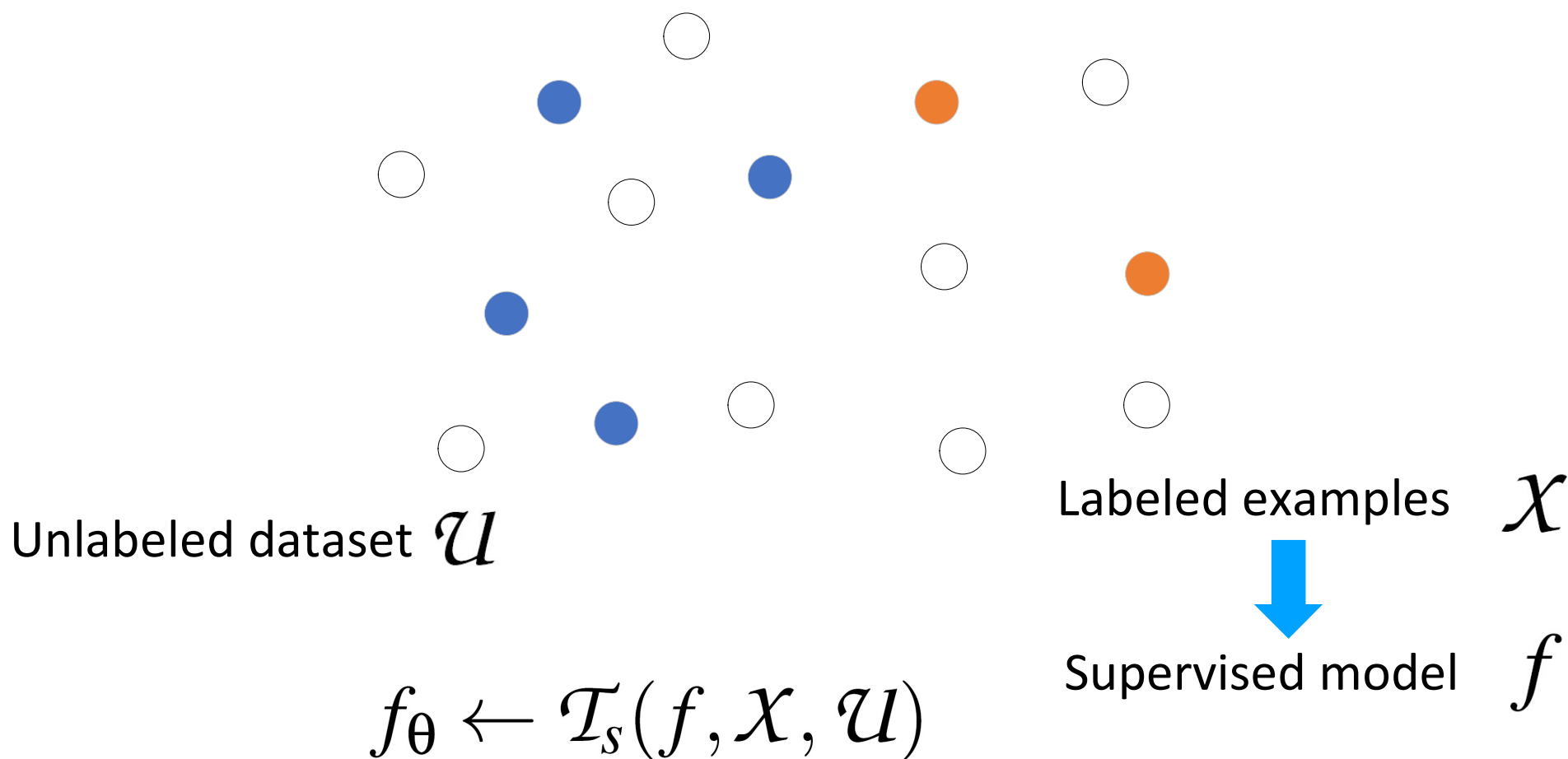


Poisoning the Unlabeled Dataset of Semi-Supervised Learning

Nicholas Carlini, Google
USENIX Security Symposium, 2021

Semi-supervised learning

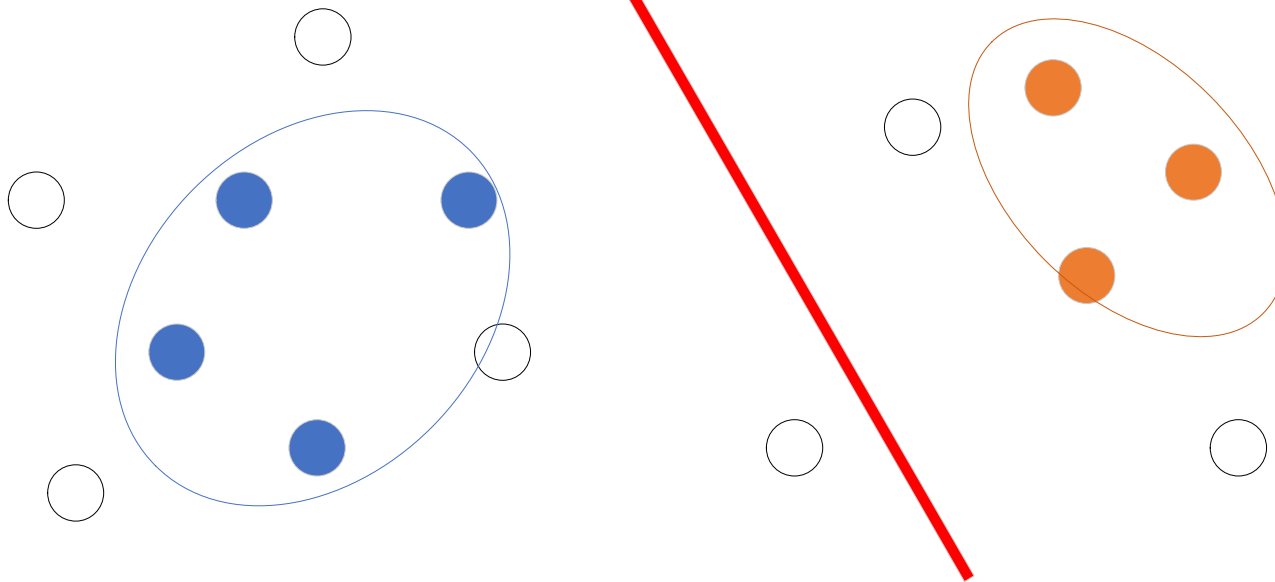
- Collecting “Labeled” Data is Expensive.



Semi-supervised learning

- Transform into Fully-supervised Problem

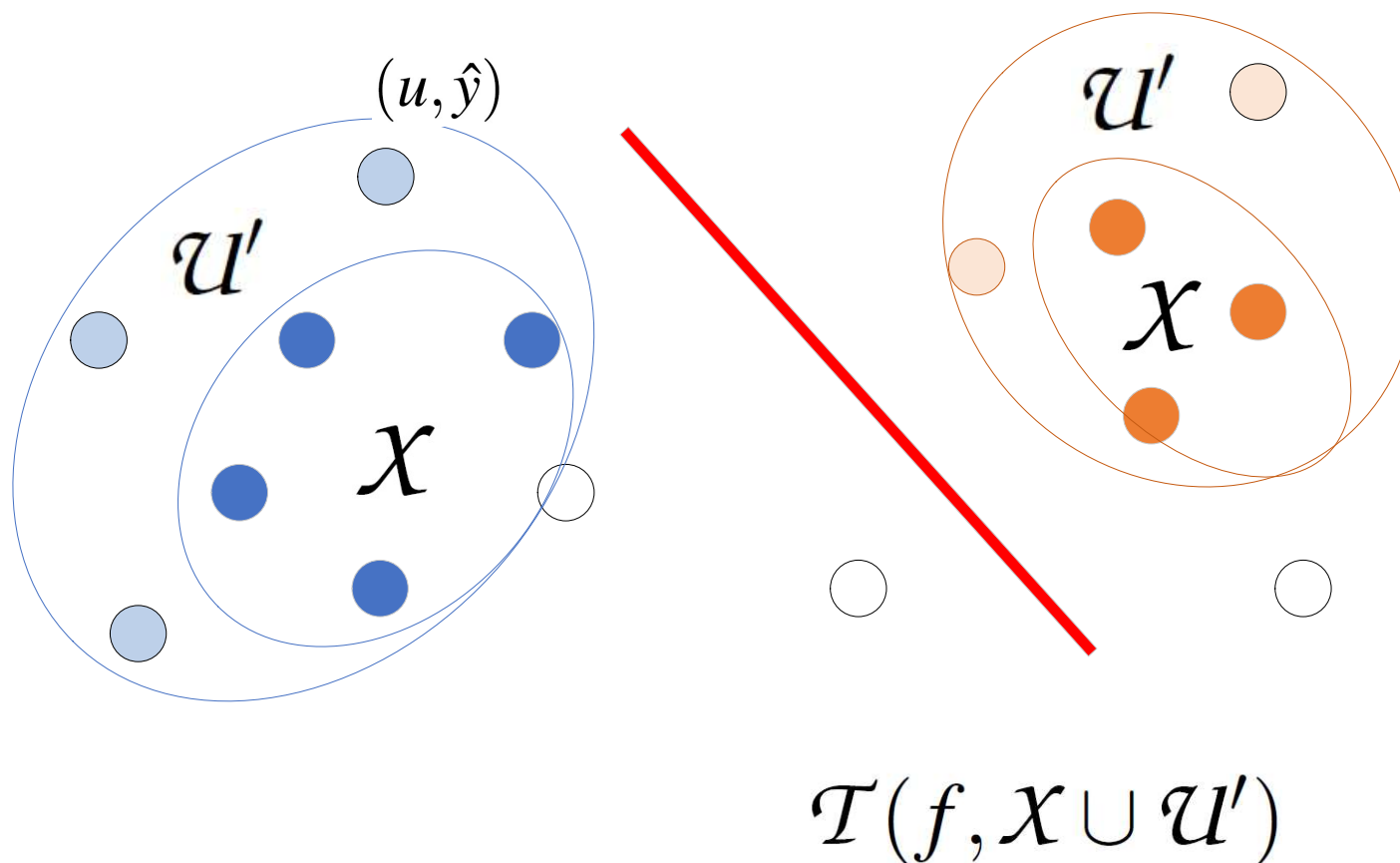
guessed label (u, \hat{y})



$$\hat{y} = f(u; \theta_i)$$

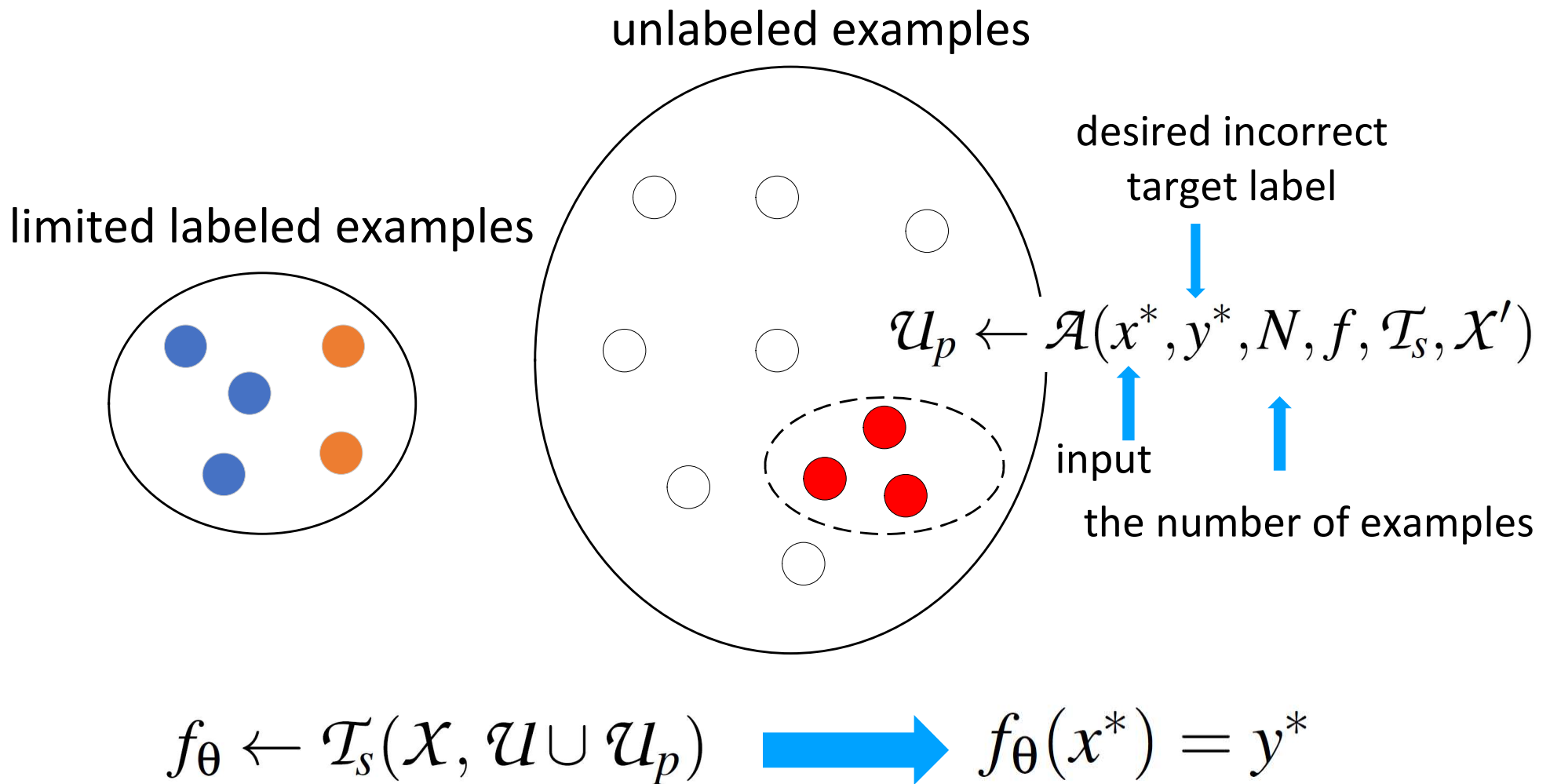
Semi-supervised learning

- Transform into Fully-supervised Problem



Threat Model

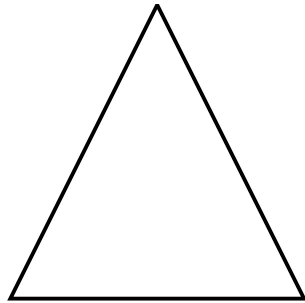
- Transform into Fully-supervised Problem



Poisoning the Unlabeled Dataset

- Problem

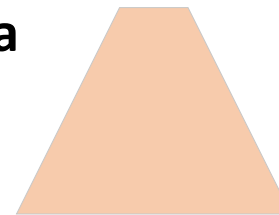
How a task should be completed



What should be done

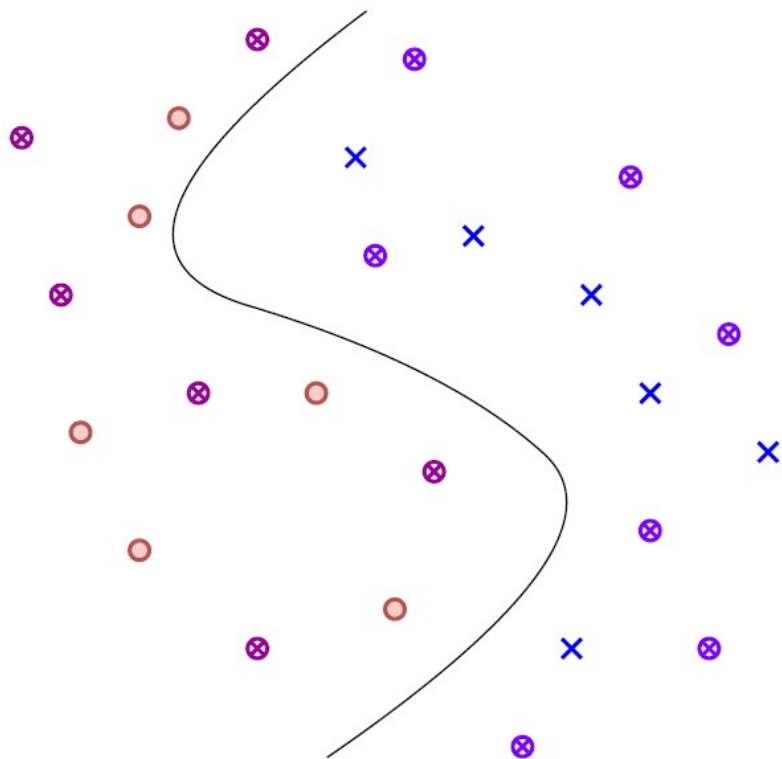


Teach itself from this unlabeled data



Poisoning the Unlabeled Dataset

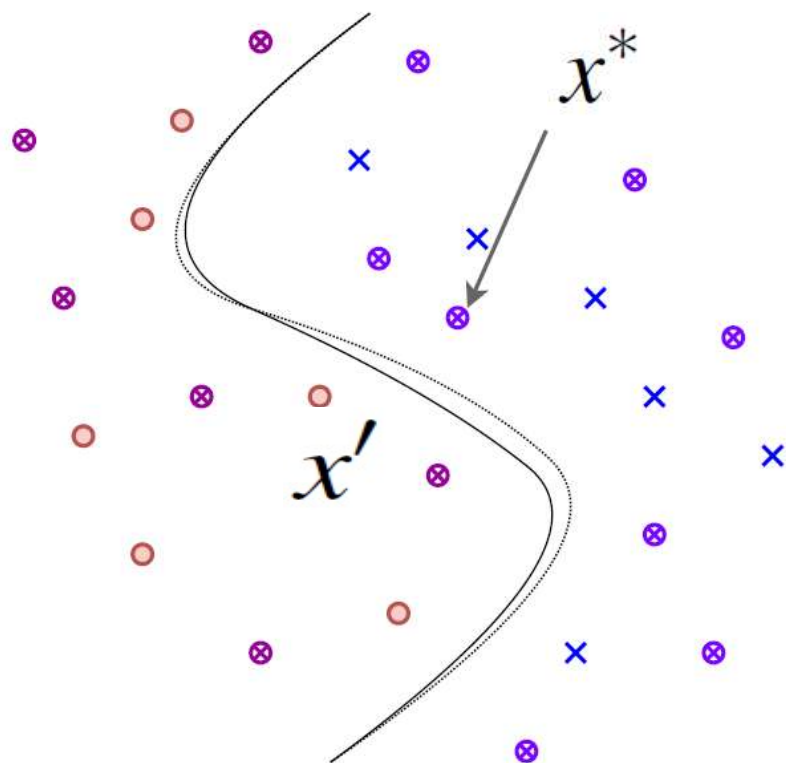
- Interpolation Consistency Poisoning



(a) A classifier trained on a semi-supervised dataset of red \odot s, blue \times s, and *unlabeled* \otimes s. During training the unlabeled \otimes s are given pseudo-labels such that the correct original decision boundary is learned.

Poisoning the Unlabeled Dataset

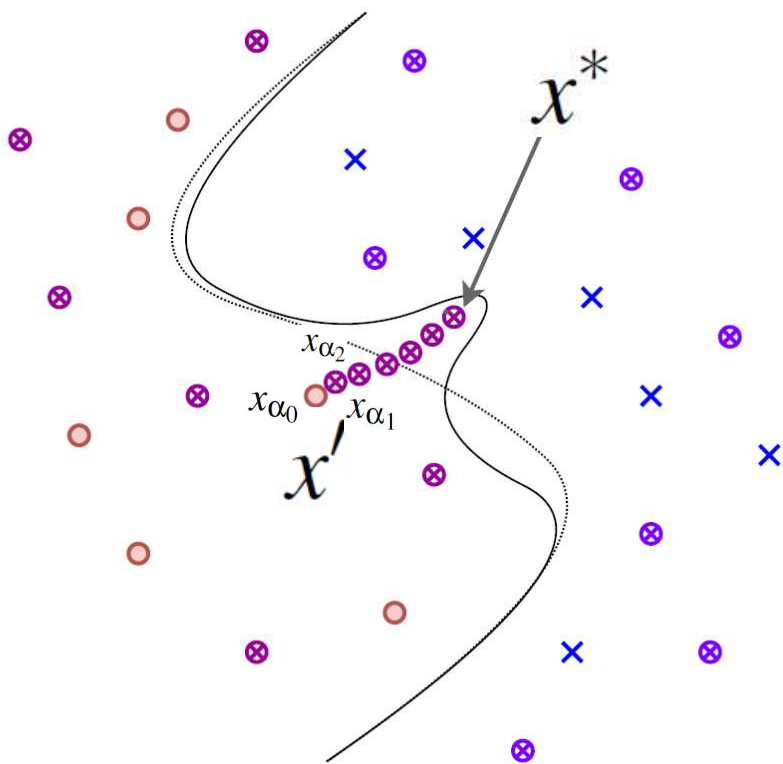
- Interpolation Consistency Poisoning



(b) When inserting just one new *unlabeled* poisoned example near the boundary, the model gives it the correct pseudo label of the blue \times s. The poisoning attempt fails, and the decision boundary remains largely unchanged.

Poisoning the Unlabeled Dataset

- Interpolation Consistency Poisoning



$$\{x_{\alpha_i}\}_{i=0}^{N-1} = \text{interp}(x', x^*, \alpha_i)$$

$$\text{interp}(x', x^*, 0) = x'$$

$$\text{interp}(x', x^*, 1) = x^*$$

$$f(x_{\alpha_0}) = f(x_{N-1}) = f(x^*) = y^*$$

Poisoning the Unlabeled Dataset

- **Interpolation Strategy**

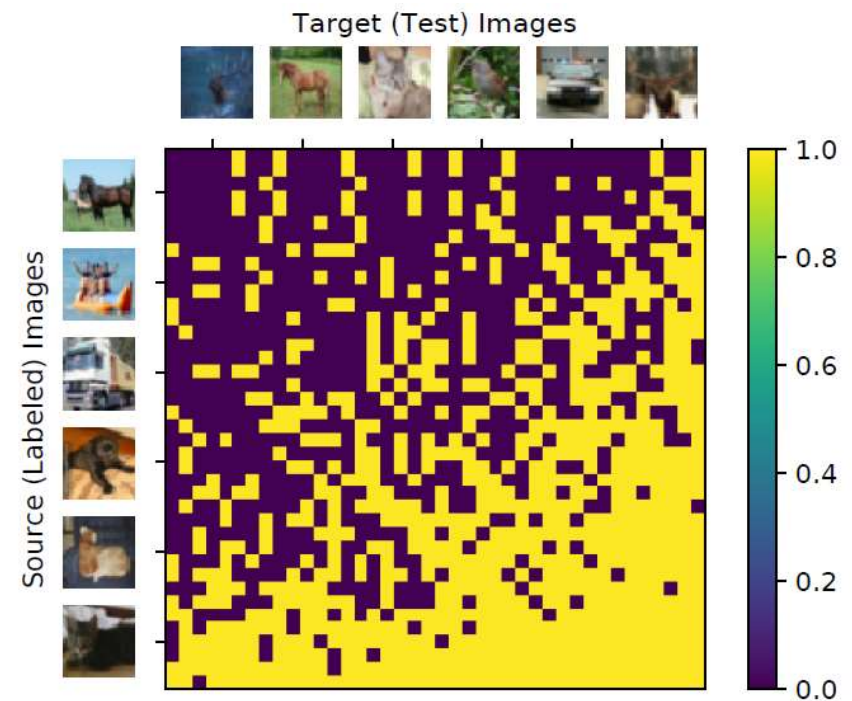
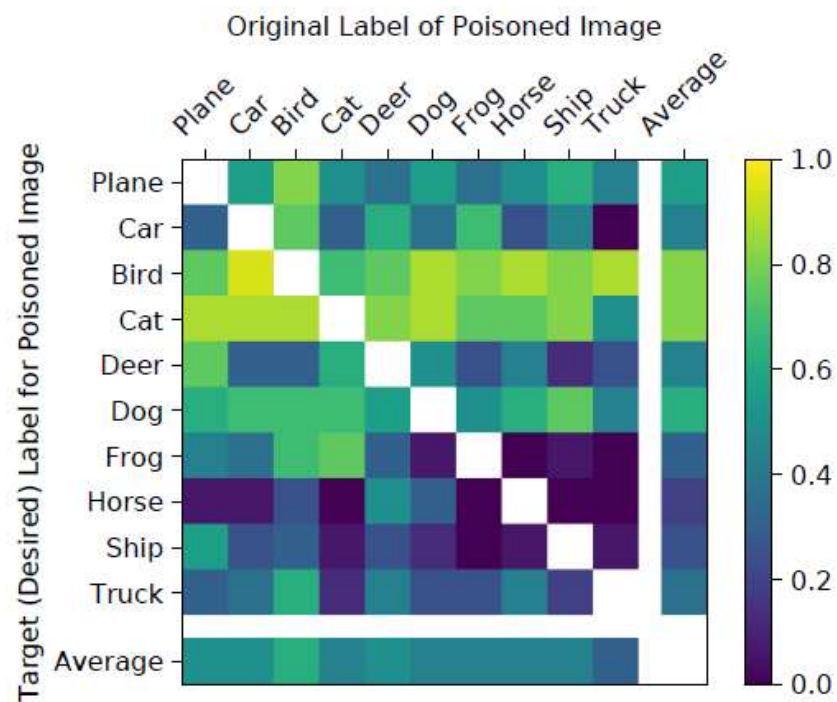
$$\text{interp}(x', x^*, \alpha) = x' \cdot (1 - \alpha) + x^* \cdot \alpha$$

- **Density of poisoned samples.**

$$\hat{\rho}(x) = \rho(x) \cdot \left(\int_0^1 \rho(x) dx \right)^{-1} \quad \Pr[p < \alpha < q] = \int_p^q \hat{\rho}(x) dx$$

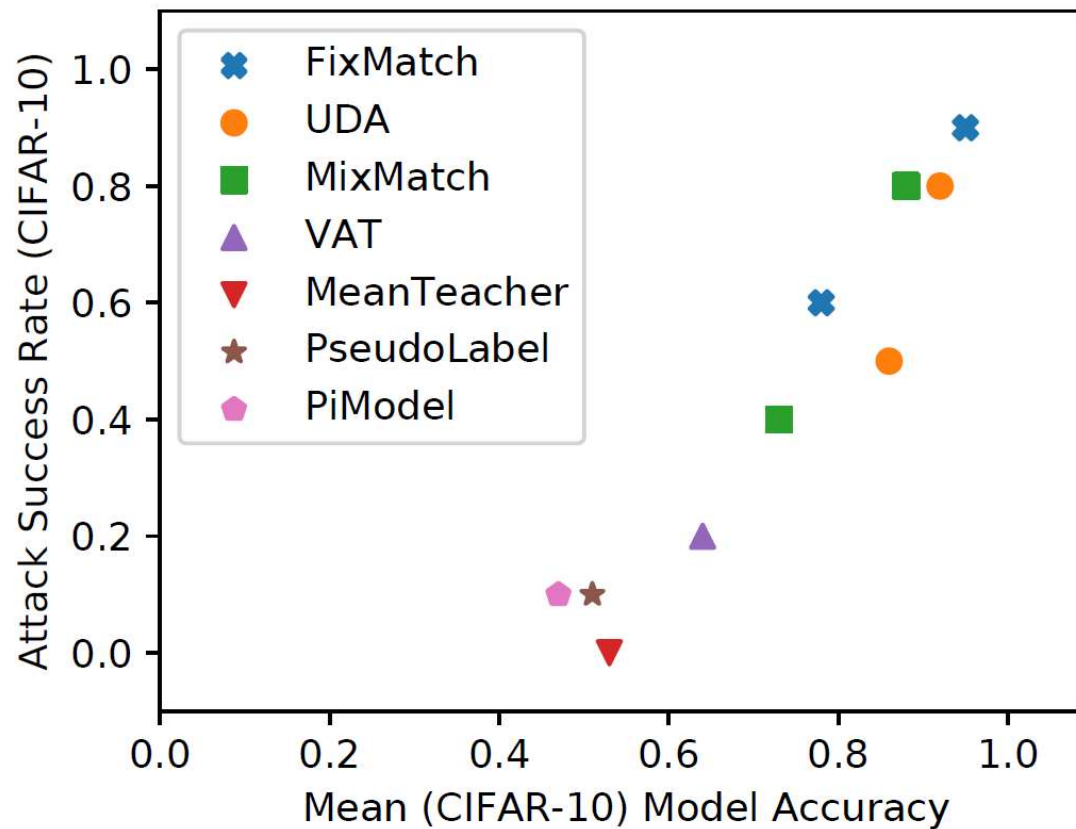
Evaluation

- Evaluation across source- and target-image



Evaluation

- **Evaluation across training techniques**



Evaluation

- Evaluation across datasets

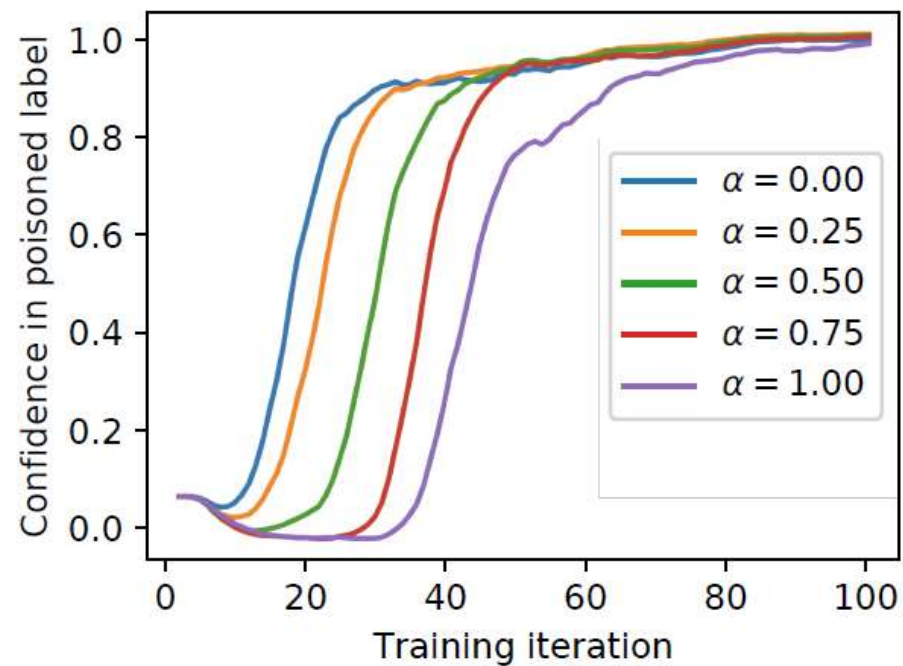
Dataset (% poisoned)	CIFAR-10			SVHN			STL-10		
	0.1%	0.2%	0.5%	0.1%	0.2%	0.5%	0.1%	0.2%	0.5%
MixMatch	5/8	6/8	8/8	4/8	5/8	5/8	4/8	6/8	7/8
UDA	5/8	7/8	8/8	5/8	5/8	6/8	-	-	-
FixMatch	7/8	8/8	8/8	7/8	7/8	8/8	6/8	8/8	8/8

- Evaluation across number of labeled examples

Dataset (# labels)	CIFAR-10			SVHN		
	40	250	4000	40	250	4000
MixMatch	5/8	4/8	1/8	6/8	4/8	5/8
UDA	5/8	5/8	2/8	5/8	4/8	4/8
FixMatch	7/8	7/8	7/8	7/8	6/8	7/8

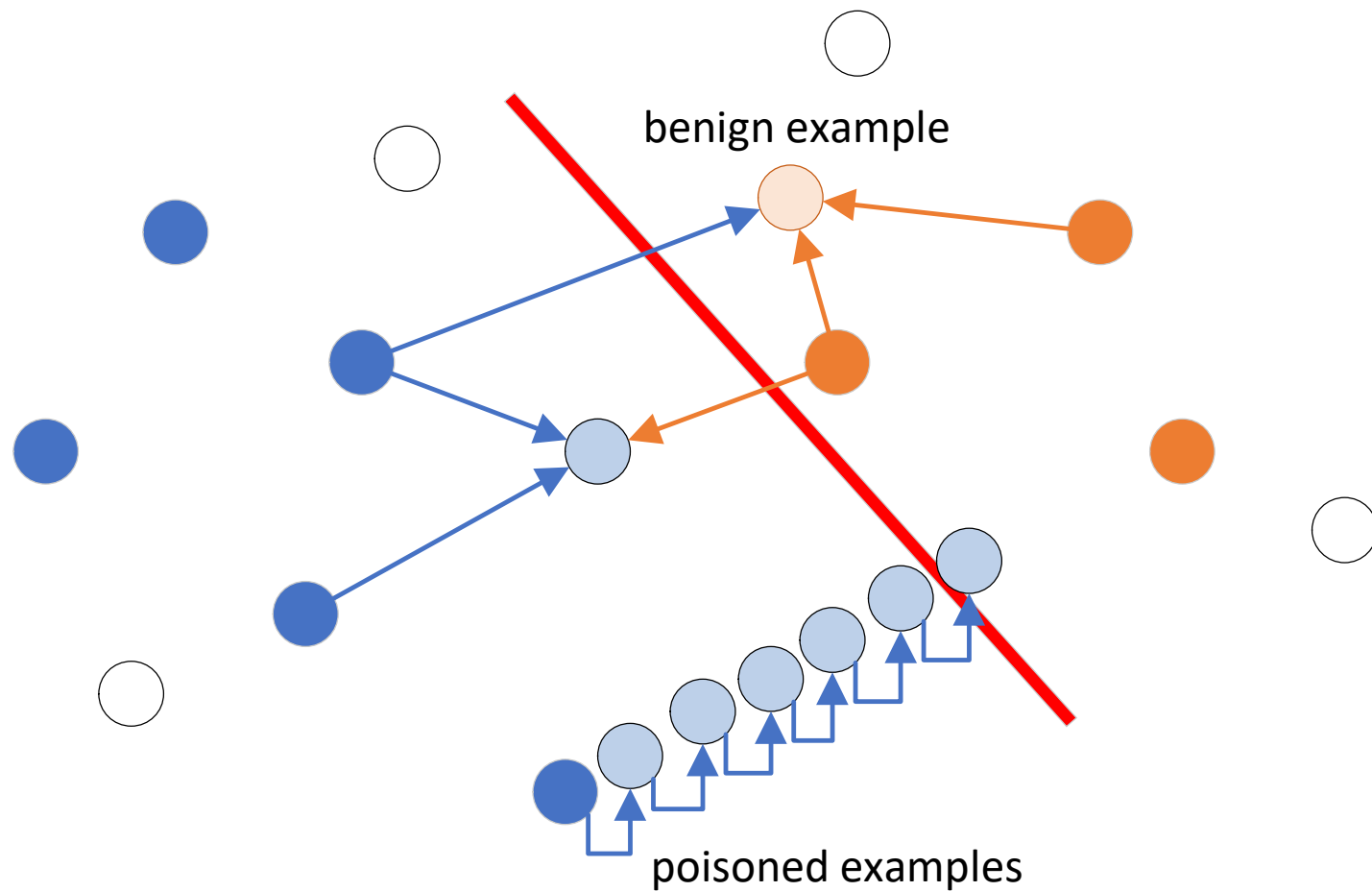
Evaluation

- Why does this attack work?



Defense

- **Monitoring Training Dynamics**



Defense

- **Computing Pairwise Influence**

Difference in the model f predictions on example j from i epoch to $i+1$.

$$\partial f_{\theta_i}(u_j) = f_{\theta_{i+1}}(u_j) - f_{\theta_i}(u_j)$$

Model's collection of prediction difference from epoch a to epoch b .

$$\mu_j^{(a,b)} = [\partial f_{\theta_a}(u_j) \quad \partial f_{\theta_{a+1}}(u_j) \quad \dots \quad \partial f_{\theta_{b-1}}(u_j) \quad f_{\theta_b}(u_j)]$$

The influence of example u_i on u_j

$$\text{Influence}(u_i, u_j) = \|\mu_i^{(0,K-2)} - \mu_j^{(1,K-1)}\|_2^2$$

Defenses

- Identifying Poisoned Example

