

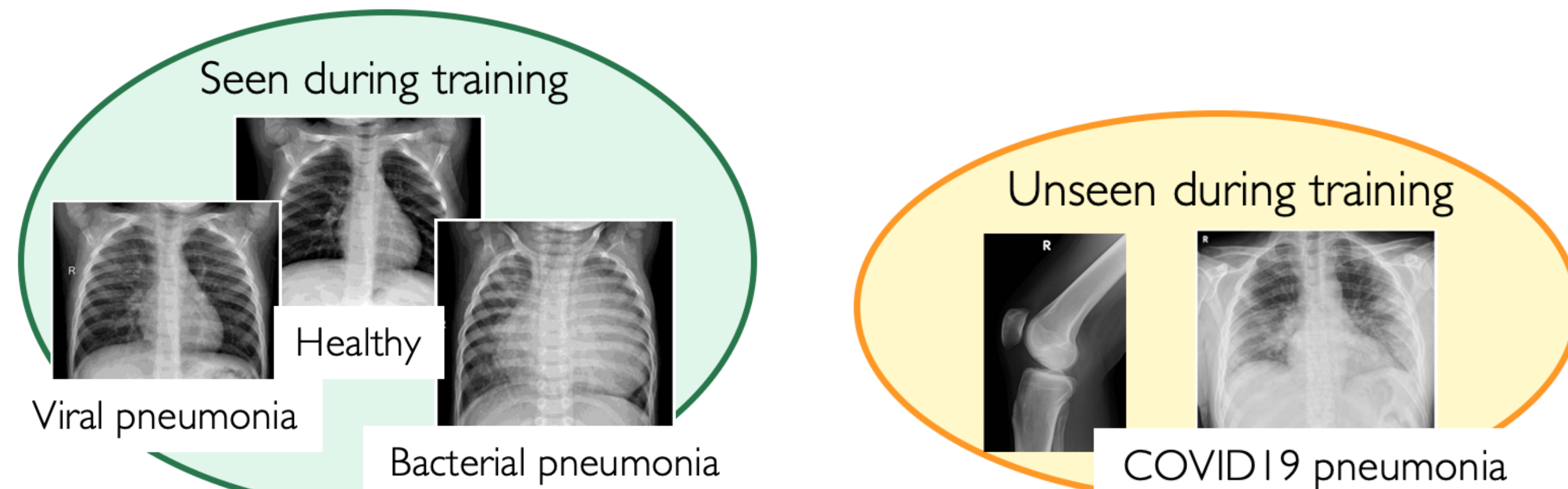
# Novel disease detection using ensembles with regularized disagreement

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## NOVEL CLASSES AS OOD DATA

**Problem:** Classifier predictions are incorrect on novel classes.  
→ Flag data from unseen classes as out-of-distribution (OOD).



→ Novel classes are often similar to in-distribution (ID) classes  
⇒ difficult to distinguish ID and OOD data.

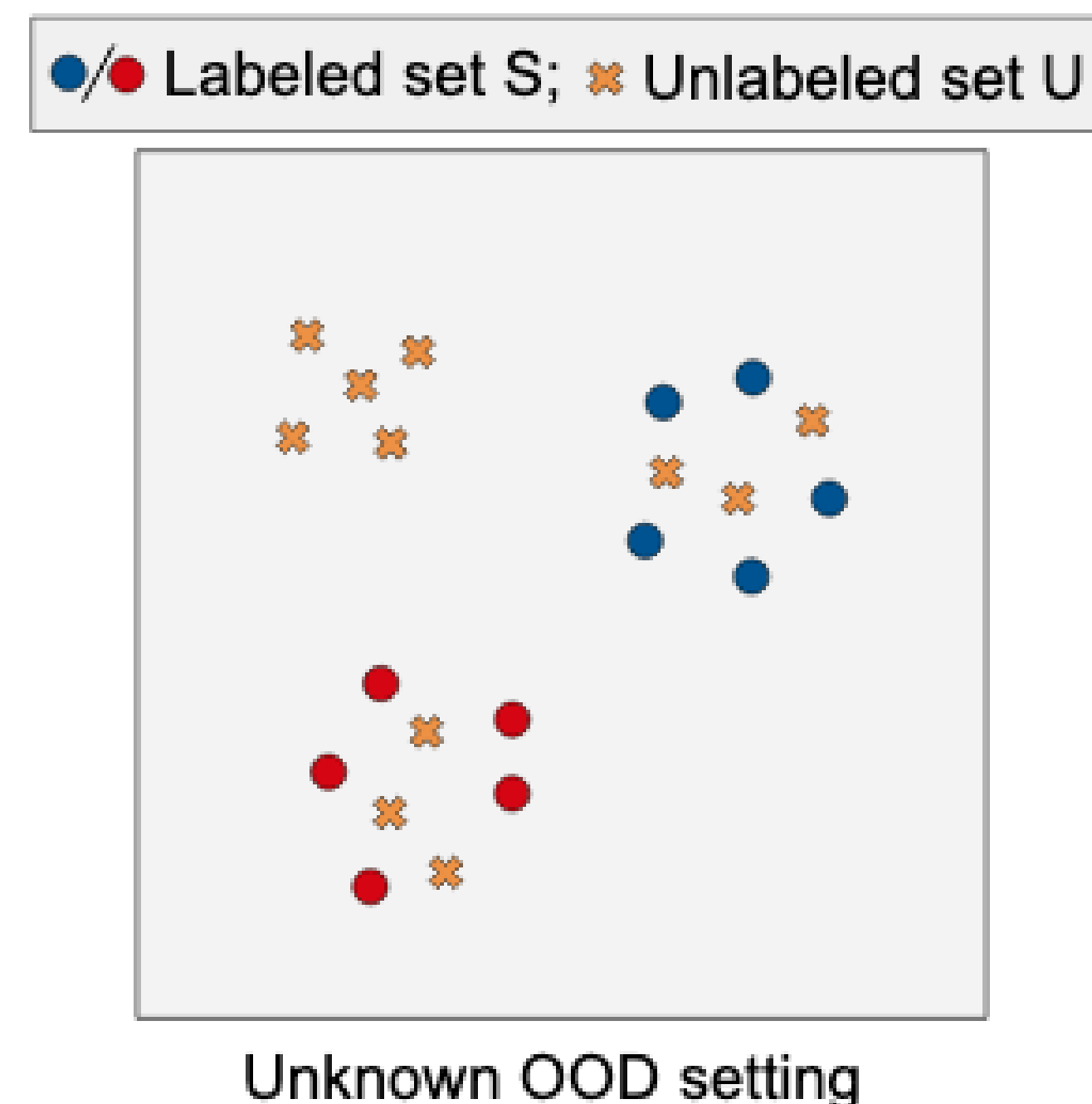
Existing OOD detection methods (assuming different access to OOD data) **perform poorly on novel-class detection.**

## OUR SETTING

### Available data:

- ▶ Labeled set with ID samples.  
→ e.g. the training set for the prediction task.
- ▶ Unlabeled set with unknown mixture of ID and OOD data.  
→ e.g. hospital collects all X-rays performed during the day.

### Unknown OOD setting:



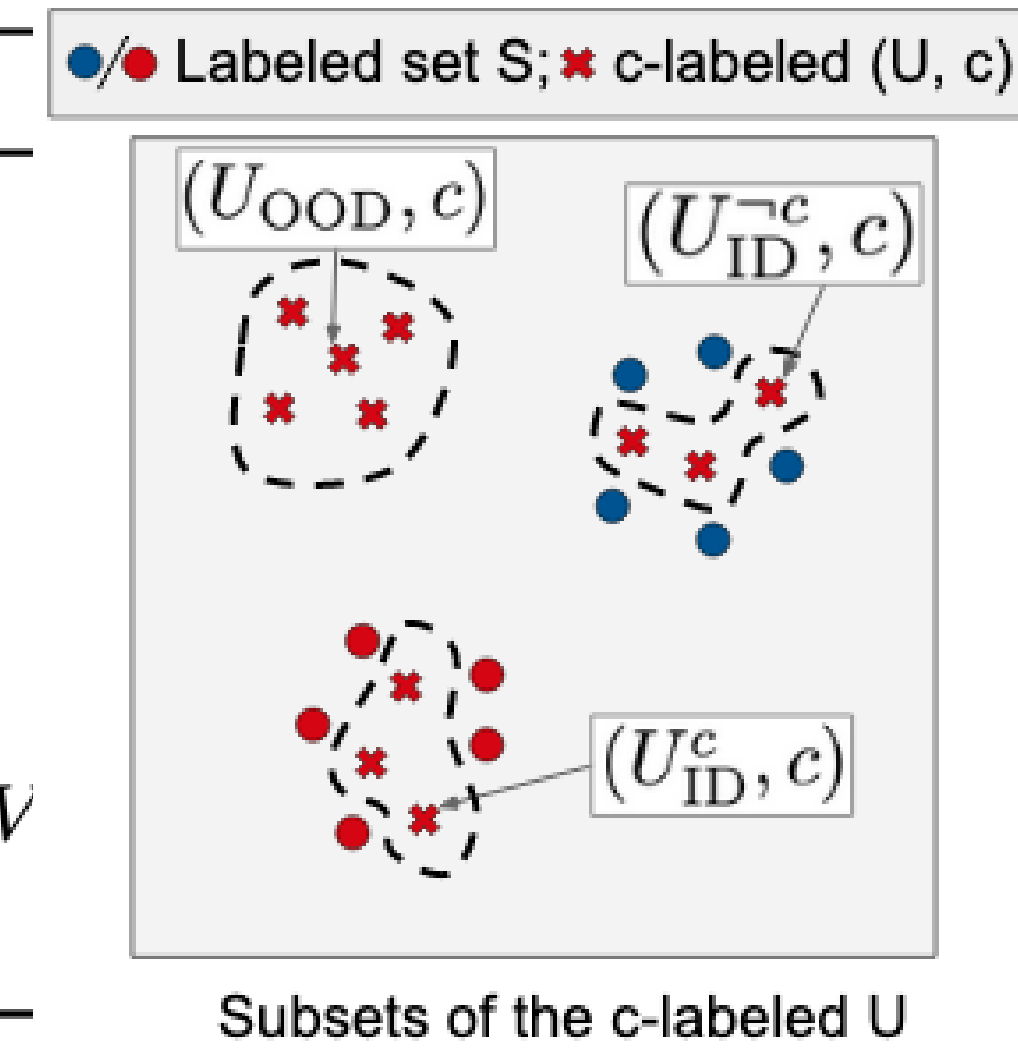
Previous methods that employ the Unknown OOD setting (e.g. nnPU, MCD) **fail to leverage unlabeled data effectively.**

## OUR APPROACH

**Idea:** Train an **Ensemble w/ Regularized Disagreement.**

### Algorithm 1: Fine-tuning the ERD ensemble

**Input:** Train set  $S$ , Validation set  $V$ , Unlabeled set  $U$ , Weights  $W$  pretrained on  $S$ , Ensemble size  $K$   
**Result:** ERD ensemble  $\{f_{y_i}\}_{i=1}^K$   
Sample  $K$  different labels  $\{y_1, \dots, y_K\}$  from  $\mathcal{Y}$   
**for**  $c \leftarrow \{y_1, \dots, y_K\}$  **do** // fine-tune  $K$  models  
   $f_c \leftarrow \text{Initialize}(W)$   
   $(U, c) \leftarrow \{(x, c) : x \in U\}$   
   $f_c \leftarrow \text{FinetuneWithEarlyStopping}(f_c, S \cup (U, c); V)$   
**return**  $\{f_{y_i}\}_{i=1}^K$

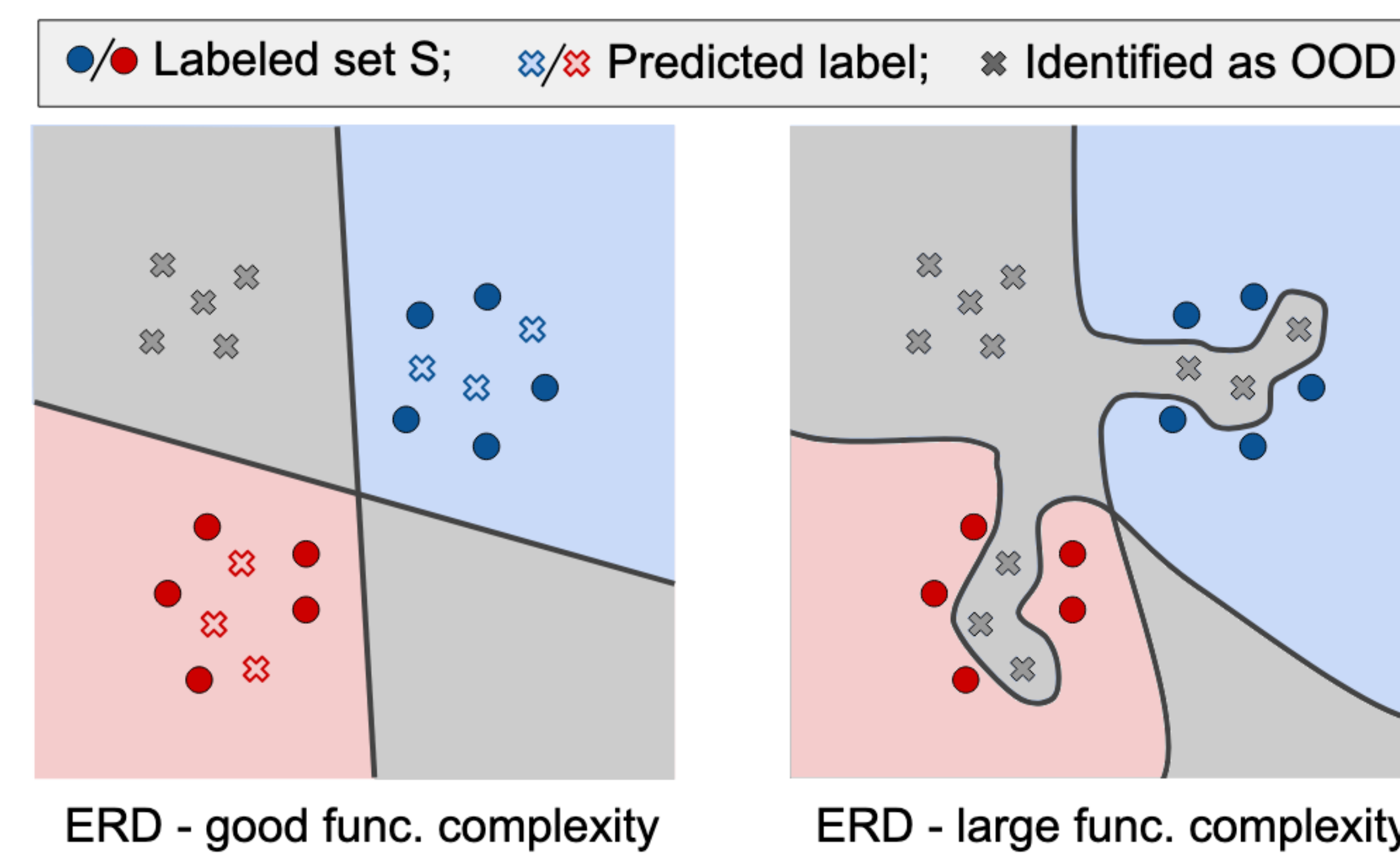


### At test time:

- ▶ For a test sample  $x$ , use outputs  $f_1(x), \dots, f_k(x)$  to compute the **average pairwise disagreement score** (details later).  
→ Flag as OOD samples with score larger than threshold  $\tau$ .

## KEY INGREDIENTS

- 1) Regularization:** Prevent complex models from interpolating on  $S \cup (U, c)$ .  
→ We **early stop** at epoch with highest ID validation accuracy.



### 2) Average pairwise disagreement score:

$$(\text{Avg} \circ \rho)(f_1(x), \dots, f_K(x)) := \frac{2}{K(K-1)} \sum_{i \neq j} \rho(f_i(x), f_j(x))$$

→ e.g.  $\rho$  = total variation distance

- ▶ Unlike prior OOD metrics (e.g. entropy of average predictor), our score exploits ensemble diversity.

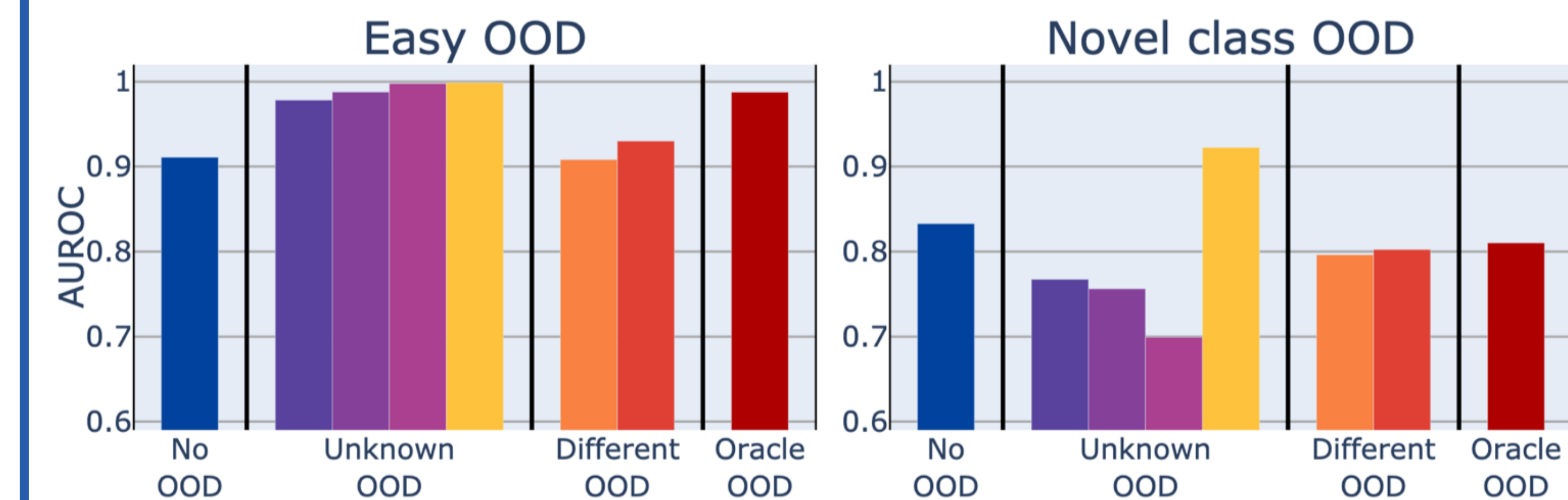
## EXPERIMENTS

**Evaluation metric:** Area under the ROC curve (AUROC).  
→ higher is better.

→ TP = correctly identified OOD; FP = ID flagged as OOD.

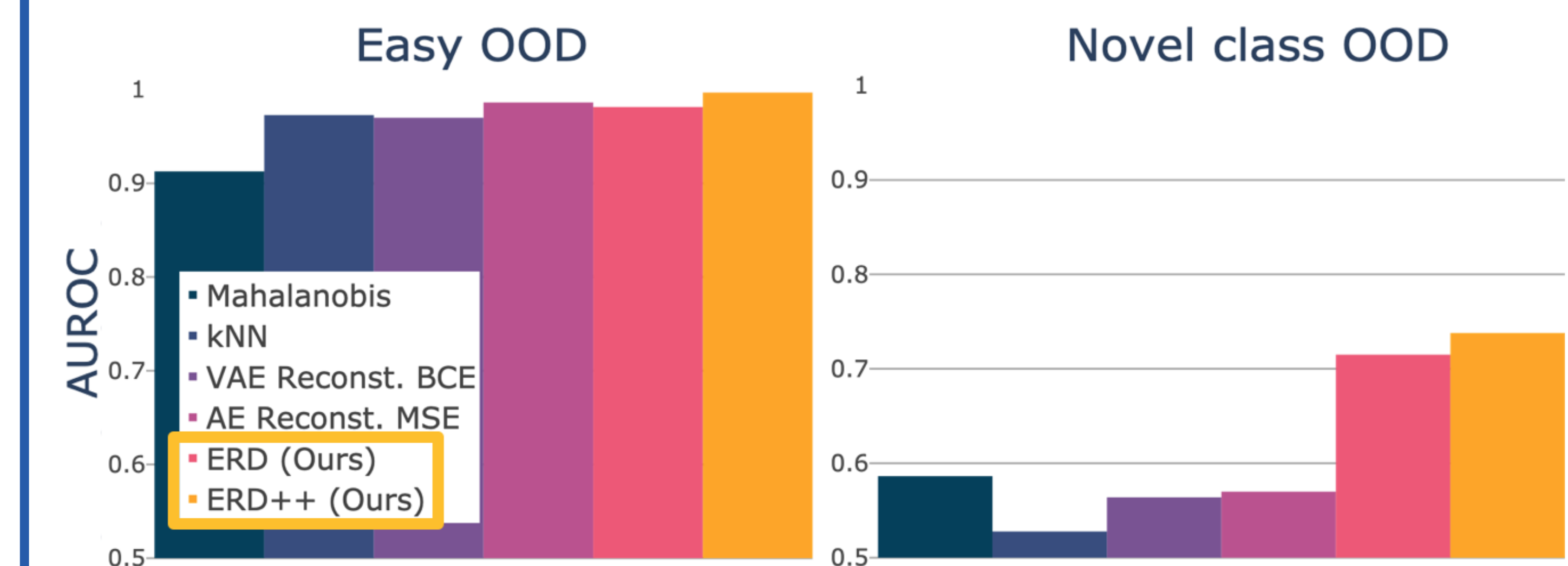
### 1) Natural images

*Easy OOD:* SVHN vs CIFAR10, CIFAR10 vs SVHN etc  
*Novel class OOD:* CIFAR100[0-49] vs CIFAR100[50-99] etc



### 2) Medical images

*Easy OOD:* Chest X-ray vs Knee X-ray etc  
*Novel class OOD:* Novel disease as OOD



→ **Image modalities:** Frontal and lateral chest X-rays and retinal images.

→ **ERD++:** our approach trained from random initializations instead of pretrained weights.