSRPAN [1]: Residual pyramid attention network for multimodal medical image fusion and Feature Energy Ratio Strategy to fuse feature maps
 - MSDRA [2]: Double residual attention network for multimodal medical image fusion and uses weighted L1 Norm to fuse feature maps.
 - EH-DRAN [3]: Dilated residual attention network + edge enhancer to extract multi-scale features and enhance edge details. A family of parameter-free fusion strategies is proposed to fuse feature maps in the latent space.



Limitations...

1. limited by their inherent local receptive fields, which restrict their ability to capture long-range spatial dependencies.
2. Most of the methods are in two-stage, which create another layer of computation

[1] Fu, J., Li, W., Du, J., & Huang, Y. (2021). A multiscale residual pyramid attention network for medical image fusion. *Biomedical Signal Processing and Control*, 66, 102488.
[2] Li, W., Peng, X., Fu, J., Wang, G., Huang, Y., & Chao, F. (2022). A multiscale double-branch residual attention network for anatomical–functional medical image fusion. *Computers in biology and medicine*, 141, 105005.
[3] Zhou M, Zhang Y, Xu X, Wang J, Khalvati F. Edge-Enhanced Dilated Residual Attention Network for Multimodal Medical Image Fusion. In 2024 IEEE International Conference on Bioinformatics and Biomedicine (BIBM) 2024 Dec 3 (pp. 4108-4111). IEEE.

Related Works

- Transformer-based:
 - SwinFusion [4]: Combining a CNN feature extractor with a cross-domain transformer model to fuse local and global information.
 - MRSCFusion [5]: Combining a multiscale CNN model and applied residual Swin Transformer layers to fuse cross-domain information.
 - MACTFusion [6]: Light-weight cross modality transformer with window and grid attention.



Limitations...

1. Self-Attention mechanism requires quadratic computational complexity, limiting the practical applications on large image
2. Applying the fusion method to real clinical diagnosis tasks is not fully explored.

[4] Ma, J., Tang, L., Fan, F., Huang, J., Mei, X., & Ma, Y. (2022). SwinFusion: Cross-domain long-range learning for general image fusion via swin transformer. *IEEE/CAA Journal of Automatica Sinica*, 9(7), 1200-1217.

[5] Xie, X., Zhang, X., Ye, S., Xiong, D., Ouyang, L., Yang, B., ... & Wan, Y. (2023). MRSCFusion: Joint residual Swin transformer and multiscale CNN for unsupervised multimodal medical image fusion. *IEEE Transactions on Instrumentation and Measurement*.

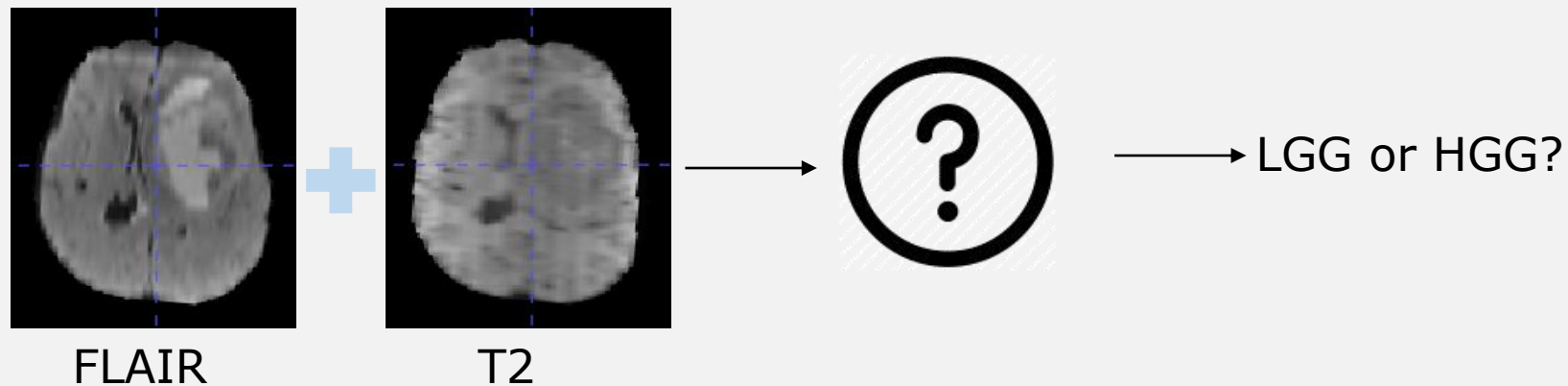
[6] Xie, X., Zhang, X., Tang, X., Zhao, J., Xiong, D., Ouyang, L., ... & Teo, K. L. (2024). MACTFusion: Lightweight Cross Transformer for Adaptive Multimodal Medical Image Fusion. *IEEE Journal of Biomedical and Health Informatics*.

Problem

- Two most common fusion tasks in medical imaging: MRI-CT and MRI-SPECT fusion tasks (MRI-CT and MRI-SPECT from Harvard Whole Brain Atlas datasets)



- To further validate the effectiveness of the fusion method, we apply it to a downstream clinical brain tumor pathology classification task between Low-Grade and High-Grade Gliomas (BraTS 2019 dataset).





Method

- Preliminaries
- Proposed Method
- Scanning Strategy

Preliminaries

Mamba is a framework that model sequential data through a hidden state that evolves over time. The transformation from a 1-D sequence $x(t)$ to an output $y(t)$ is achieved through a hidden state $h(t)$. The implementation is typically achieved through linear ODEs:

$$h'(t) = Ah(t) + Bx(t), y(t) = Ch(t)$$

To discretize, we use zero-order hold:

$$\bar{A} = \exp(\Delta A), \bar{B} = (\Delta A)^{-1}(\exp(\Delta A) - I) \cdot \Delta B$$

We can rewrite the first equation as:

$$h_t = \bar{A}h_{t-1} + \bar{B}x_t, y_t = Ch_t$$

Selective-scan:

1. can focus on or ignore particular information, like self-attention in Transformer
2. B, C, Δ is dynamic to the input, allowing the model to filter relevant information and focus on important context within long sequences
3. B : filter out irrelevant data or to focus on important data to allow only relevant data to enter into the new state.
4. C : selective to decide which information from the state is required to materialize the output
5. Δ : weighting between the importance of new samples compared to the previous state.

Proposed Method

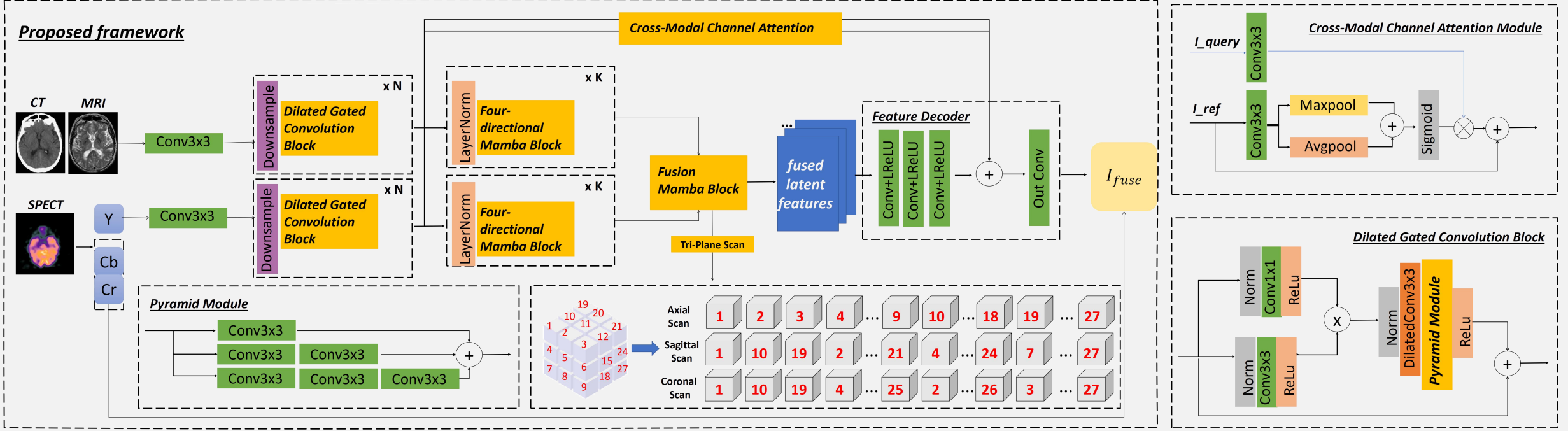
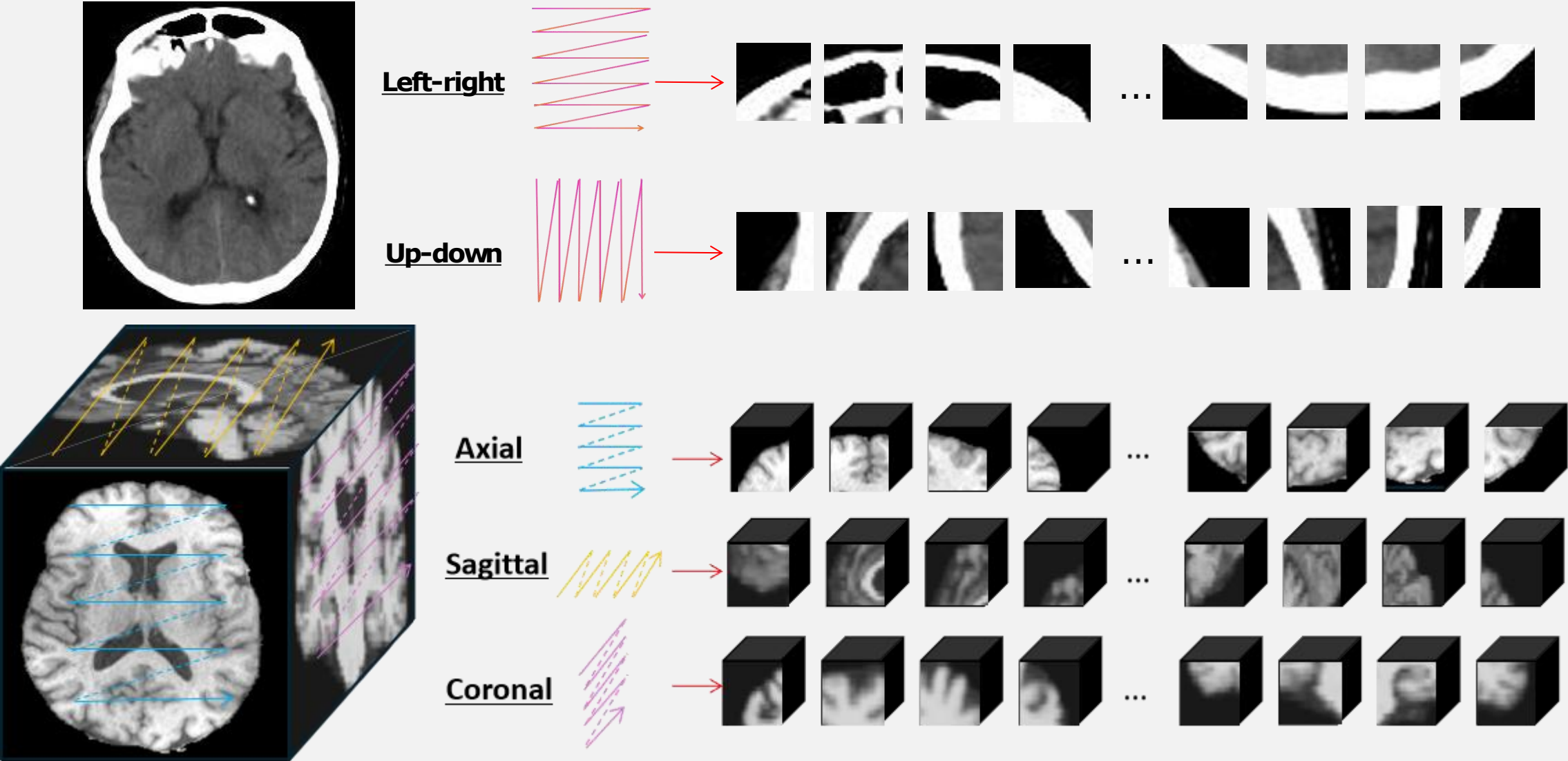


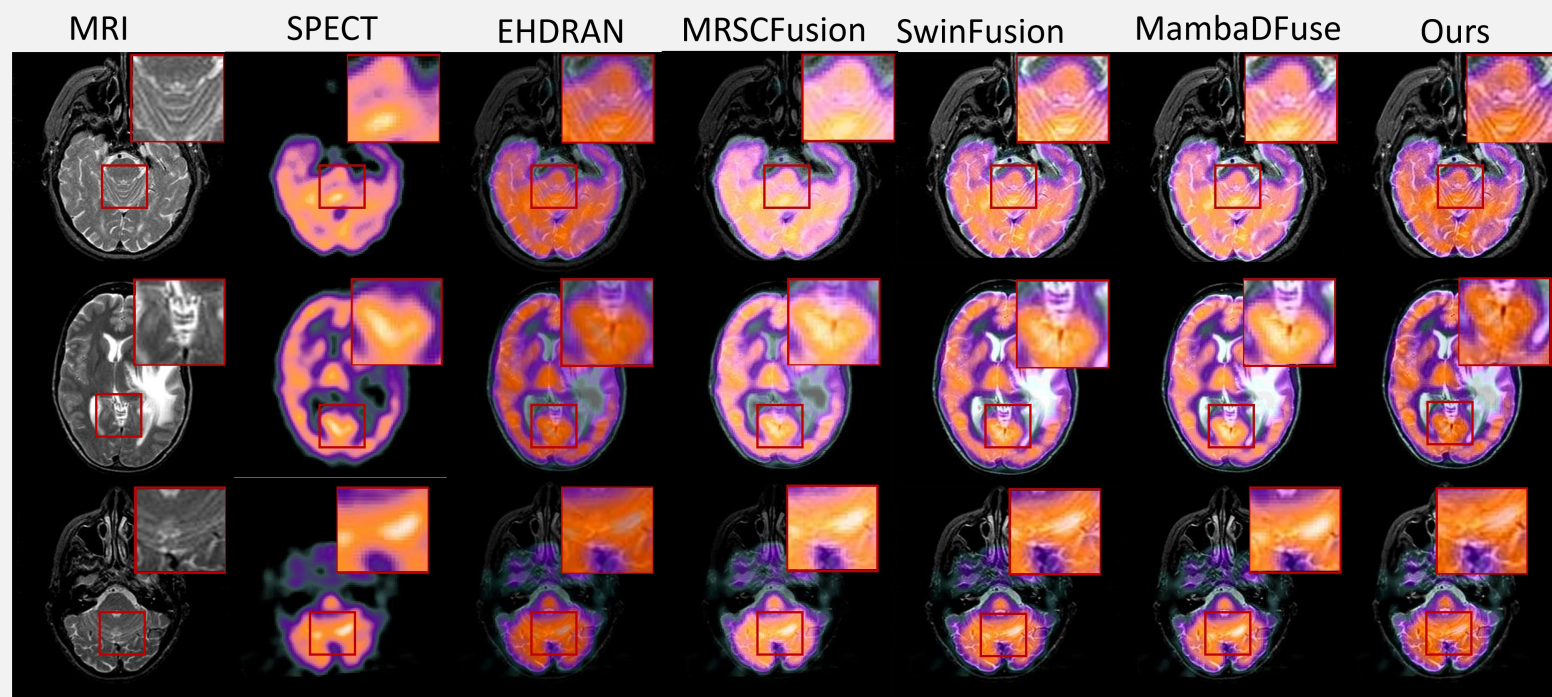
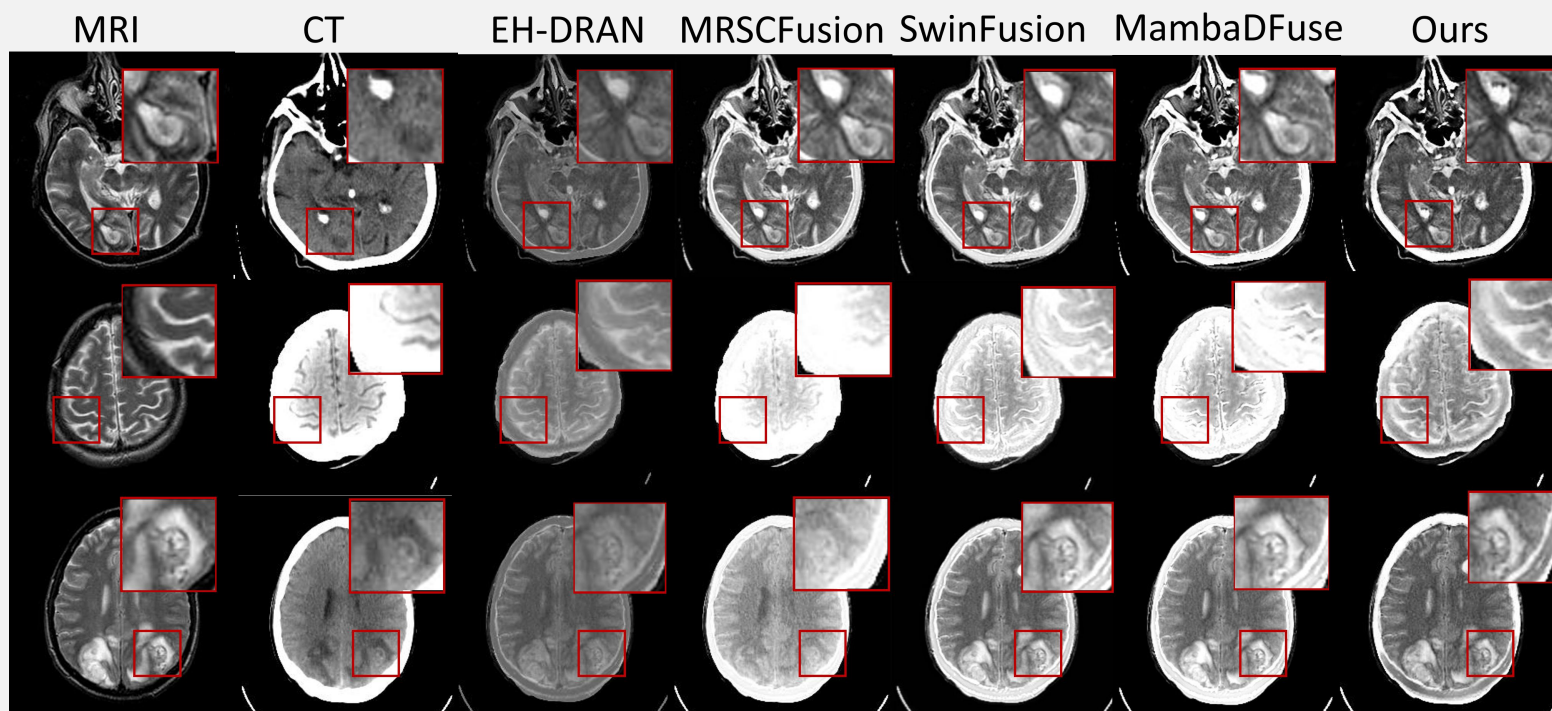
Fig. 1. An overview of the proposed framework. DilatedConv3x3 represents dilated convolution with kernel size 3×3. All Conv+LReLU layers in the decoder have 3×3 convolution followed by Leaky-ReLU. For 3D implementation, all operations are replaced with their corresponding 3D alternatives (i.e., Conv3×3 to Conv3×3×3).

- Pixel loss, ensure appropriate intensity information is retained between fused and input images $L_{pixel} = \|I_f - \max(I_1, I_2)\|_1$
- Gradient loss, preverse fine-grained texture details from input images, $L_{grad} = \|\nabla I_f - \max(\nabla I_1, \nabla I_2)\|_2$
- Similarity loss, preserve structure similarity between fused and input images, $L_{ssim} = \frac{1}{2}(1 - SSIM(I_f, I_1)) + \frac{1}{2}(1 - SSIM(I_f, I_2))$

2D vs. 3D Scan in Mamba



Result



Quantitative Results

Dataset	Method	PSNR $\uparrow$	SSIM $\uparrow$	FMI $\uparrow$	FSIM $\uparrow$	EN $\uparrow$
MRI-CT	EH-DRAN	16.830$\pm$0.490	0.753 $\pm$ 0.007	0.883 $\pm$ 0.005	<u>0.820$\pm$0.003</u>	10.727 $\pm$ 0.531
	SwinFusion	14.962 $\pm$ 0.173	0.768 $\pm$ 0.007	<u>0.882$\pm$0.002</u>	0.810 $\pm$ 0.001	8.445 $\pm$ 0.078
	MRSCFusion	14.476 $\pm$ 0.205	0.713 $\pm$ 0.012	0.877 $\pm$ 0.006	0.791 $\pm$ 0.010	7.544 $\pm$ 0.232
	MambaDFuse	15.873 $\pm$ 0.289	<u>0.771$\pm$0.007</u>	0.882 $\pm$ 0.005	0.817 $\pm$ 0.004	<u>15.018$\pm$0.167</u>
	ClinicalFMamba (Ours)	<u>16.519$\pm$0.352</u>	0.783$\pm$0.005	0.883$\pm$0.003	0.820$\pm$0.001	15.213$\pm$0.069
MRI-SPECT	EH-DRAN	<u>21.455$\pm$0.071</u>	0.736 $\pm$ 0.002	0.876$\pm$0.004	<u>0.843$\pm$0.003</u>	11.970 $\pm$ 0.538
	SwinFusion	17.557 $\pm$ 0.021	0.728 $\pm$ 0.004	0.808 $\pm$ 0.007	0.819 $\pm$ 0.011	13.066 $\pm$ 0.428
	MRSCFusion	18.412 $\pm$ 0.211	0.734 $\pm$ 0.012	0.827 $\pm$ 0.009	0.814 $\pm$ 0.006	9.87 $\pm$ 0.600
	MambaDFuse	21.021 $\pm$ 0.034	<u>0.748$\pm$0.004</u>	0.845 $\pm$ 0.006	0.829 $\pm$ 0.002	<u>14.126$\pm$0.439</u>
	ClinicalFMamba (Ours)	21.561$\pm$0.067	0.759$\pm$0.009	<u>0.856$\pm$0.003</u>	0.848$\pm$0.002	14.871$\pm$0.334

	PSNR $\uparrow$	MS-SSIM $\uparrow$	EN $\uparrow$
EH-DRAN-3D	30.653 $\pm$ 0.554	0.814 $\pm$ 0.095	17.402 $\pm$ 1.134
ClinicalFMamba-3D (Ours)	33.937$\pm$0.361	0.859$\pm$0.045	20.468$\pm$1.541

Fusion Time Analysis

For 2D model on BraTS-2D dataset, our model has 4.05M parameters and achieves an average time of **0.1s** per image pair with 128x128 resolution

For 3D model on BraTS-3D dataset, our model has 6.01M parameters and achieves an average time of **7.3s** per image volume with 128x128x128 resolution

Ablations

MRI-CT	PSNR	SSIM	FMI	FSIM	EN
w/o CMCA	15.967±0.349	0.761±0.007	0.876±0.004	0.813±0.005	14.621±0.075
with CMCA	16.519±0.352	0.783±0.005	0.883±0.003	0.820±0.001	15.213±0.069

BraTS-3D	PSNR	MS-SSIM	EN
w/o 3D-scan	26.882±0.324	0.833±0.073	19.558±1.374
with 3D-can	33.937±0.361	0.859±0.045	20.468±1.541

Quantitative Results on BraTS-classification

Dataset	Method	AUC $\uparrow$	F1-Score $\uparrow$	Accuracy $\uparrow$
BraTS-2D	T2	0.722 ± 0.021	0.703 ± 0.018	0.604 ± 0.037
	FLAIR	0.727 ± 0.024	0.701 ± 0.008	0.611 ± 0.017
	T2+FLAIR	0.723 ± 0.028	0.717 ± 0.012	0.640 ± 0.015
	EH-DRAN	0.769 ± 0.003	0.723 ± 0.006	0.640 ± 0.011
	ClinicalFMamba (Ours)	0.790 ± 0.013	0.778 ± 0.023	0.665 ± 0.004
BraTS-3D	T2-3D	0.647 ± 0.022	0.560 ± 0.029	0.635 ± 0.041
	FLAIR-3D	0.641 ± 0.110	0.529 ± 0.223	0.566 ± 0.010
	T2-3D+FLAIR-3D	0.636 ± 0.072	0.540 ± 0.145	0.630 ± 0.027
	EH-DRAN-3D	0.646 ± 0.015	0.571 ± 0.037	0.657 ± 0.016
	ClinicalFMamba-3D (Ours)	0.652 ± 0.038	0.584 ± 0.023	0.647 ± 0.013

Conclusion

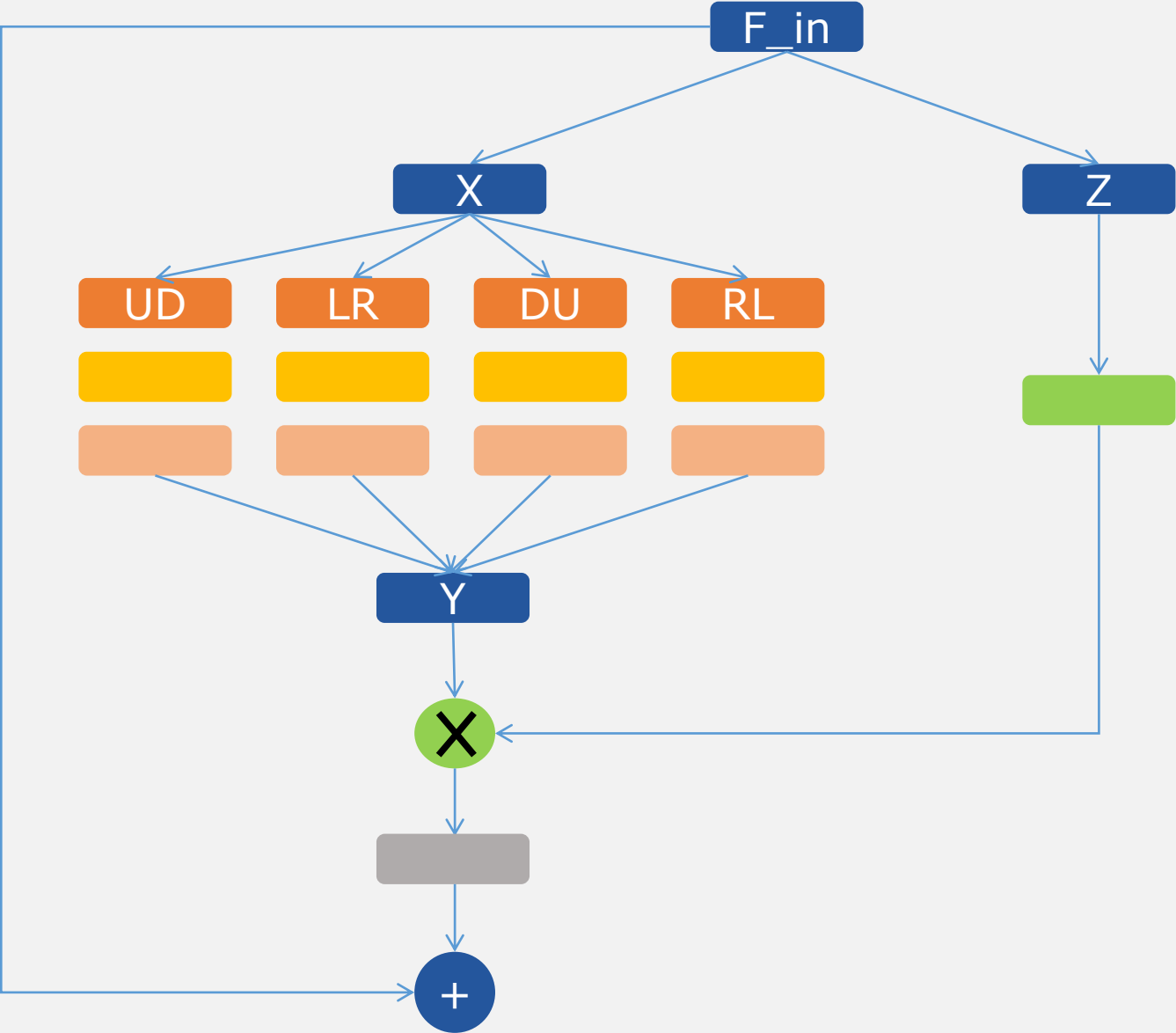
- Novel CNN-Mamba architecture for effective 2D and 3D medical image fusion
 - We propose dilated gated convnets for multiscale feature learning and cross-modal channel attention for cross-modal information fusion and decode.
 - Introduced a tri-plane scanning strategy for 3D medical image fusion.
 - The framework is able to generate fused images in real-time, and we validate the clinical usage of the fused images through brain tumor classification task.
 - Extensive evaluated on three datasets to valid the effectiveness of the proposed approach.
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- Future Work: Explore pure Mamba-based methods and extend to other diseases and downstream tasks like (3D) segmentation.

Thank you!

Arxiv preprint:



Appendix



 flatten  SSM Block  unflatten  SiLU  Conv2D