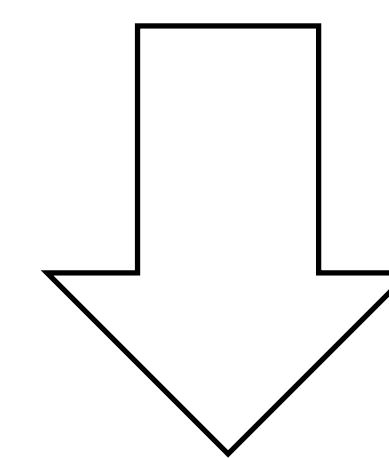
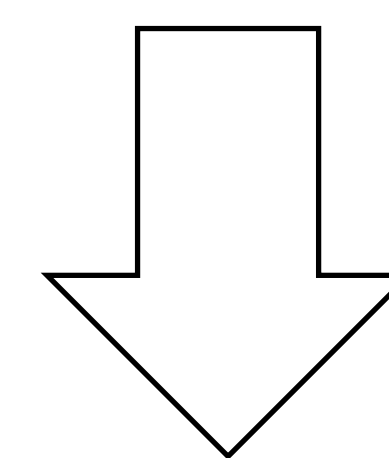


## Introduction

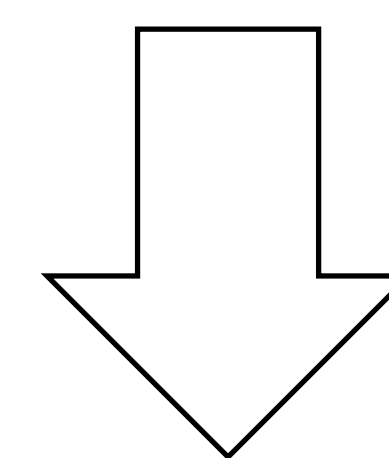
Recently, FixMatch [1] utilizes the confidence-based threshold to select more accurate pseudo-labels and proves the superiority of this technique.



We try to ask — *is the confidence-based threshold really necessary for pseudo-labeling?*



We consider only using distribution alignment (DA) to improve the pseudo-labels without additional hyperparameters.

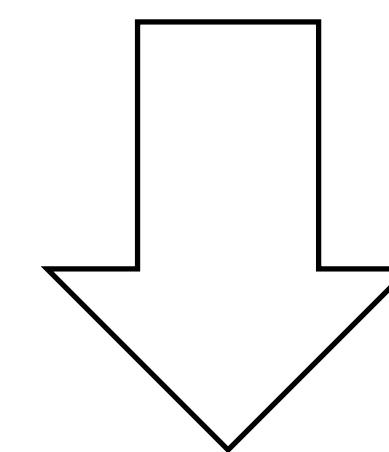


However, original DA is based on a strong assumption: **“labeled data and unlabeled data share the same distribution”**.

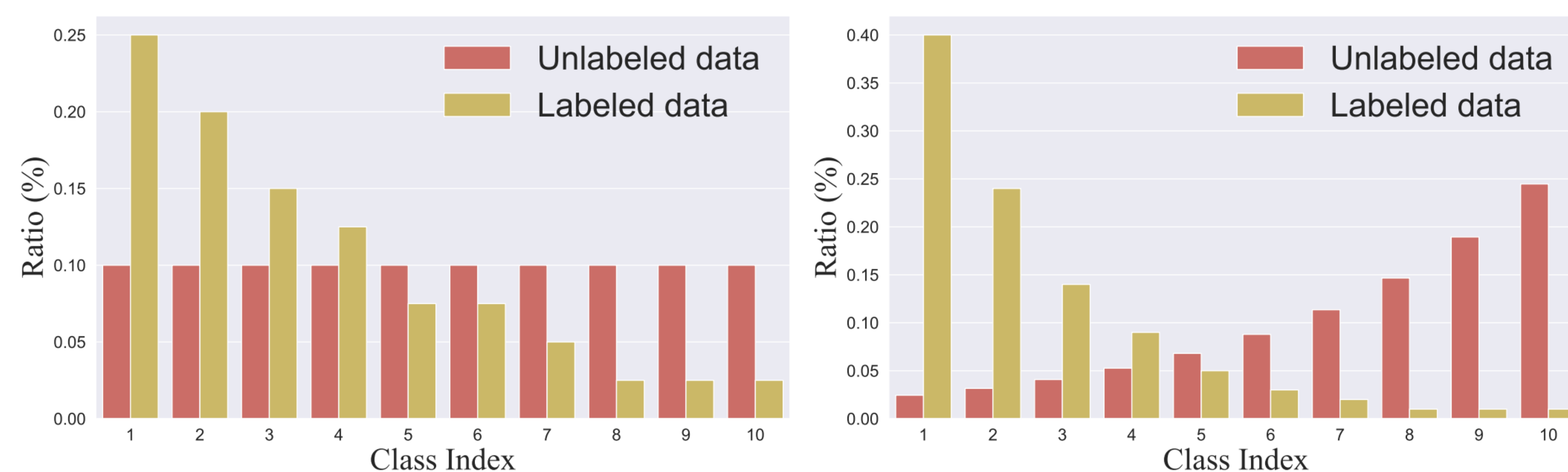


## Motivation

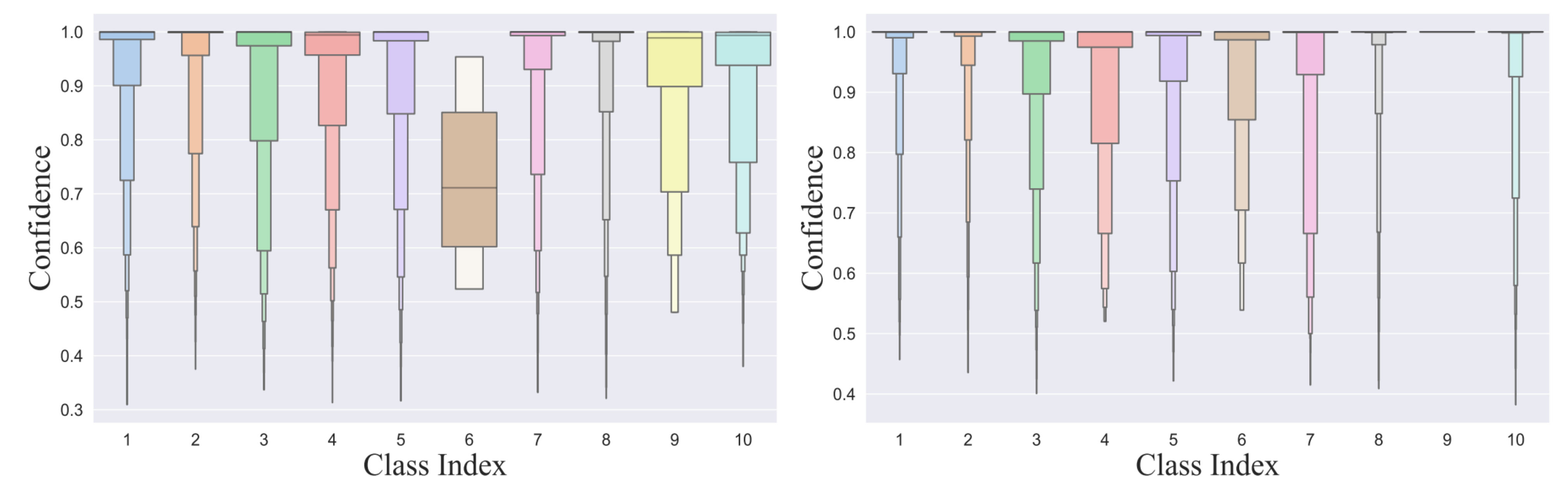
The original distribution alignment technique fails in the SSL with mismatched distribution, while the confidence threshold is difficult to set.



Explore a more general distribution alignment technique to address the challenges of mismatched distributions.



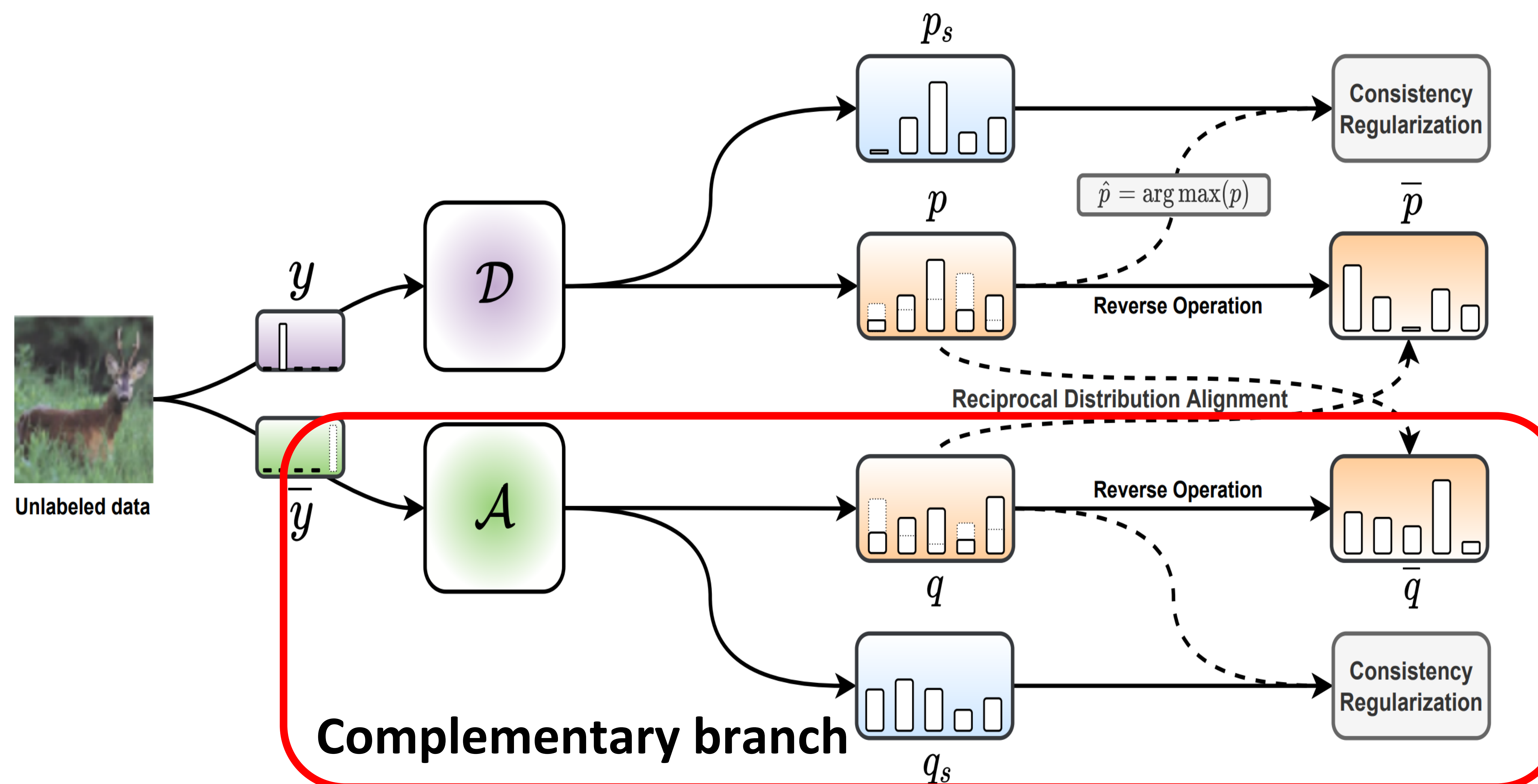
Mismatched distributions



Confidence threshold is difficult to set



A new distribution alignment technique based on **pseudo-label** and **complementary label** distribution is proposed to improve pseudo-label quality



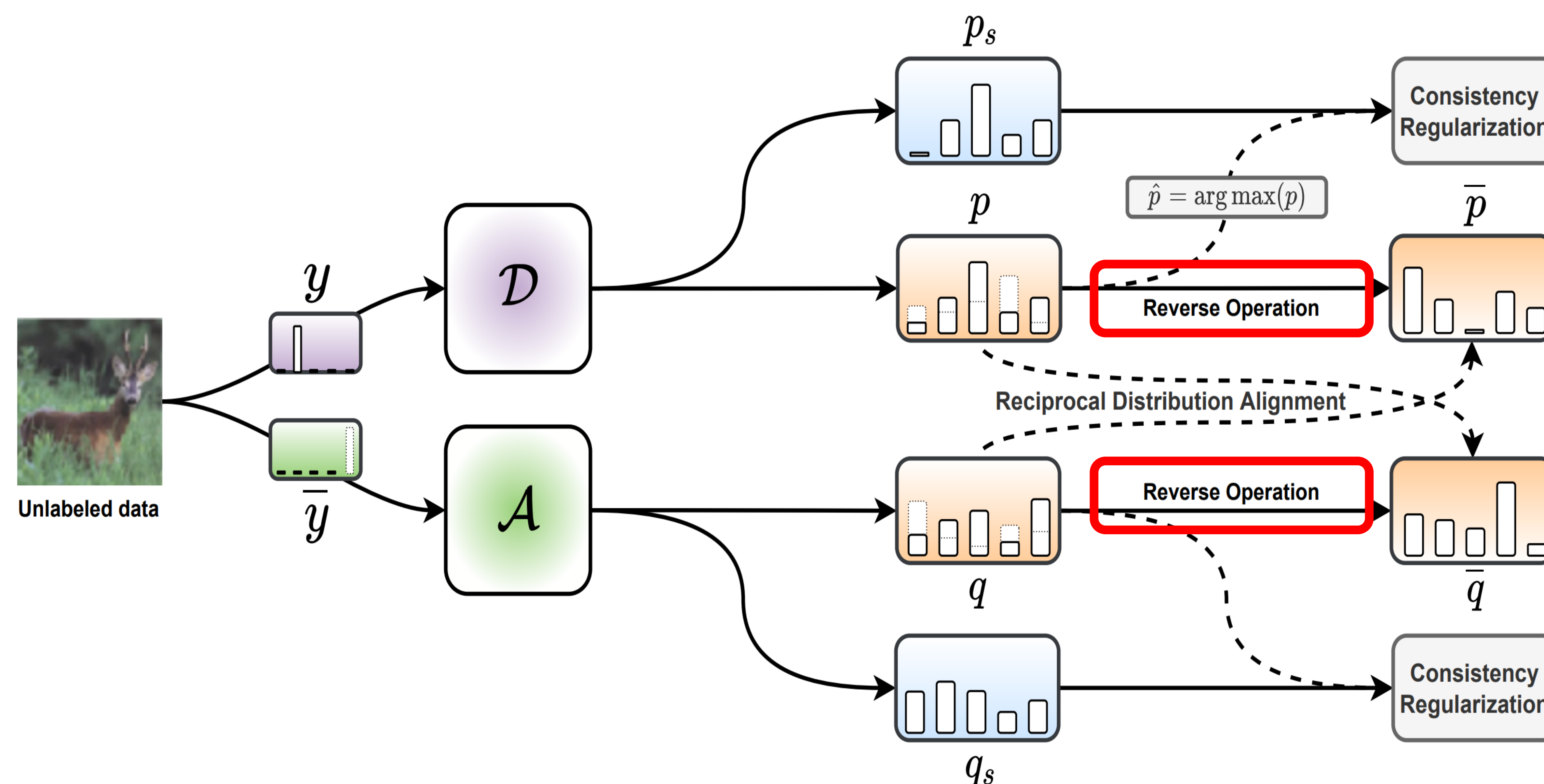
$$\mathcal{I}(y; x) = \mathcal{H}(\mathbb{E}_x[P(y|x)]) - \mathbb{E}_x[\mathcal{H}(P(y|x))]$$

Improve pseudo-labeling by maximizing input-output mutual information [2]

**How to maximize input-output mutual information?**



# RDA: Reciprocal Distribution Alignment for Robust Semi-supervised Learning



## Proposition 1 (Reverse Operation).

In the case of using  $\mathcal{A}$  to predict pseudo-labels, we have  $\bar{q} = \text{Norm}(\mathbb{1} - q)$ , where  $\mathbb{1}$  is all-one vector and  $\text{Norm}(x)$  is the normalized operation.

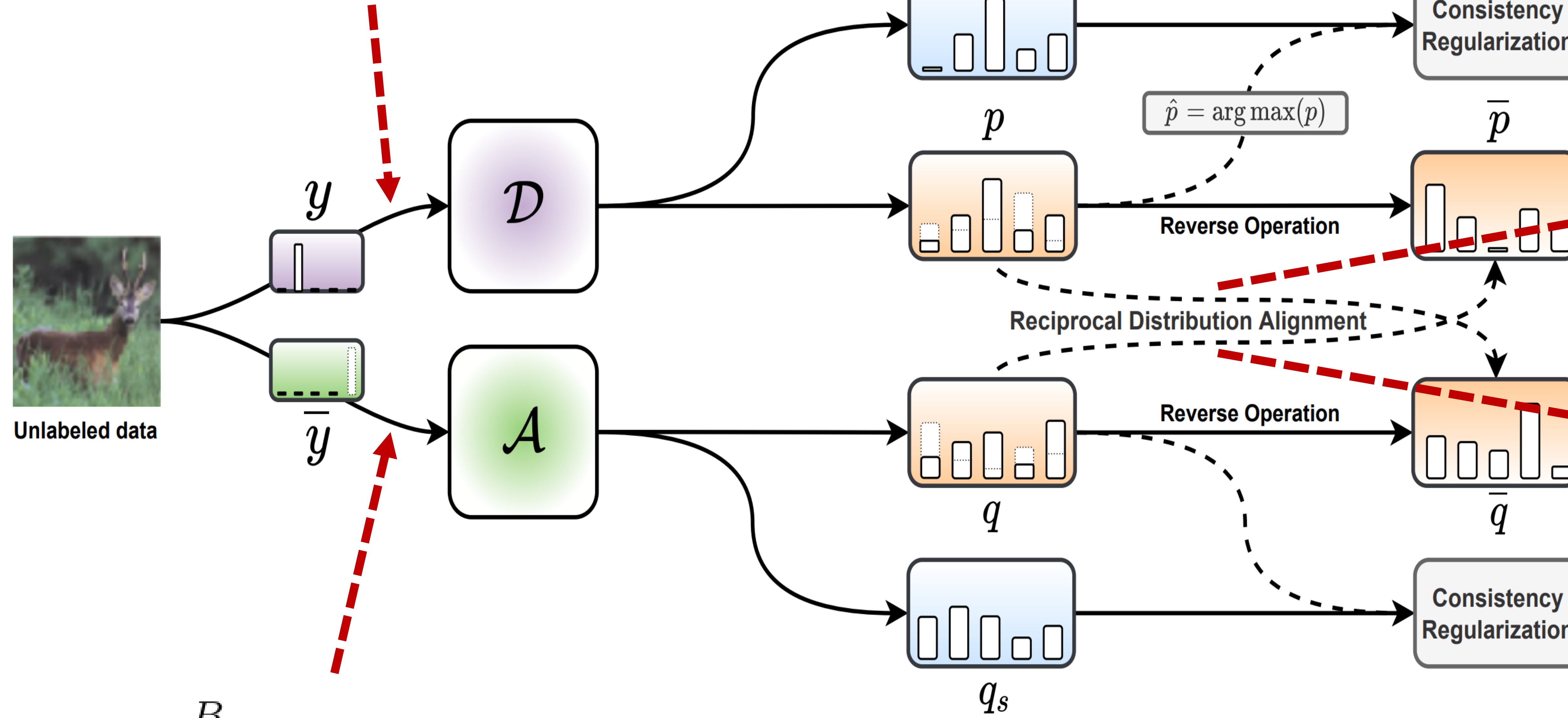
- ◆ Generate complementary labels from labeled data to train Auxiliary Classifier  $\mathcal{A}$
- ◆ Reverse the pseudo-labels and complementary labels output by the Default Classifier and Auxiliary Classifier respectively (**Reverse Operation**)
- ◆ Align the pseudo-label distribution from Default Classifier to the reversed distribution of complementary labels from Auxiliary Classifier, while aligning the complementary label distribution to the reversed distribution of pseudo-labels



# RDA: Reciprocal Distribution Alignment for Robust Semi-supervised Learning



$$\mathcal{L}_{sd} = \frac{1}{B} \sum_{n=1}^B H(y_n, P_{\mathcal{D}}(y_c | x_{w,n})),$$



$$\mathcal{L}_{cd} = \frac{1}{\mu B} \sum_{n=1}^{\mu B} H(\hat{p}_n, p_{s,n}),$$

$$\tilde{p} = \text{Norm}(p \times \frac{\Psi(\bar{q})}{\Psi(p)}),$$

$$\tilde{q} = \text{Norm}(q \times \frac{\Psi(\bar{p})}{\Psi(q)}),$$

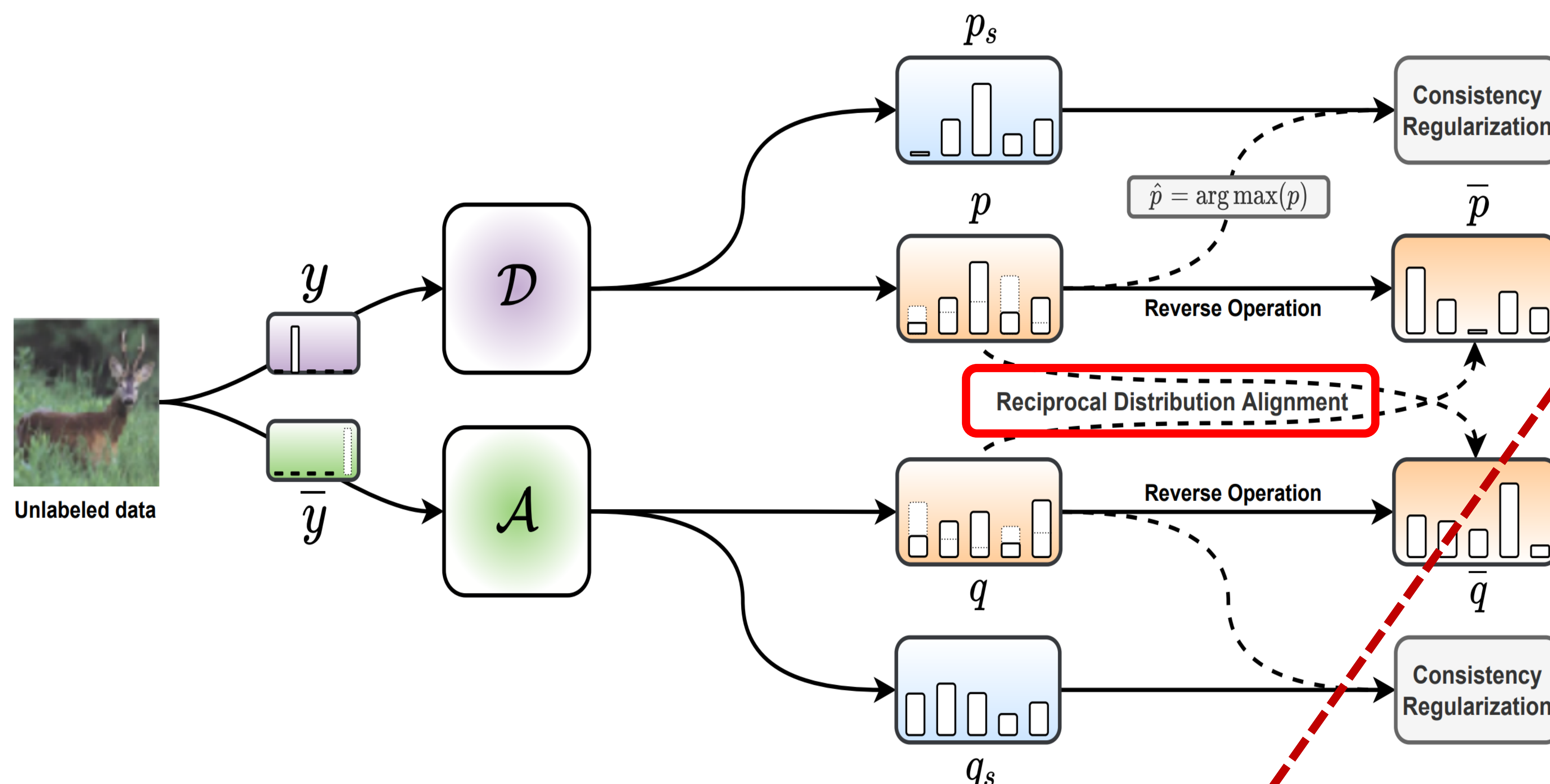
$$\mathcal{L}_{sa} = \frac{1}{B} \sum_{n=1}^B H(\bar{y}_n, P_{\mathcal{A}}(y_c | x_{w,n})),$$

$$\mathcal{L}_{ca} = \frac{1}{\mu B} \sum_{n=1}^{\mu B} H(q_n, q_{s,n}),$$

$$\mathcal{L} = \mathcal{L}_{sd} + \lambda_a \mathcal{L}_{sa} + \lambda_{cd} \mathcal{L}_{cd} + \lambda_{ca} \mathcal{L}_{ca},$$



# RDA: Reciprocal Distribution Alignment for Robust Semi-supervised Learning



**Theorem 1.** For pseudo-label  $p$  and the reversed pseudo-label  $\bar{p}$  obtained by *Reverse Operation*, we show that the entropy of  $\bar{p}$  is larger than that of  $p$ :  $\mathcal{H}(\bar{p}) \geq \mathcal{H}(p)$ , where  $\mathcal{H}(\cdot)$  refers to the entropy.

$$\max_{\mathcal{D}, \mathcal{A}} h(\mathcal{D}, \mathcal{A}) = \mathcal{H}[\mathbb{E}_u(p)] + \mathcal{H}[\mathbb{E}_u(q)] \implies \mathcal{H}[\mathbb{E}_u(p)] + \mathcal{H}[\mathbb{E}_u(q)] \leq \mathcal{H}[\mathbb{E}_u(\bar{p})] + \mathcal{H}[\mathbb{E}_u(\bar{q})].$$

We show theoretically that the input-output mutual information can be maximized by *reciprocal distribution alignment*



## ◆ Conventional SSL setting: CIFAR10, mini-ImageNet, STL-10

| Method      | CIFAR-10          |                   |                   |                   | mini-ImageNet     | STL-10            |
|-------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|
|             | 20 labels         | 40 labels         | 80 labels         | 100 labels        | 1000 labels       | 1000 labels       |
| MixMatch*   | 27.84±10.63       | 51.90±11.76       | 80.79±1.28        | -                 | -                 | 38.02±8.29        |
| AlphaMatch† | -                 | 91.35±3.38        | -                 | -                 | -                 | -                 |
| FixMatch    | 84.97±10.37       | 89.18±1.54        | 91.99±0.71        | 93.14±0.76        | 39.03±0.66        | 65.38±0.42*       |
| CoMatch     | 88.43±7.22        | 93.21±1.55        | 94.08±0.31        | <b>94.55±0.27</b> | 43.72±0.58        | 79.80±0.38*       |
| RDA         | <b>92.03±2.01</b> | <b>94.13±1.22</b> | <b>94.24±0.42</b> | <b>94.35±0.25</b> | <b>46.91±1.16</b> | <b>82.63±0.54</b> |

In the conventional SSL setting, where the labeled and unlabeled data have the same distribution and are uniformly distributed, RDA achieves superior performance.

## ◆ Mismatched distribution scenarios: CIFAR-10/100, mini-ImageNet

- The labeled data is imbalanced, the unlabeled data is balanced
- The labeled data is balanced, the unlabeled data is imbalanced
- The labeled data and unlabeled data are imbalanced and mismatched



# RDA: Reciprocal Distribution Alignment for Robust Semi-supervised Learning



| Method         | CIFAR-10          |                   |                   |                   | CIFAR-100         |                   | mini-ImageNet     |                   |
|----------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|-------------------|
|                | 40 labels         |                   | 100 labels        |                   | 400 labels        | 1000 labels       | 1000 labels       |                   |
|                | $N_0 = 10$        | 20                | 40                | 80                | 40                | 80                | 40                | 80                |
| FixMatch       | 85.72±0.93        | 76.53±3.03        | 93.01±0.72        | 71.57±1.88        | 25.66±0.46        | 40.22±1.00        | 36.20±0.36        | 28.33±0.41        |
| FixMatch w. DA | 71.23±1.25        | 47.85±1.99        | 56.78±1.28        | 34.18±0.86        | 22.66±1.53        | 31.06±0.51        | 33.87±0.40        | 23.53±0.72        |
| CoMatch        | 60.27±3.22        | 39.48±2.20        | 52.82±2.03        | 26.91±0.75        | 23.97±0.62        | 28.35±1.20        | 30.24±1.37        | 21.47±0.86        |
| RDA            | <b>92.57±0.53</b> | <b>81.78±6.44</b> | <b>94.23±0.36</b> | <b>79.00±2.67</b> | <b>30.86±0.78</b> | <b>41.29±0.43</b> | <b>42.73±0.84</b> | <b>36.73±1.01</b> |

| Method | CIFAR-10              |                   |                        |                   | mini-ImageNet           |
|--------|-----------------------|-------------------|------------------------|-------------------|-------------------------|
|        | 40 labels, $N_0 = 10$ |                   | 100 labels, $N_0 = 40$ |                   | 1000 labels, $N_0 = 40$ |
|        | $\gamma = 2$          | 5                 | 5                      | 10                | 10                      |
|        | FixMatch              | 74.97±5.80        | 64.62±6.13             | 58.72±3.61        | 57.49±4.56              |
| RDA    | <b>88.58±4.05</b>     | <b>79.90±2.80</b> | <b>79.33±1.37</b>      | <b>70.93±2.91</b> | <b>25.99±0.19</b>       |

| Method   | CIFAR-10 ( $\gamma_l = 100$ ) |                   |                   |                   | STL-10 ( $\gamma_l \neq \gamma_u$ ) |                   |
|----------|-------------------------------|-------------------|-------------------|-------------------|-------------------------------------|-------------------|
|          | $\gamma_u = 1$                | 50                | 150               | 100 (reversed)    | $\gamma_l = 10$                     | 20                |
| FixMatch | 68.90±1.95                    | 73.90±0.25        | 69.60±0.60        | 65.50±0.05        | 72.90±0.09                          | 63.40±0.21        |
| DARP     | 85.40±0.55                    | 77.30±0.17        | 72.90±0.24        | 74.90±0.51        | 77.80±0.33                          | 69.90±0.40        |
| RDA      | <b>93.35±0.24</b>             | <b>79.77±0.06</b> | <b>74.48±0.24</b> | <b>79.25±0.52</b> | <b>87.21±0.44</b>                   | <b>83.21±0.52</b> |

In the mismatched scenario, RDA still achieves a superior performance advantage.