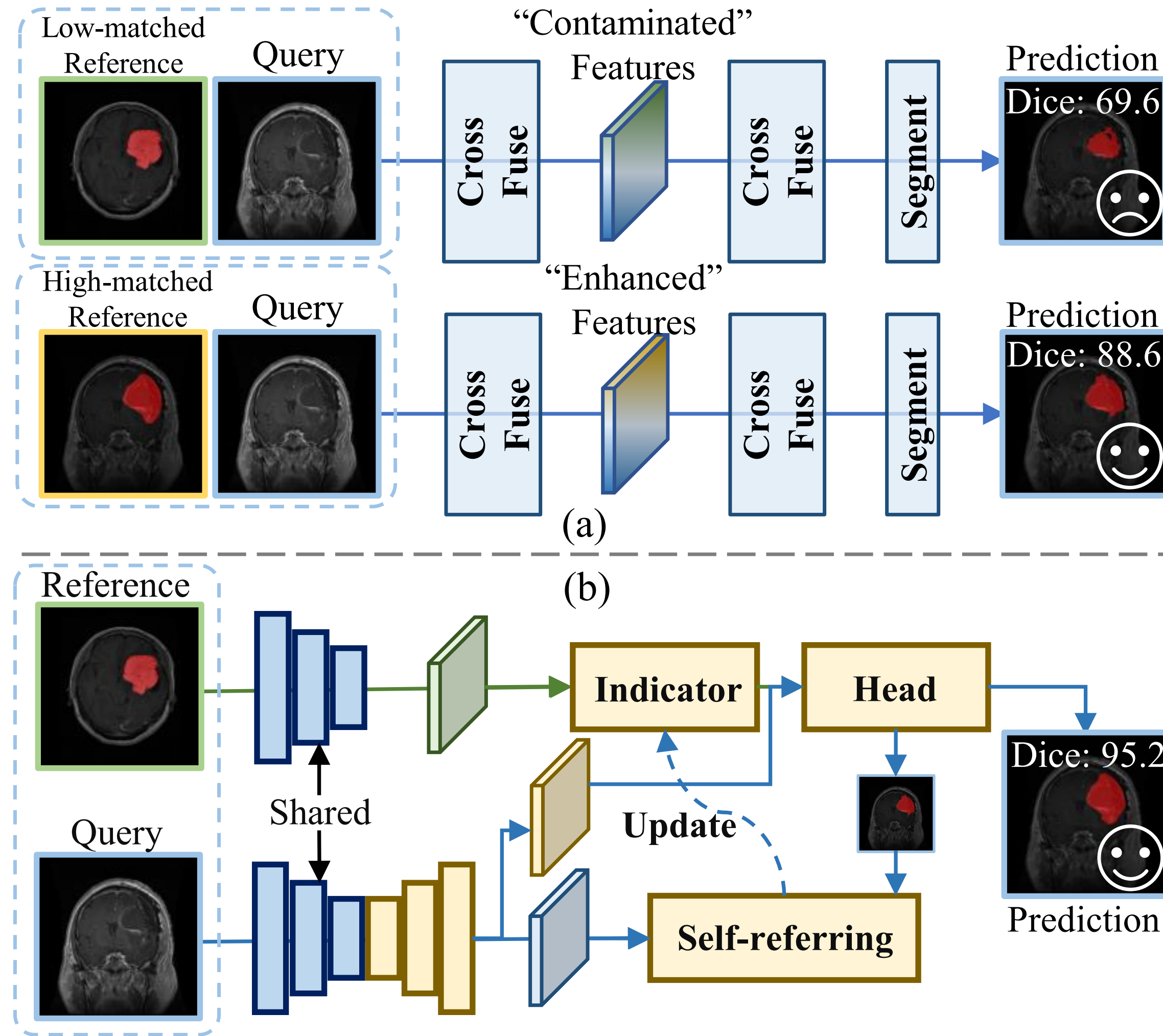
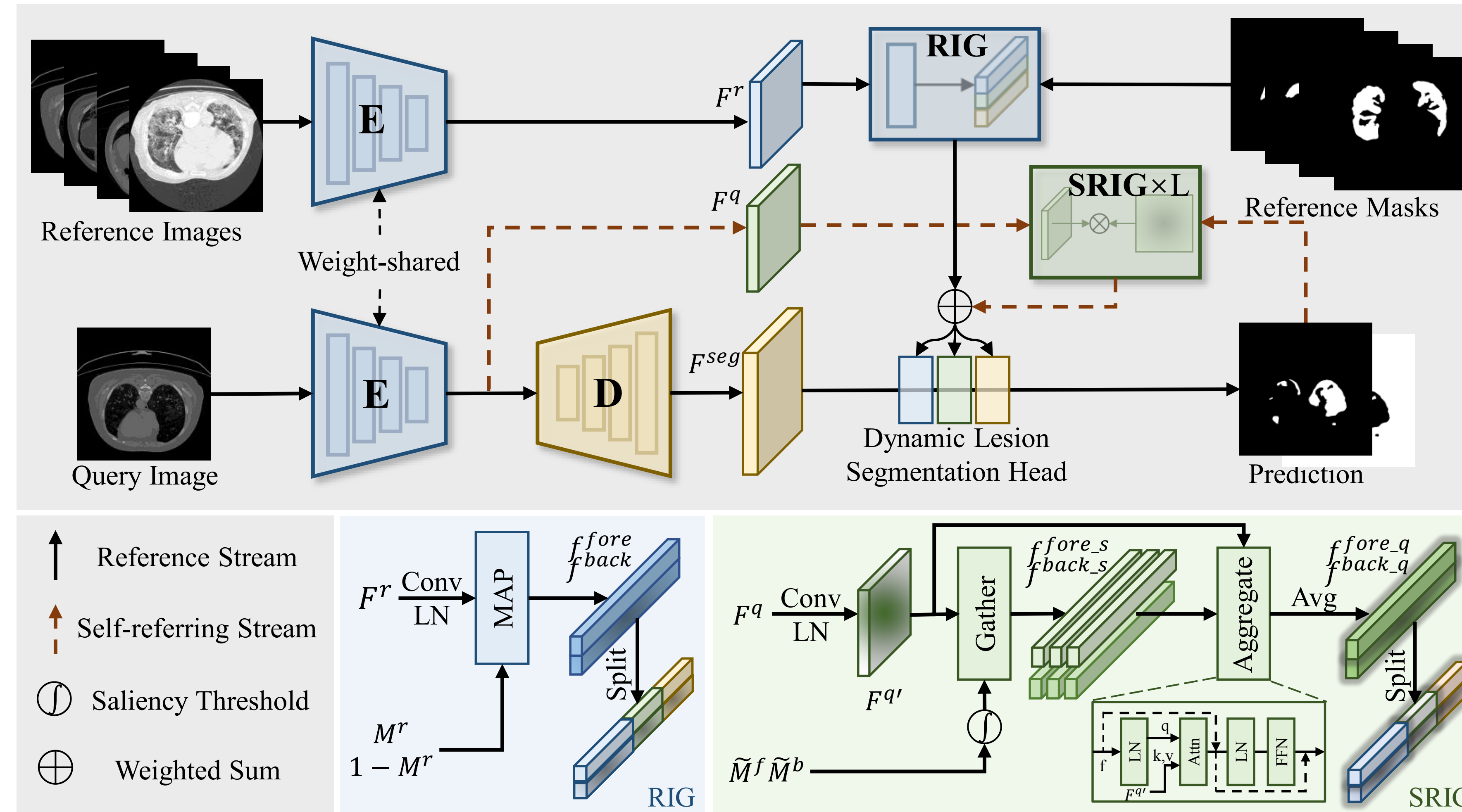


## Motivation/Contributions



## Method: SR-ICL

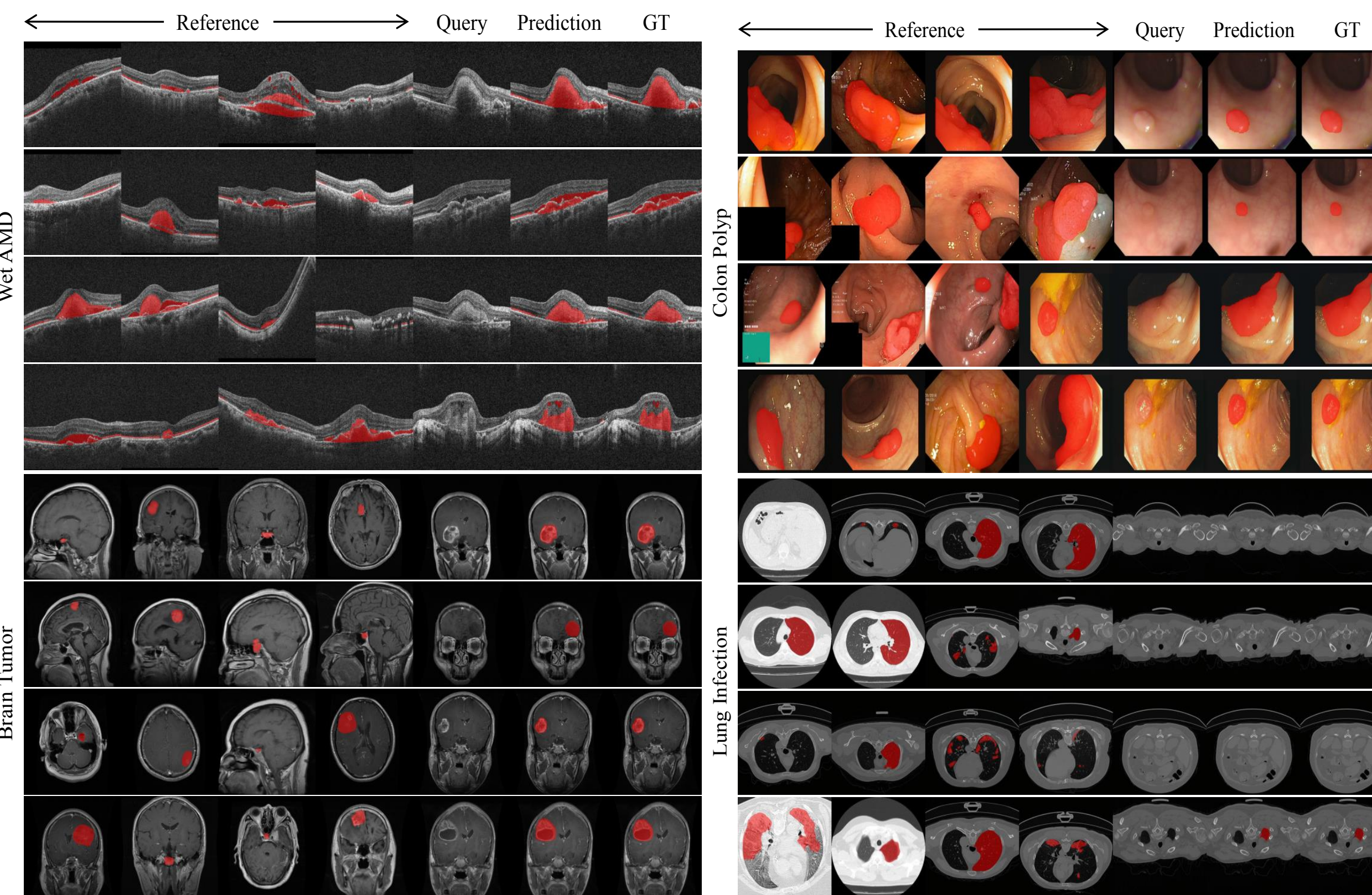


**Algorithm 1** "Separate BP - Average Update"

Training iteration with  $N = 12$  and  $B = 1$ .

**Require:** A batch  $D = \{D_t\}_{t=1}^N$  and  $Q = \{Q_t\}_{t=1}^N$ .  $D_t$  and  $Q_t$  are  $N$  and  $B$  image-mask pairs randomly selected from training set of task  $t$ .

- 1: create reference image tensor  $I_r \in \mathbb{R}^{8 \times N \times 3 \times H \times W}$  from  $D$
- 2: create reference mask tensor  $M_r \in \mathbb{R}^{8 \times N \times 1 \times H \times W}$  from  $D$
- 3: create query image tensor  $I_q \in \mathbb{R}^{8 \times B \times 3 \times H \times W}$  from  $Q$
- 4: create query mask tensor  $M_q \in \mathbb{R}^{8 \times B \times 1 \times H \times W}$  from  $Q$
- 5:  $avg\_grads \leftarrow []$  // empty list
- 6: **for** each  $param$  in  $SR-ICL.parameters()$  **do**
- 7:  $zero\_tensor \leftarrow torch.zeros\_like(param)$
- 8:  $avg\_grads.append(zero\_tensor)$
- 9: **end for**
- 10:  $L \leftarrow 0$
- 11: **for**  $t \leftarrow 1, 8$  **do**
- 12:  $R_t^I \leftarrow I_r[t, ...]$  // reference images
- 13:  $R_t^M \leftarrow M_r[t, ...]$  // reference masks
- 14:  $P_t^I \leftarrow I_q[t, ...]$  // query images
- 15:  $P_t^M \leftarrow M_q[t, ...]$  // query masks
- 16:  $P_t \leftarrow SR-ICL(R_t^I, R_t^M, P_t^I)$  // get prediction
- 17:  $L \leftarrow Loss(P_t, P_t^M)$
- 18:  $zero\_grad(Optimizer)$
- 19:  $backward(L)$  // get gradients
- 20: **for** each  $(avg\_grad, param)$  **in**  $(avg\_grads, SR-ICL.parameters())$  **do**
- 21:  $avg\_grad \leftarrow avg\_grad + param.grad$
- 22: **end for**
- 23: **end for**
- 24: **for** each  $(param, avg\_grad)$  **in**  $(SR-ICL.parameters(), avg\_grads)$  **do**
- 25:  $param.grad \leftarrow (avg\_grad)$  // average gradients
- 26: **end for**
- 27:  $step(Optimizer)$  // update parameters



## Results:

Methods	Methods	Wet AMD Dice mIoU	Brain Tumor Dice mIoU	Adenocarcinoma Dice mIoU	Thyroid Nodule Dice mIoU	Polyp Dice mIoU	Lung Infection Dice mIoU	Breast Lesion Dice mIoU	Skin Lesion Dice mIoU
<b>Specialized Models (One model for one task)</b>									
TRSRD-Net [2]	ICISN24	-	-	-	85.40	84.65	-	-	-
LDNet [57]	MICCAI22	-	-	-	-	-	64.25	74.41	-
WeakPolyp [52]	MICCAI23	-	-	-	-	-	74.90	80.66	-
Inf-Net [16]	TMI20	-	-	-	-	-	43.24	52.85	-
DECOR-Net [19]	ISBI23	-	-	-	-	-	40.25	69.49	-
AAU-Net [8]	TMI22	-	-	-	-	-	-	-	47.45 65.15
CMU-Net [44]	ISBI23	-	-	-	-	-	-	-	54.52 83.02
MALUNet [36]	BIBM22	-	-	-	-	-	-	-	86.32 85.37
EGE-UNet [37]	MICCAI23	-	-	-	-	-	-	-	85.88 84.98
UNet [35]	MICCAI15	76.01	80.88	60.44	73.79	91.94	75.33	80.16	83.94
TransUNet [9]	MIA24	76.72	81.29	57.92	72.46	91.85	74.23	78.24	82.90
RollingUNet [27]	AAAI24	77.13	81.54	61.38	74.71	92.26	75.05	78.96	83.17
<b>Generalist Models (One model performs new tasks without fine-tuning)</b>									
SegGPT [50]	ICCV23	39.42	62.68	24.43	58.61	73.60	61.71	21.33	57.64
UniverSeg [5]	CVPR23	45.90	63.66	32.83	61.29	80.85	45.66	60.95	73.63
DSC-ICL [18]	TMI24	-	-	-	-	-	-	76.03	-
<b>Unified Models (One model for all tasks)</b>									
UNet [35]	MICCAI15	75.29	80.46	54.95	70.35	91.21	72.66	78.41	82.88
TransUNet [9]	MIA24	75.46	80.43	56.59	71.77	90.34	69.16	78.57	83.21
RollingUNet [27]	AAAI24	76.46	81.11	60.48	74.43	92.17	71.62	82.26	85.24
SegGPT [50]†	ICCV23	71.66	78.64	44.17	67.30	94.24	80.14	79.69	85.18
Spider [61]†	ICML24	78.51	82.66	71.59	79.86	93.65	80.30	86.52	88.40
Ours	-	<b>80.54</b>	<b>83.84</b>	<b>74.29</b>	<b>81.50</b>	<b>94.96</b>	<b>81.98</b>	<b>87.91</b>	<b>89.18</b>
								<b>83.26</b>	<b>86.53</b>
								<b>82.36</b>	<b>87.21</b>
								<b>84.92</b>	<b>87.08</b>
								<b>90.85</b>	<b>87.29</b>

Segmentation Task	Dataset	Modality	#Train	#Val
Wet AMD	AMD-SD [20]	OCT	2346	703
Brain Tumor	BT [10, 11]	MR-T1	2298	766
Adenocarcinoma	EBHI-Seg [39]	Pathology image	636	159
Thyroid Nodule	TNUI 2021 [63]	Ultrasound	966	276
Colon Polyp	Five datasets [4, 23, 40, 43, 47]	Endoscopy image	1450	798
Lung Infection	COVID-19 data [16]	CT	894	383
Breast Lesion	BUSI [1]	Ultrasound	486	161
Skin Lesion	ISIC 2018 [13]	Dermoscopy image	1886	808

Method	Task Number	Batchsize	Memory
Balance FP - Unify BP	8	6/2	16.8G
Balance FP - Unify BP	8	12/4	OOM
Ours	8	6/2	5.7G
Ours	8	12/4	8.3G
Ours	16	12/4	8.3G

## Conclusion

- The self-referring mechanism improves the robustness to the reference information.
- SR-ICL achieves good performance with weak annotations.
- The training strategy enables SR-ICL to balance any number of datasets even with one single RTX 3090 GPU.

## Future Work

- Dynamic selection of the reference images.
- Large-scale pre-training on medical data.
- Extend to 3D medical segmentation dataset.

- We propose a self-referring mechanism that adaptively extracts information from the query itself to overcome the negative impact caused by low-matched reference sets.
- We design reference indicator generation to efficiently utilize reference information instead of cross-fusion mechanisms which heavily rely on reference sets, facilitating the self-referring mechanism.
- Our designs successfully apply ICL to unified MLS. SR-ICL has impressive performance even with weak reference annotations such as boxes and points. Furthermore, it has relatively low memory requirements during training.