



# Towards Realistic Long-Tailed Semi-Supervised Learning: Consistency Is All You Need

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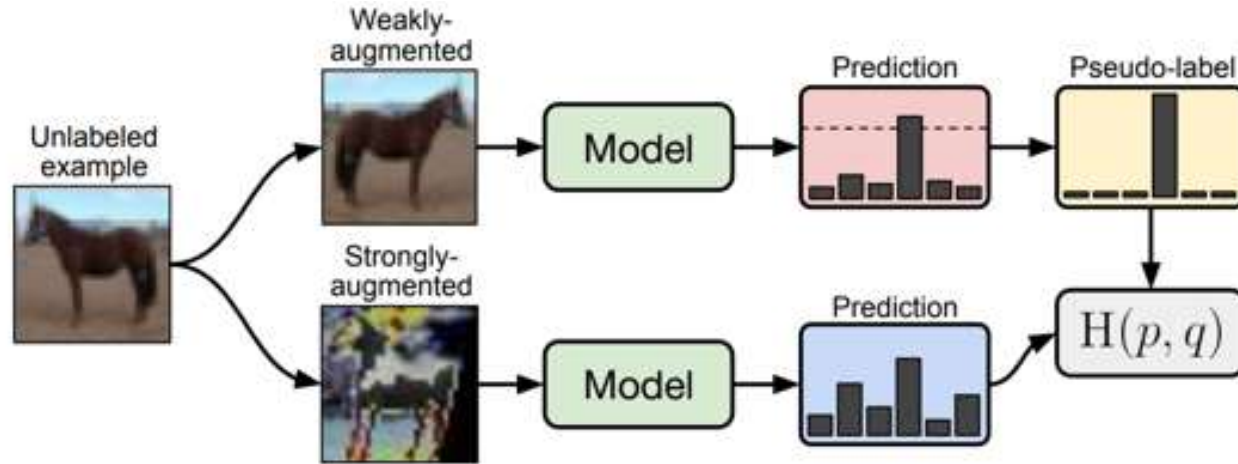
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# Long-Tailed Semi-Supervised Learning (LTSSL)

Common semi-supervised learning method: FixMatch



Labeled(Cross-entropy)  
 +  
 Unlabeled(Consistency regularization)

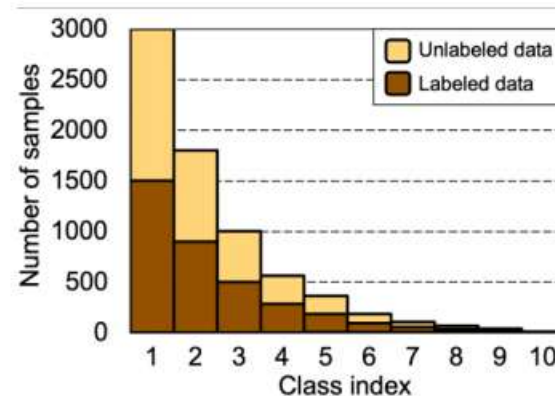
$$\min_{\theta \in \Theta} \underbrace{\sum_{i=1}^N \ell(f(x_i^{(l)}; \theta), y_i^{(l)})}_{\text{supervised } (\mathcal{L}_{\text{labeled}})} + \underbrace{\sum_{j=1}^M \Omega(x_j^{(u)}; \theta)}_{\text{unsupervised}}$$

Recent progress on SSL has revealed promising performance in various tasks

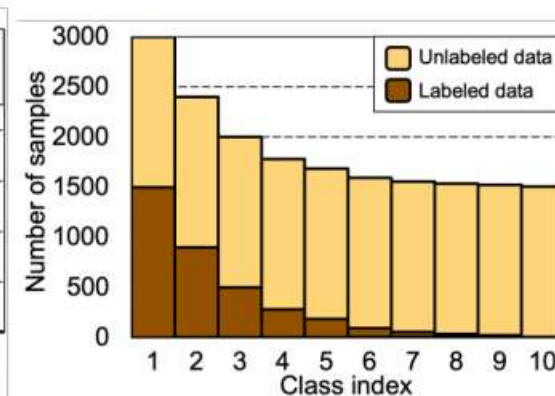


However, most existing SSL algorithms assume the datasets are class-balanced

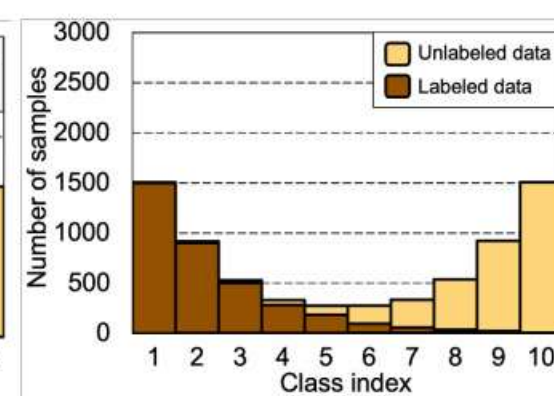
Three typical types of class distribution of unlabeled data



(a) Consistent class distribution



(b) Uniform class distribution

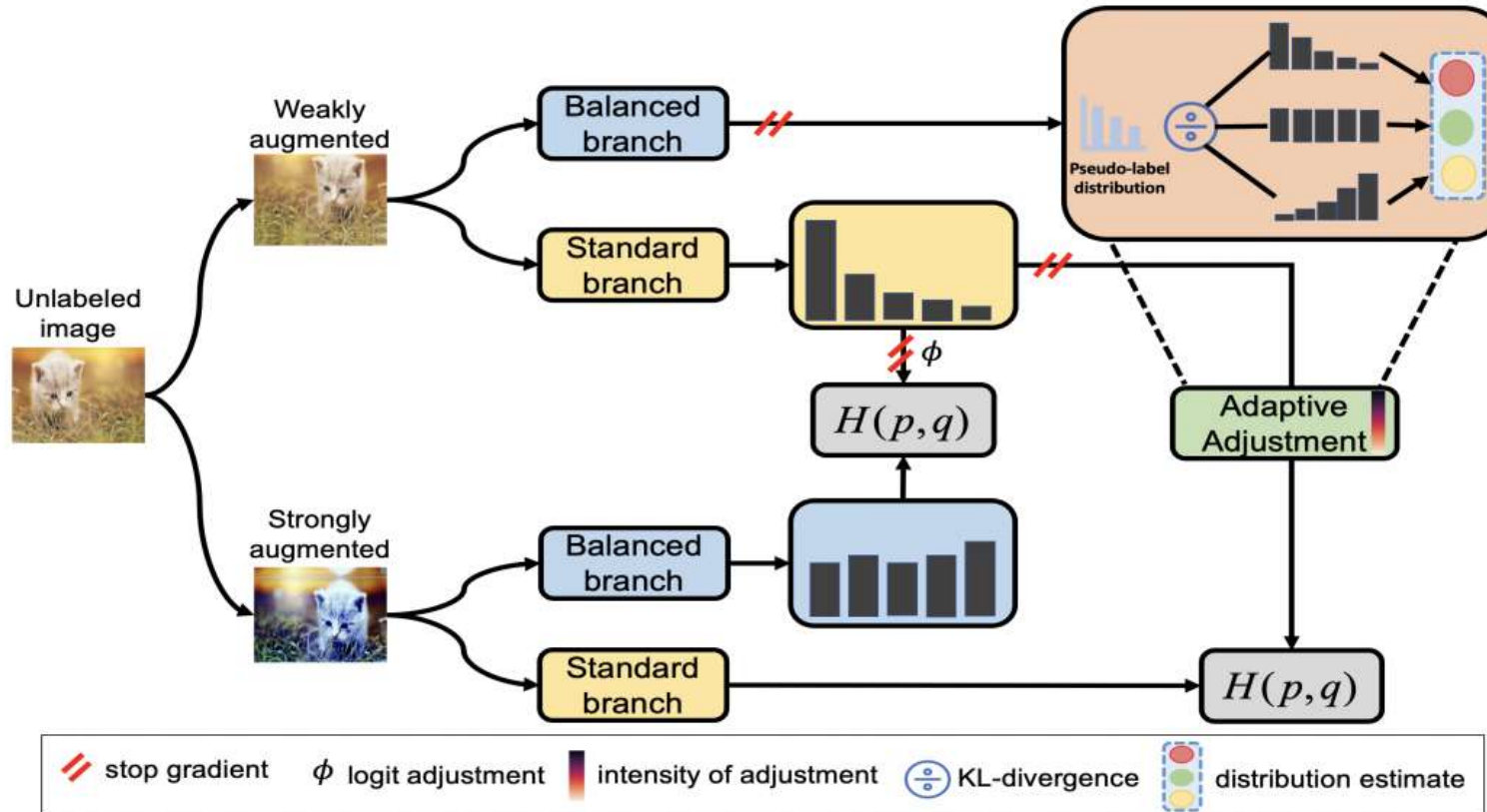


(c) Reversed class distribution

# Adaptive Consistency Regularizer (ACR)

Two findings:

- 1) Pseudo-labels biased towards minority classes can benefit the classifier learning;
- 2) Pseudo-label distribution that approximates the true distribution helps learn better feature extractor.



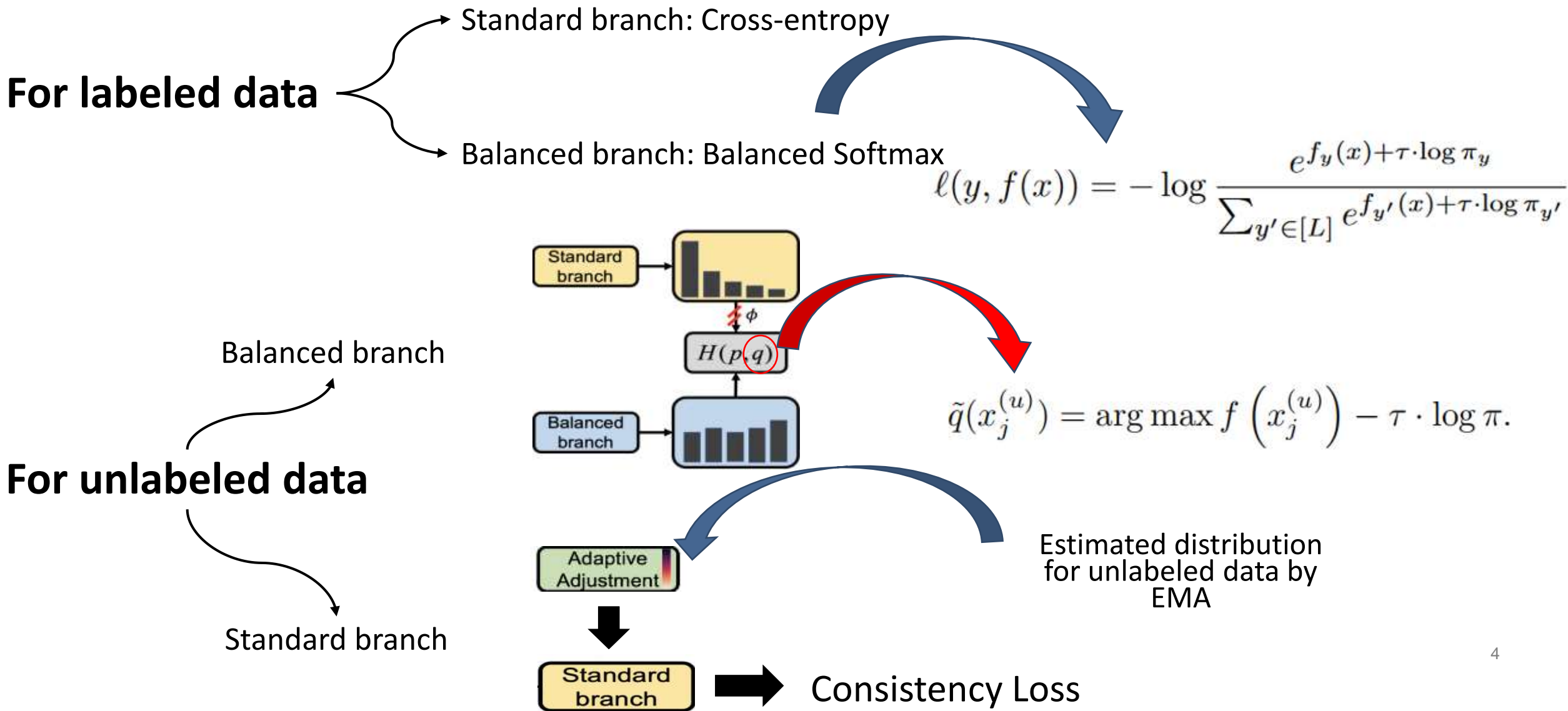
## Balanced branch

Adjust pseudo-labels appropriately biased toward the minority class via logit adjustment

## Standard branch

Refine the original pseudo-labels to match the true class distribution of unlabeled data and enhance their accuracy

# Adaptive Consistency Regularizer (ACR)





# Adaptive Consistency Regularizer (ACR)

- $\pi_{con}$  Anchor distribution for consistent
- $\pi_{uni}$  Anchor distribution for uniform
- $\pi_{rev}$  Anchor distribution for reversed
- $\pi_{est}$  Estimated distribution for unlabeled data

Calculate distances to each anchor distribution



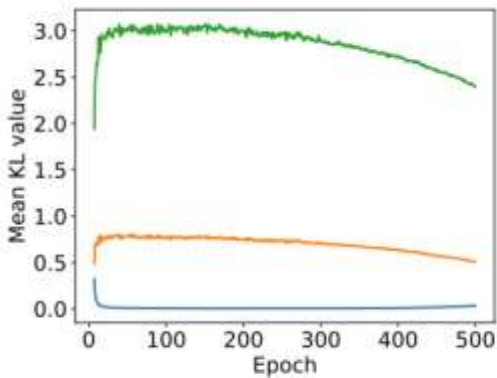
$$dist_{con} = \frac{1}{2} (D_{KL}(\pi_{con} || \pi_{est}) + D_{KL}(\pi_{est} || \pi_{con}))$$

$$dist_{uni} = \frac{1}{2} (D_{KL}(\pi_{uni} || \pi_{est}) + D_{KL}(\pi_{est} || \pi_{uni}))$$

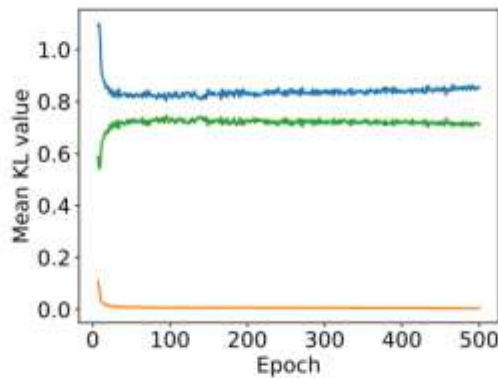
$$dist_{rev} = \frac{1}{2} (D_{KL}(\pi_{rev} || \pi_{est}) + D_{KL}(\pi_{est} || \pi_{rev})),$$

$$\tau(t) = \frac{2e^{dist_{con}^{(t-1)}}}{e^{dist_{con}^{(t-1)}} + e^{dist_{uni}^{(t-1)}} + e^{dist_{rev}^{(t-1)}}}$$

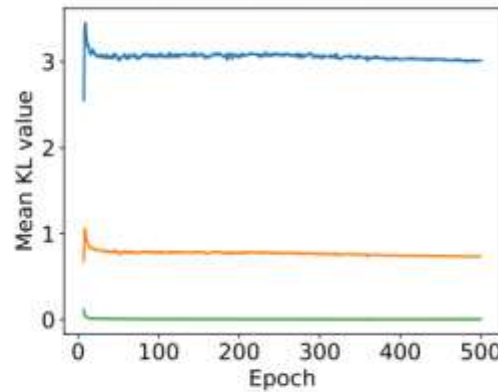
— Consistent    — Uniform    — Reversed



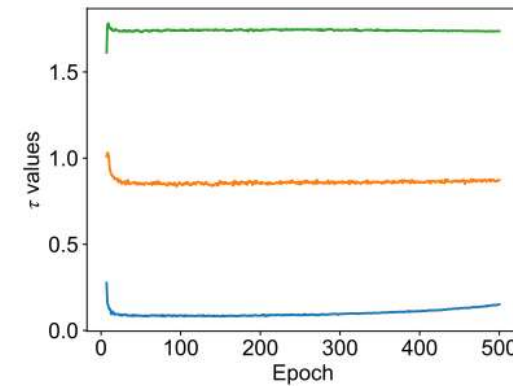
(a) Distance for *consistent* setting



(b) Distance for *uniform* setting



(c) Distance for *reversed* setting



(d)  $\tau$  values for LA

# Adaptive Consistency Regularizer (ACR)

## Sample mask generation

$$\mathcal{L}_{\text{b-con}} = \sum_{j=1}^M \tilde{M}(x_j^{(u)}) \ell \left( \tilde{f}(\mathcal{A}(x_j^{(u)})), \tilde{q}_{j\delta} \right),$$

$$\tilde{M}(x_j^{(u)}) = \mathbb{I} \left( \max \left( \delta(\tilde{f}(x_j^{(u)})) \right) \geq \rho \right) \vee$$

$$\mathbb{I} \left( \max \left( \delta(f(x_j^{(u)})) - \tau \cdot \log \pi \right) \geq \rho \right),$$

$\tilde{M}(x_j^{(u)})$  Sample mask for  $x_j^{(u)}$  in balanced branch

$\delta$  Softmax function

$\mathbb{I}$  Indicator function

$\rho$  Predefined threshold

$\pi$  Distribution of labeled data

In this way, we can

- (1) select more samples for the minority classes by considering the balanced branch's output;
- (2) obtain more confident samples through the newly constructed sample mask, which is beneficial for consistency loss to work.

# Experimental Results

## Test accuracy for consistent setting

Algorithm	CIFAR10-LT				CIFAR100-LT			
	$\gamma = \gamma_l = \gamma_u = 100$		$\gamma = \gamma_l = \gamma_u = 150$		$\gamma = \gamma_l = \gamma_u = 10$		$\gamma = \gamma_l = \gamma_u = 20$	
	$N_1 = 500$ $M_1 = 4000$	$N_1 = 1500$ $M_1 = 3000$	$N_1 = 500$ $M_1 = 4000$	$N_1 = 1500$ $M_1 = 3000$	$N_1 = 50$ $M_1 = 400$	$N_1 = 150$ $M_1 = 300$	$N_1 = 50$ $M_1 = 400$	$N_1 = 150$ $M_1 = 300$
Supervised	47.3 ± 0.95	61.9 ± 0.41	44.2 ± 0.33	58.2 ± 0.29	29.6 ± 0.57	46.9 ± 0.22	25.1 ± 1.14	41.2 ± 0.15
w/ LA [22]	53.3 ± 0.44	53.3 ± 0.21	49.5 ± 0.40	67.1 ± 0.78	30.2 ± 0.44	48.7 ± 0.89	26.5 ± 1.31	44.1 ± 0.42
FixMatch [29]	67.8 ± 1.13	77.5 ± 1.32	62.9 ± 0.36	72.4 ± 1.03	45.2 ± 0.55	56.5 ± 0.06	40.0 ± 0.96	50.7 ± 0.25
w/ DARP [14]	74.5 ± 0.78	77.8 ± 0.63	67.2 ± 0.32	73.6 ± 0.73	49.4 ± 0.20	58.1 ± 0.44	43.4 ± 0.87	52.2 ± 0.66
w/ CReST+ [34]	76.3 ± 0.86	78.1 ± 0.42	67.5 ± 0.45	73.7 ± 0.34	44.5 ± 0.94	57.4 ± 0.18	40.1 ± 1.28	52.1 ± 0.21
w/ DASO [25]	76.0 ± 0.37	79.1 ± 0.75	70.1 ± 1.81	75.1 ± 0.77	49.8 ± 0.24	59.2 ± 0.35	43.6 ± 0.09	52.9 ± 0.42
FixMatch+LA [22]	75.3 ± 2.45	82.0 ± 0.36	67.0 ± 2.49	78.0 ± 0.91	47.3 ± 0.42	58.6 ± 0.36	41.4 ± 0.93	53.4 ± 0.32
w/ DARP [14]	76.6 ± 0.92	80.8 ± 0.62	68.2 ± 0.94	76.7 ± 1.13	50.5 ± 0.78	59.9 ± 0.32	44.4 ± 0.65	53.8 ± 0.43
w/ CReST+ [34]	76.7 ± 1.13	81.1 ± 0.57	70.9 ± 1.18	77.9 ± 0.71	44.0 ± 0.21	57.1 ± 0.55	40.6 ± 0.55	52.3 ± 0.20
w/ DASO [25]	77.9 ± 0.88	82.5 ± 0.08	70.1 ± 1.68	79.0 ± 2.23	50.7 ± 0.51	60.6 ± 0.71	44.1 ± 0.61	55.1 ± 0.72
FixMatch+ABC [18]	78.9 ± 0.82	83.8 ± 0.36	66.5 ± 0.78	80.1 ± 0.45	47.5 ± 0.18	59.1 ± 0.21	41.6 ± 0.83	53.7 ± 0.55
w/ DASO [25]	80.1 ± 1.16	83.4 ± 0.31	70.6 ± 0.80	80.4 ± 0.56	50.2 ± 0.62	60.0 ± 0.32	44.5 ± 0.25	55.3 ± 0.53
FixMatch w/ ACR (ours)	<b>81.6 ± 0.19</b>	<b>84.1 ± 0.39</b>	<b>77.0 ± 1.19</b>	<b>80.9 ± 0.22</b>	<b>55.7 ± 0.12</b>	<b>65.6 ± 0.16</b>	<b>48.0 ± 0.75</b>	<b>58.9 ± 0.36</b>

ACR outperforms all algorithms even though most these methods are particularly developed based on the assumption that labeled and unlabeled data share the same class distribution



# Experimental Results

## Test accuracy for inconsistent settings

Algorithm	CIFAR10-LT ( $\gamma_l \neq \gamma_u$ )				STL10-LT ( $\gamma_u = \text{N/A}$ )			
	$\gamma_u = 1$ (uniform)		$\gamma_u = 1/100$ (reversed)		$\gamma_l = 10$		$\gamma_l = 20$	
	$N_1 = 500$ $M_1 = 4000$	$N_1 = 1500$ $M_1 = 3000$	$N_1 = 500$ $M_C = 4000$	$N_1 = 1500$ $M_C = 3000$	$N_1 = 150$ $M = 100k$	$N_1 = 450$ $M = 100k$	$N_1 = 150$ $M = 100k$	$N_1 = 450$ $M = 100k$
FixMatch [29]	73.0 $\pm$ 3.81	81.5 $\pm$ 1.15	62.5 $\pm$ 0.94	71.8 $\pm$ 1.70	56.1 $\pm$ 2.32	72.4 $\pm$ 0.71	47.6 $\pm$ 4.87	64.0 $\pm$ 2.27
w/ DARP [14]	82.5 $\pm$ 0.75	84.6 $\pm$ 0.34	70.1 $\pm$ 0.22	80.0 $\pm$ 0.93	66.9 $\pm$ 1.66	75.6 $\pm$ 0.45	59.9 $\pm$ 2.17	72.3 $\pm$ 0.60
w/ CReST [34]	83.2 $\pm$ 1.67	87.1 $\pm$ 0.28	70.7 $\pm$ 2.02	80.8 $\pm$ 0.39	61.7 $\pm$ 2.51	71.6 $\pm$ 1.17	57.1 $\pm$ 3.67	68.6 $\pm$ 0.88
w/ CReST+ [34]	82.2 $\pm$ 1.53	86.4 $\pm$ 0.42	62.9 $\pm$ 1.39	72.9 $\pm$ 2.00	61.2 $\pm$ 1.27	71.5 $\pm$ 0.96	56.0 $\pm$ 3.19	68.5 $\pm$ 1.88
w/ DASO [25]	86.6 $\pm$ 0.84	88.8 $\pm$ 0.59	71.0 $\pm$ 0.95	80.3 $\pm$ 0.65	70.0 $\pm$ 1.19	78.4 $\pm$ 0.80	65.7 $\pm$ 1.78	75.3 $\pm$ 0.44
w/ ACR (ours)	<b>92.1</b> $\pm$ 0.18	<b>93.5</b> $\pm$ 0.11	<b>85.0</b> $\pm$ 0.09	<b>89.5</b> $\pm$ 0.17	<b>77.1</b> $\pm$ 0.24	<b>83.0</b> $\pm$ 0.32	<b>75.1</b> $\pm$ 0.70	<b>81.5</b> $\pm$ 0.25

## CIFAR100-LT ( $\gamma_l \neq \gamma_u$ )

Algorithm	$\gamma_u = 1$ (uniform)		$\gamma_u = 1/10$ (reversed)	
	$N_1 = 50$ $M_1 = 400$	$N_1 = 150$ $M_1 = 300$	$N_1 = 50$ $M_C = 400$	$N_1 = 150$ $M_C = 300$
	FixMatch [29]	45.5 $\pm$ 0.71	58.1 $\pm$ 0.72	44.2 $\pm$ 0.43
w/ DARP [14]	43.5 $\pm$ 0.95	55.9 $\pm$ 0.32	36.9 $\pm$ 0.48	51.8 $\pm$ 0.92
w/ CReST [34]	43.5 $\pm$ 0.30	59.2 $\pm$ 0.25	39.0 $\pm$ 1.11	56.4 $\pm$ 0.62
w/ CReST+ [34]	43.6 $\pm$ 1.60	58.7 $\pm$ 0.16	39.1 $\pm$ 0.77	56.4 $\pm$ 0.78
w/ DASO [25]	53.9 $\pm$ 0.66	61.8 $\pm$ 0.98	51.0 $\pm$ 0.19	60.0 $\pm$ 0.31
w/ ACR (ours)	<b>66.0</b> $\pm$ 0.25	<b>73.4</b> $\pm$ 0.22	<b>57.0</b> $\pm$ 0.46	<b>67.6</b> $\pm$ 0.12

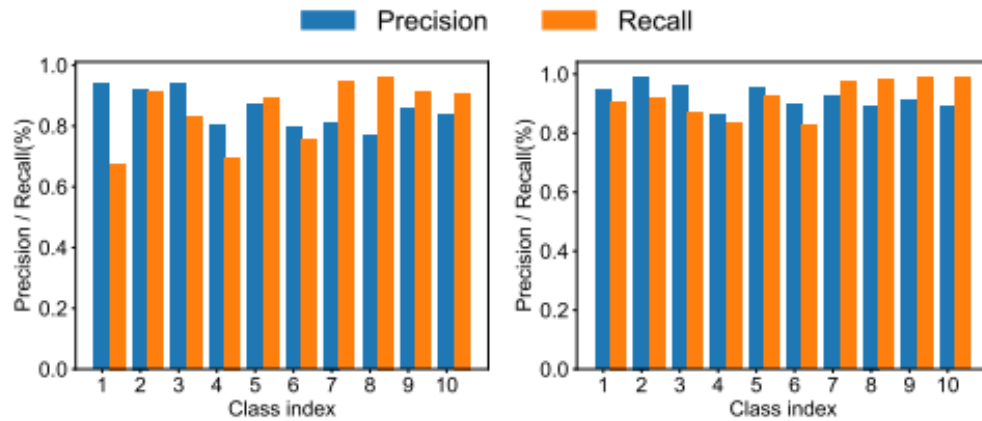
## Test accuracy on ImageNet-127

Algorithm	32 $\times$ 32	64 $\times$ 64
FixMatch [29]	29.7	42.3
w/ DARP [14]	30.5	42.5
w/ DARP+cRT [14]	39.7	51.0
w/ CReST+ [34]	32.5	44.7
w/ CReST++LA [22]	40.9	55.9
w/ CoSSL [9]	43.7	53.9
w/ TRAS [35]	46.2	54.1
w/ ACR (ours)	<b>57.2</b>	<b>63.6</b>

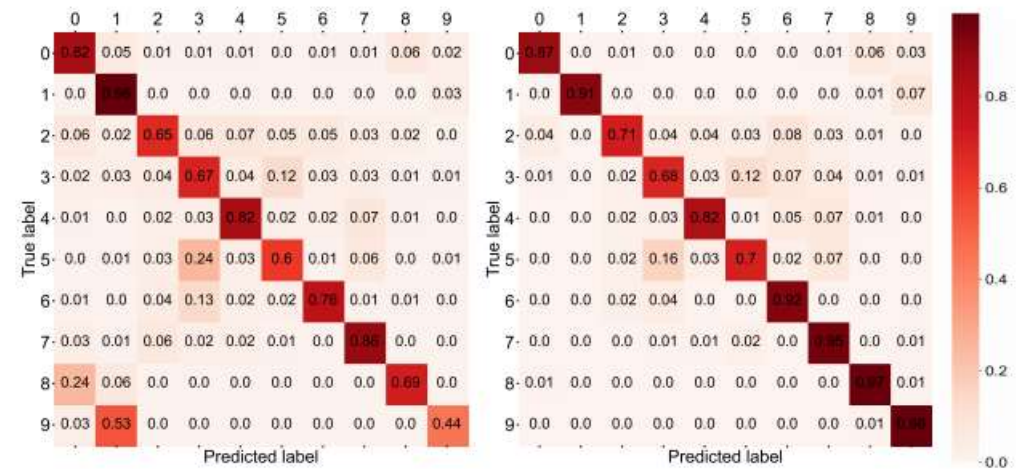
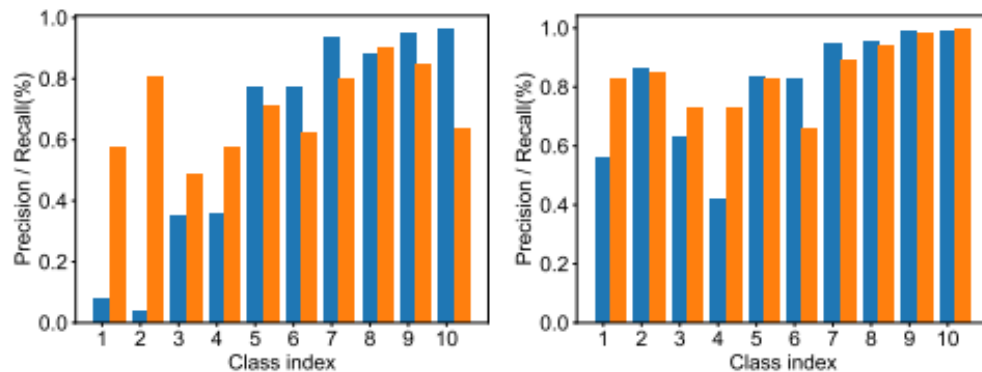
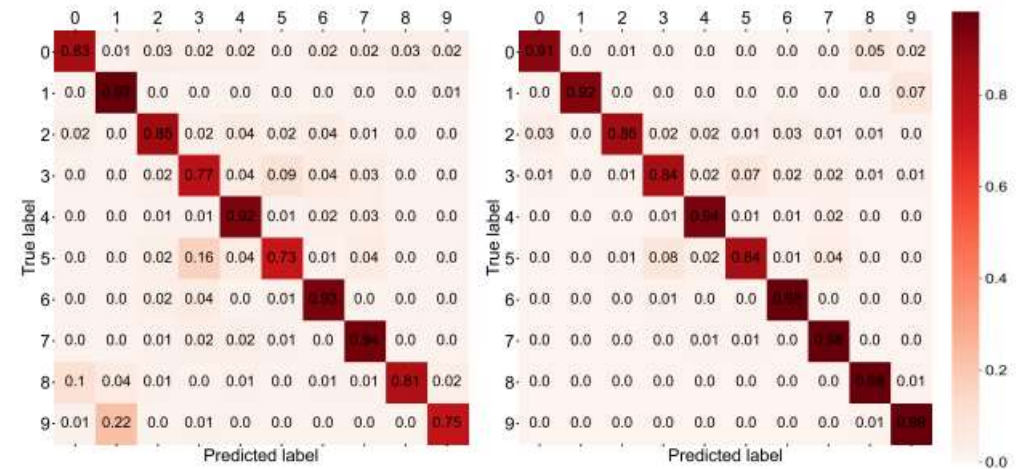


# Experimental Results

The precision and recall of pseudo-labels

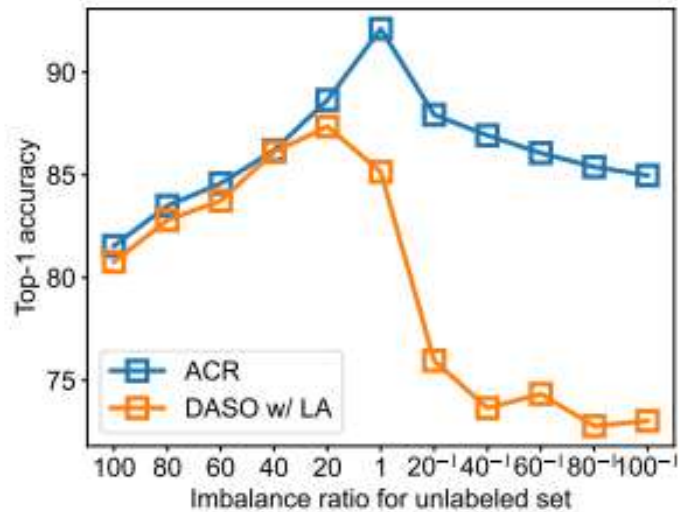


Confusion matrices

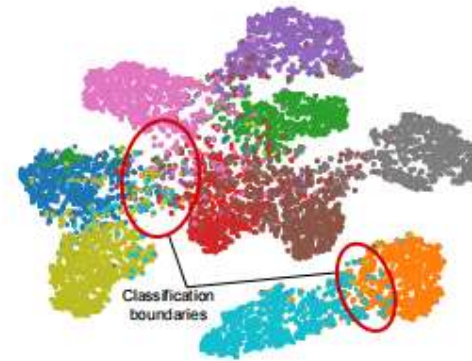


# Experimental Results

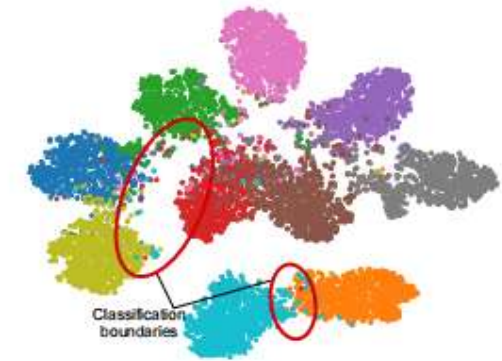
More settings:



The t-SNE visualization:



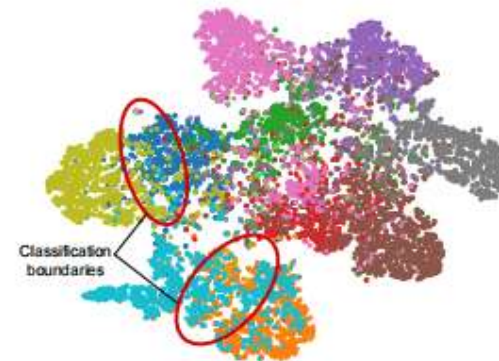
(a) DASO for *uniform*



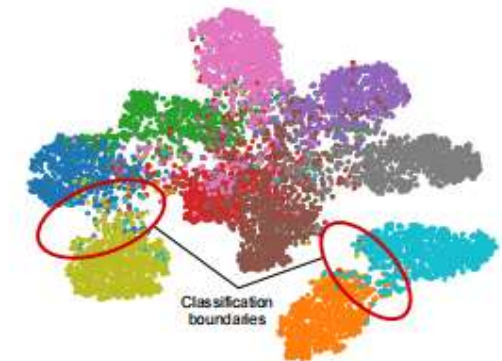
(b) ACR for *uniform*

Ablation studies:

Ablations	CIFAR10-LT			CIFAR100-LT		
	Con	Uni	Rev	Con	Uni	Rev
ACR(ours)	81.6	92.1	85.0	55.7	66.0	57.0
w/o sample mask principle	81.7	91.1	84.6	55.0	63.7	55.0
w/o adaptive LA	76.8	92.4	85.1	53.5	62.8	56.1
w/o LA for balanced branch	74.3	90.6	83.5	54.5	66.2	56.7
w/o balanced softmax	76.7	93.0	84.8	55.3	65.6	57.3
w/o gradients from balanced branch	73.7	92.3	85.2	54.3	65.2	56.7
w/o labeled data in unlabeled set	81.0	92.7	79.9	56.1	66.4	56.8



(c) DASO for *reversed*



(d) ACR for *reversed*

## Conclusion

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We presents a simple and effective method by minimizing the adaptive consistency regularizer (ACR) for long-tailed semi-supervised learning with unknown class distributions of the unlabeled data.

- Benefit classifier learning by generating pseudo-labels that are properly biased towards minority classes.
- Benefit representation learning by generating pseudo-labels whose distribution approximates the true class distribution.

**Thanks!**