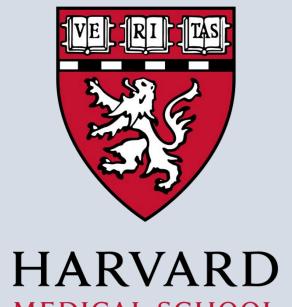
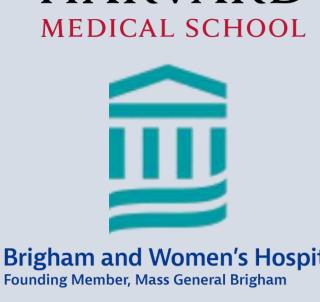


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Medical Image Registration Meets Vision Foundation Model: Prototype Learning and Contour Awareness

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Hao Xu¹, Tengfei Xue¹, Jianan Fan¹, Dongnan Liu¹, Yuqian Chen^{2,4}, Fan Zhang³,

Carl-Fredrik Westin^{2,4}, Ron Kikinis^{2,4}, Lauren J. O'Donnell^{2,4}, Weidong Cai¹

¹ University of Sydney, Australia ² Harvard Medical School, USA ³ UESTC, China ⁴ BWH, USA

Introduction

- Medical image registration methods, which rely solely on intensity-based similarity, often struggle in cases with complex anatomy or ambiguous boundaries.
- Segment Anything Model (SAM) provides high-quality segmentation masks, enabling direct incorporation of structural priors into the registration process.
- We propose a novel SAM-assisted registration framework that incorporates prototype learning and contour awareness to enhance medical image registration.

Methodology

1. SAM Mask Generation

 Segmentation masks are generated for both the fixed and moving images using SAM with text prompts. These masks serve as anatomical priors to guide the registration process.

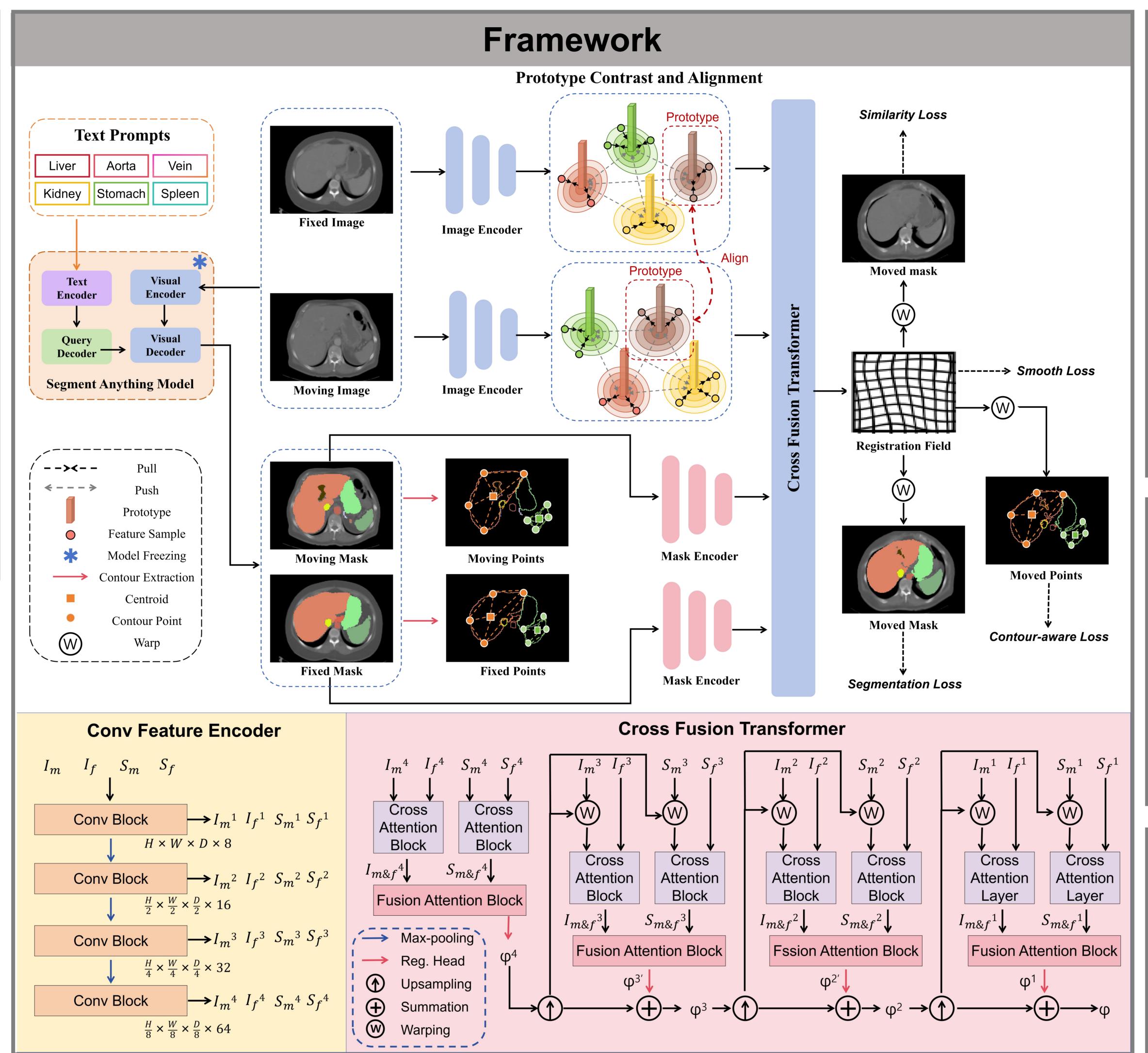
2. Prototype Contrast and Alignment

- Extract feature prototypes of different anatomical regions of the image according to the SAM mask.
- Introduce contrast loss to aggregate features of similar regions and align prototypes of the same region in different images.

3. Contour-Aware Loss

- Extract the contour point sets of the fixed and moving mask.
- Introduce the Chamfer Loss to minimize the Euclidean distance between the fixed and moved point sets.

$$\mathcal{L}_{\text{contour}} = \frac{1}{|C_m|} \sum_{i \in C_m} \min_{j \in C_f} ||i - j||^2 + \frac{1}{|C_f|} \sum_{j \in C_f} \min_{i \in C_m} ||j - i||^2$$



Evaluation on the Abdomen CT dataset

Methods	DSC (%) ↑									SDlogJ ↓				
Wiethous	Spl	Kid R	Kid L	Eso	Liv	Sto	Aor	IVC	Vei	Pan	Adr R	Adr L	AVG	SDIOG3 +
Initial	30.5	23.0	27.2	10.9	50.2	19.2	21.9	20.1	1.9	7.1	3.7	3.8	17.1	<u> </u>
VoxelMorph [1]	61.1	55.3	55.6	30.7	70.1	31.2	44.4	44.0	16.7	19.3	18.3	14.4	38.4 ± 16.6	0.143
TransMorph [4]	60.3	54.3	54.0	30.1	70.7	33.5	45.4	46.1	19.3	18.5	18.2	16.3	39.0 ± 16.2	0.254
TransMatch [5]	65.3	58.4	56.3	33.4	72.3	36.4	49.3	54.5	21.3	20.6	19.9	18.7	42.2 ± 14.7	0.101
CorrMLP [21]	68.2	60.1	63.7	38.4	73.4	45.6	51.2	56.9	24.7	29.1	24.6	24.0	46.7 ± 13.2	0.099
SAM Masks [33]	92.8	88.1	89.5	72.2	95.5	89.4	88.4	84.5	68.5	79.7	64.2	62.6	81.2	-
VoxelMorph [1] + SAM	64.2	61.1	59.6	40.7	70.3	44.4	52.1	53.5	19.2	34.1	24.8	25.9	45.8 ± 12.7	0.065
TransMorph [4] + SAM	71.0	68.7	68.6	49.2	73.1	47.4	66.3	61.1	21.6	38.2	33.9	31.7	52.6 ± 12.2	0.088
TransMatch [5] + SAM	74.0	70.4	69.6	70.2	74.3	54.3	67.1	67.9	32.8	52.9	46.3	47.1	60.5 ± 11.4	0.095
CorrMLP $[21]$ + SAM	77.1	72.8	75.7	55.1	77.7	57.2	76.1	73.1	35.1	50.0	44.6	41.8	61.3 ± 11.3	0.103
Ours	83.9	79.3	76.0	55.2	83.8	82.5	74.5	73.2	36.9	56.4	44.8	48.5	66.3 ± 10.6	0.091

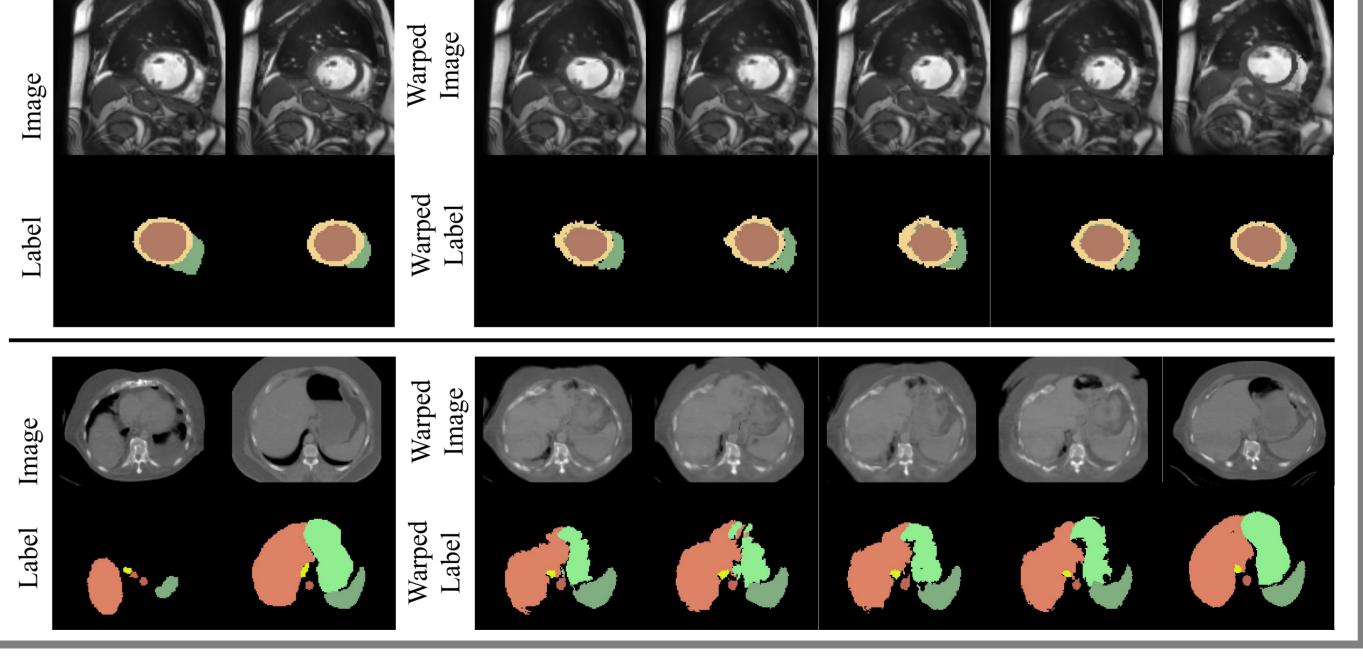
Evaluation on the ACDC MRI dataset

Methods		SDlogJ ↓			
Methods	LV	Myo	RV	AVG	ppioga 1
Initial	58.1	35.8	74.5	56.1	-
VoxelMorph [1]	83.2	60.1	80.5	74.6 ± 7.1	0.041
TransMorph [4]	82.5	58.8	80.4	73.9 ± 7.4	0.031
TransMatch [5]	82.3	58.6	82.1	74.4 ± 6.9	0.037
CorrMLP [21]	82.8	72.9	83.2	79.7 ± 4.6	0.054
SAM Masks [33]	89.2	79.2	76.2	81.6	-
VoxelMorph [1] + SAM					0.077
TransMorph [4] + SAM	86.0	60.7	81.3	76.0 ± 5.5	0.026
TransMatch [5] + SAM	87.6	61.8	82.8	77.5 ± 5.1	0.073
CorrMLP $[21]$ + SAM	83.4	74.2	84.2	80.6 ± 4.2	0.047
Ours	91.9	77.6	83.9	$\textbf{84.6}\pm\textbf{3.7}$	0.049

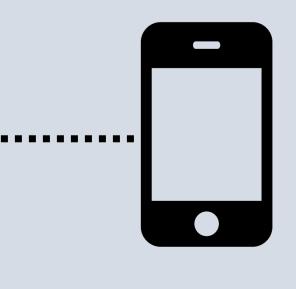
Ablation Study

Dataset	$L_{prototype}$	$L_{contour}$	DSC (%) ↑	$SDlogJ \downarrow$
	X	X	62.8	0.089
Abdomen	✓	X	63.3	0.086
	1	/	66.3	0.091
	X	X	83.0	0.053
ACDC	✓	X	83.5	0.045
	✓	✓	84.6	0.049









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