





上海人工智能实验室 Shanghai Artificial Intelligence Laboratory



# **CAT: Coordinating Anatomical-Textual Prompts for Multi-Organ and Tumor Segmentation**

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# Motivation

- Textual-prompted methods utilize textual representations from referred text phrases to guide the segmentation process. Data scarcity due to long-tailed distribution hinders the effective learning of alignments between textual and visual representations.
- Visual prompts provides a more intuitive and direct method to enhance the segmentation process but fail to convey the general concept of each object, leading to a performance drop when confronted with various scenarios in medical domains, especially for tumors.





The T-stage primarily indicates the extent to which the tumor invades its nearby tissues. A more advanced stage suggests deeper invasion and potentially greater spread within the local area. Colon tumors can present with varying densities and might be associated with surrounding inflammation, adjacent organ invasion, or regional lymph node enlargement.

**Colon Tumor in Different Cancer Stages** 

# Method

- Coordinating Anatomical-Textual Prompts (CAT)
- **Dual-perspective prompting scheme**: employ the cropped volumes derived from the anatomical structure and enhance the textual prompts with more comprehensive knowledge
- ShareRefiner: refine segmentation queries and prompt queries
- **PromptRefer**: updates segmentation queries by integrating both types of prompt queries



#### **Experiments-Organ Segmentation**

Methods	Liv.	R_Kid.	Spl.	Pan.	Aor.	IVC	RAG	LAG	Gal.	Eso.	Sto.	Duo.	L_Kid.	Avg.
SAM 22	86.0	87.6	84.5	53.4	77.5	44.5	19.4	33.9	52.4	35.2	68.0	44.4	82.6	59.2
MedSAM [17]	93.0	90.0	89.1	73.5	82.5	76.5	36.0	48.7	56.4	64.7	84.0	53.9	89.7	72.2
SAM-Med2D [19]	91.4	83.7	83.9	58.8	60.6	18.6	10.6	27.1	32.9	28.1	72.9	45.4	86.0	53.8
SAM-Med3D [40]	85.4	84.2	84.7	46.9	60.4	44.5	32.6	35.3	56.0	32.6	46.9	27.4	84.9	55.5
SegVol [39]	83.9	71.7	75.9	69.4	83.1	80.3	42.1	49.7	55.6	69.6	81.1	55.6	75.1	68.7
CT-SAM3D [41]	95.6	95.0	96.1	83.6	94.5	91.8	78.4	82.5	88.4	82.9	92.3	73.2	94.8	88.4
Universal <sup>†</sup> [13]	97.4	95.5	96.4	73.7	84.9	84.4	72.9	73.4	86.0	76.8	88.5	74.5	96.9	84.7
ZePT* [12]	96.7	95.6	96.6	84.3	90.0	84.4	67.2	66.8	79.6	74.2	85.2	59.1	97.2	82.8
CAT	97.7	96.3	97.1	89.2	90.5	88.0	73.6	74.3	83.0	80.1	88.2	73.4	97.3	86.8

Organ segmentation performance on FLARE22. The results(%) are evaluated by DSC. Liv.-Liver, R\_Kid.-Right Kidney, Spl.-Spleen, Pan.-Pancreas, Aor.-Aorta, IVC-Inferior Vena Cava, RAG-Right Adrenal Gland, LAG-Left Adrenal Gland, Gal.-Gallbladder, Eso.-Esophagus, Sto.-Stomach, Duo.-Duodenum, L\_Kid.-Inferior Vena Cava.





Ground Truth

## **Experiments-Tumor Segmentation**

			MSD D	Dataset (Tu	mor in A	In-house Data (Colon Tumor)								
Method	Liver		Pancreas		Hepatic Vessel		Colon		T1	T2	T3	T4	Av	/g.
	$  DSC\uparrow HD95\downarrow$		DSC↑	HD95↓	$  DSC\uparrow HD95\downarrow$		DSC↑	HD95↓	DSC↑		SC↑		$  DSC\uparrow HD95 \downarrow$	
nnUNet* [62] Swin UNETR* [48]	66.42 68.67	42.29 42.54	43.50 41.77	25.80 22.87	66.90 63.32	47.59 44.02	41.41 39.35	153.06 161.26	19.51   21.40	45.06 33.32	44.87 46.11	45.54 52.72	43.00 45.92	150.48 168.25
SAM-Med3D <sup>†</sup> [40] SegVol <sup>†</sup> [39]	44.78 66.20	-	40.05 46.36	-	44.86 68.57	-	39.23 60.63	-	34.28 36.93	42.65 42.63	50.20 <b>60.17</b>	42.65 49.83	47.11 50.28	-
Universal <sup>†</sup> [13] ZePT <sup>*</sup> [12] CAT	65.68 68.58 <b>72.73</b>	63.31 43.23 <b>34.64</b>	45.72 44.39 <b>49.67</b>	16.58 19.47 <b>15.56</b>	66.31 68.12 <b>70.11</b>	51.47 33.94 <b>33.44</b>	42.26 40.38 48.31	115.40 113.07 <b>108.26</b>	7.11 23.87 30.62	43.28 34.64 <b>45.61</b>	46.52 50.81 55.85	53.08 51.09 <b>57.37</b>	47.14 46.28 <b>53.35</b>	140.28 155.83 <b>80.96</b>

Segmentation performance (%) of tumors on MSD and In-house dataset.





Ground Truth

## **Experiments-Qualitative Comparison**



#### **Experiments-Ablation Study**

	V	<b>Variant</b>			(	Organ (%	)		Tumor (%)					
AP	TP	Hard	Mask	Pan.	an.   RAG   LAG		Eso.	Duo.	Liver   Pancreas		HepVes.	Colon	T4	
				78.18	69.63	69.08	76.99	54.39	66.37	42.05	62.20	39.85	51.17	
$\checkmark$				83.55	72.80	71.65	79.31	60.45	64.82	45.08	68.72	43.84	53.91	
	$\checkmark$			80.62	71.02	70.34	72.81	57.31	69.13	44.31	65.18	40.16	52.32	
$\checkmark$	$\checkmark$			83.50	72.71	69.96	78.99	64.08	72.49	44.55	69.40	44.50	55.84	
	$\checkmark$		$\checkmark$	86.74	72.41	69.00	77.68	59.99	69.12	43.23	67.75	41.32	54.33	
$\checkmark$	$\checkmark$	$\checkmark$		87.36	74.46	74.02	75.39	70.80	72.64	48.49	69.02	47.29	53.67	
$\checkmark$	$\checkmark$		$\checkmark$	88.49	73.24	74.51	80.76	70.26	72.18	46.46	69.97	46.65	58.49	
$\checkmark$	$\checkmark$	$\checkmark$	$\checkmark$	88.28	74.42	72.50	79.30	71.26	70.95	45.52	69.51	46.07	56.41	
$\checkmark$	✓	✓	✓	89.24	73.69	74.63	80.10	73.46	72.73	49.67	70.11	48.31	57.37	

#### **Experiments-Visualization**



T-SNE visualization of the distribution of Features and Heatmaps.

#### Conclusion

- A promising attempt towards comprehensive medical segmentation via coordinating anatomicaltextual prompts.
- Apart from performing generic organ segmentation, CAT can identify varying tumors without human interaction.
- To effectively integrate two prompt modalities into a single model, we design ShareRefiner to refine latent prompt queries with different strategies and introduce PromptRefer with specific attention masks to assign prompts to segmentation queries for mask prediction.
- Extensive experiments indicate that the coordination of these two prompt modalities yields competitive performance on organ and tumor segmentation benchmarks. Further studies revealed the robust generalization capabilities to segment tumors in different cancer stages.

# **Thanks for Listening !**

Code Link: <u>https://github.com/zongzi3zz/CAT</u>

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