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Fine-tuning Convolutional Neural Networks for Biomedical Image Analysis: Actively and Incrementally*

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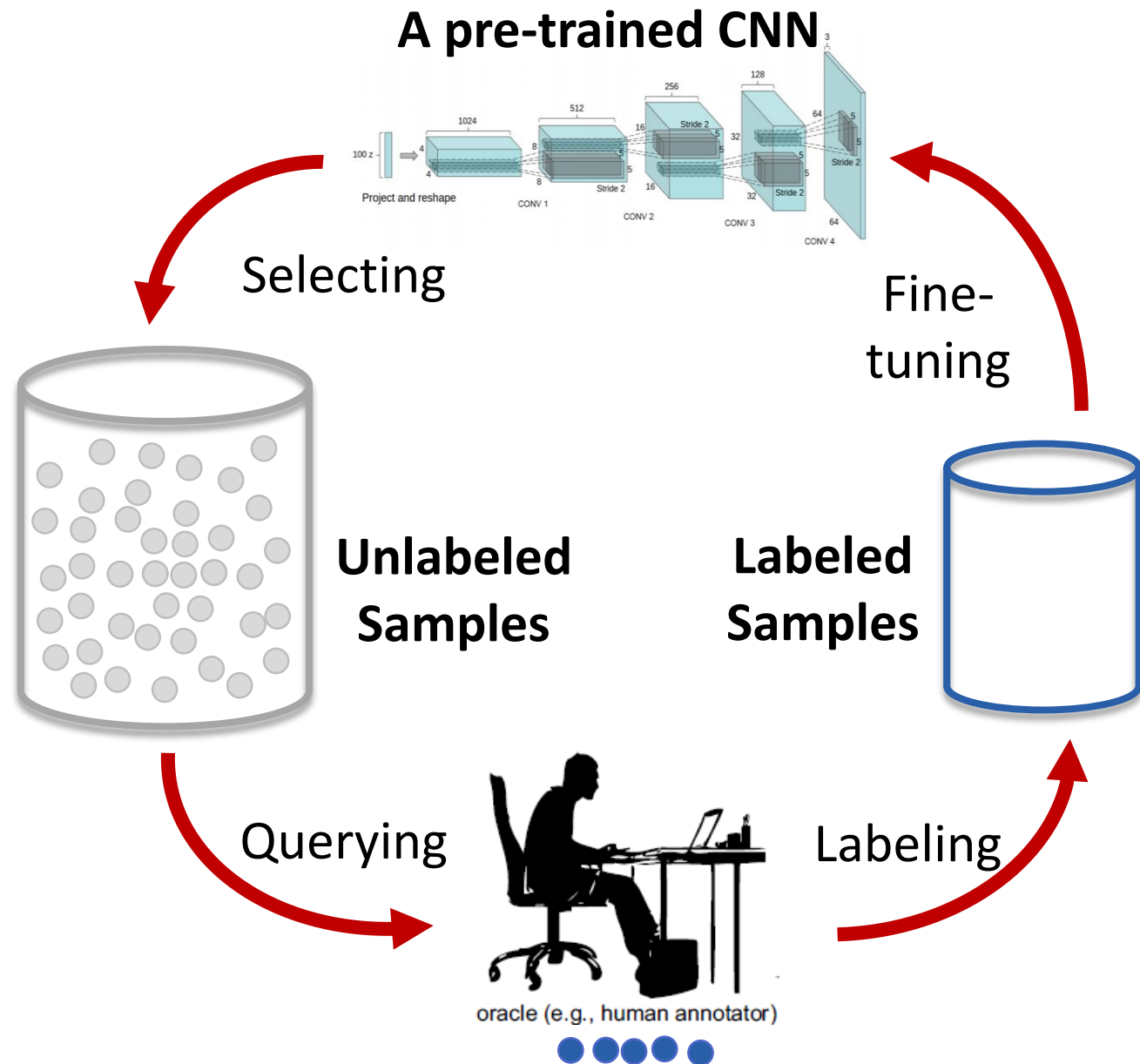
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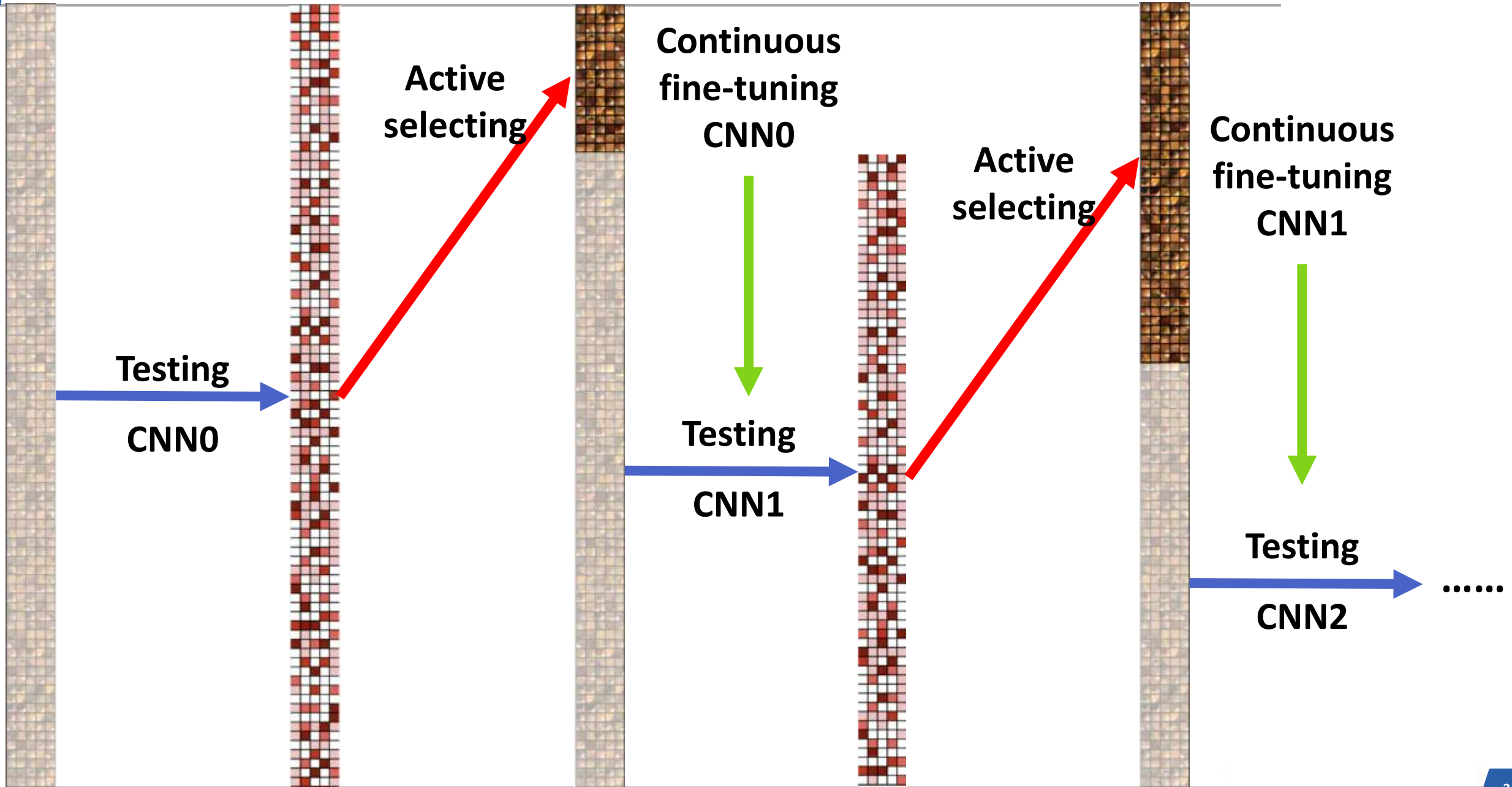
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CVPR2017



- Starting with a completely empty labeled dataset and a pre-trained CNN.
- The pre-trained CNN seeks worthy samples for annotation.
- Then incorporating newly annotated samples in each iteration to fine-tune the CNN continuously.

An Illustration

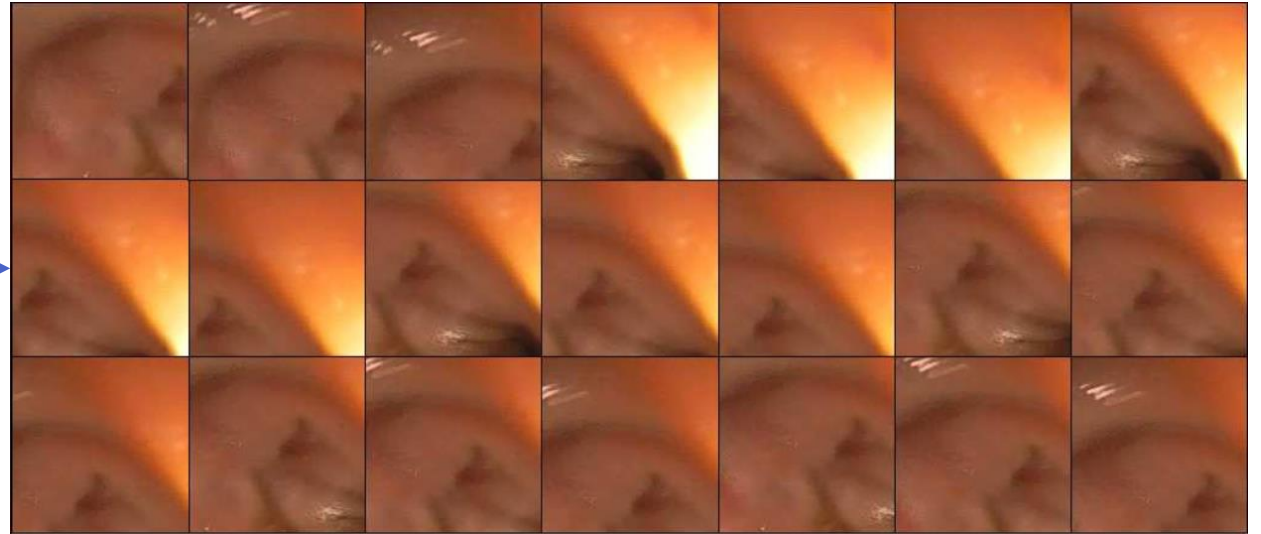


How to design query strategy ?

Each Candidate



Data
Augmentation



Multiple Patches

- These patches generated from the same candidate share the same label, and are naturally expected to have similar predictions by current CNN.
- Their **entropy** and **diversity** provide a useful indicator to the power of a candidate in elevating the performance of the current CNN.

➤ Uncertainty

- Higher uncertainty values denote higher degrees of information.
- Assuming the prediction of patch x_i^j by the current CNN is p_i^j , we define its entropy as:

$$e_i^j = - \sum_{k=1}^{|Y|} p_i^{j,k} \log p_i^{j,k}$$

- Entropy e_i^j denotes the information furnished by patch x_i^j of candidate C_i in the unlabeled pool.

➤ Diversity

- Higher diversity values denote higher degrees of prediction inconsistency among the patches within a candidate.
- Diversity between patches x_i^j and x_i^l of candidate C_i as:

$$d_i(j, l) = \sum_{k=1}^{|Y|} (p_i^{j,k} - p_i^{l,k}) \log\left(\frac{p_i^{j,k}}{p_i^{l,k}}\right)$$

- Diversity $d_i(j, l)$ estimates the amount of information overlap between patches x_i^j and x_i^l of candidate C_i .

How to combine uncertainty and diversity ?

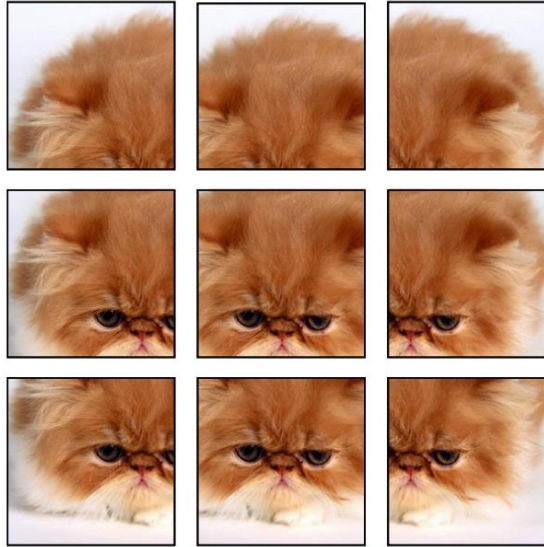
- By definition, all the entries in e_i^j and $d_i(j, l)$ are non-negative.
- Further, $d_i(j, j) = 0, \forall j$
- We combine e_i^j and $d_i(j, l)$ into a signal matrix R_i for each candidate C_i :

$$R_i(j, l) = \begin{cases} \lambda_1 e_i^j & \text{if } j = l, \\ \lambda_2 d_i(j, l) & \text{otherwise} \end{cases}$$

Candidate



Data augmentation



Prediction

0.1	0.1	0.1
0.9	0.9	0.9
0.9	0.9	0.9

- Data augmentation inevitably generates hard samples for some candidates injecting noisy labels.

Handling noisy labels via majority selection

- We compute entropy and diversity by selecting only **a portion of the patches of each candidate** according to the predictions by the current CNN.
- For each candidate C_i compute the average probabilistic predictions of all of its patches:

$$a_i = \frac{1}{m} \sum_{j=1}^m p_i^j$$

Select the **top α** percent patches, if $a_i > 0.5$
Select the **bottom α** percent patches, otherwise

Algorithm 1: Active incremental fine-tuning method.

Input:

$\mathcal{U} = \{\mathcal{C}_i\}, i \in [1, n]$ { \mathcal{U} contains n candidates}

$\mathcal{C}_i = \{x_i^j\}, j \in [1, m]$ { \mathcal{C}_i has m patches}

\mathcal{M}_0 : pre-trained CNN

b : batch size

α : patch selection ratio

Output:

\mathcal{L} : labeled candidates

\mathcal{M}_t : fine-tuned CNN model at Iteration t

Functions:

$p \leftarrow P(\mathcal{C}, \mathcal{M})$ {outputs of \mathcal{M} given $\forall x \in \mathcal{C}$ }

$\mathcal{M}_t \leftarrow F(\mathcal{L}, \mathcal{M}_{t-1})$ {fine-tune \mathcal{M}_{t-1} with \mathcal{L} }

$a \leftarrow \text{mean}(p_i)$ { $a = \frac{1}{m} \sum_{j=1}^m p_i^j$ }

Initialize:

$\mathcal{L} \leftarrow \emptyset, t \leftarrow 1$

```
1 repeat Handling noisy labels via majority selection
2   for each  $\mathcal{C}_i \in \mathcal{U}$  do
3      $p_i \leftarrow P(\mathcal{C}_i, \mathcal{M}_{t-1})$ 
4     if  $\text{mean}(p_i) > 0.5$  then
5       |  $\mathcal{S}'_i \leftarrow$  top  $\alpha$  percent of the patches of  $\mathcal{C}_i$ 
6     else Active candidate selection
7       |  $\mathcal{S}'_i \leftarrow$  bottom  $\alpha$  percent of the patches of  $\mathcal{C}_i$ 
8     end
9     Build matrix  $R_i$  using Eq. 3 for  $\mathcal{S}'_i$ 
10  end Continuous fine-tuning
11  Sort  $\mathcal{U}$  according to the numerical sum of  $R_i$ 
12  Query labels for top  $b$  candidates, yielding  $\mathcal{Q}$ 
13   $\mathcal{L} \leftarrow \mathcal{L} \cup \mathcal{Q}; \mathcal{U} \leftarrow \mathcal{U} \setminus \mathcal{Q}$ 
14   $\mathcal{M}_t \leftarrow F(\mathcal{L}, \mathcal{M}_{t-1}); t \leftarrow t + 1$ 
15 until classification performance is satisfactory;
```

➤ Six variants of AIFT

- AIFT Diversity^{1/4}
- AIFT Diversity
- AIFT Entropy^{1/4}
- AIFT Entropy
- AIFT (Entropy + Diversity)^{1/4}
- AIFT (Entropy + Diversity)

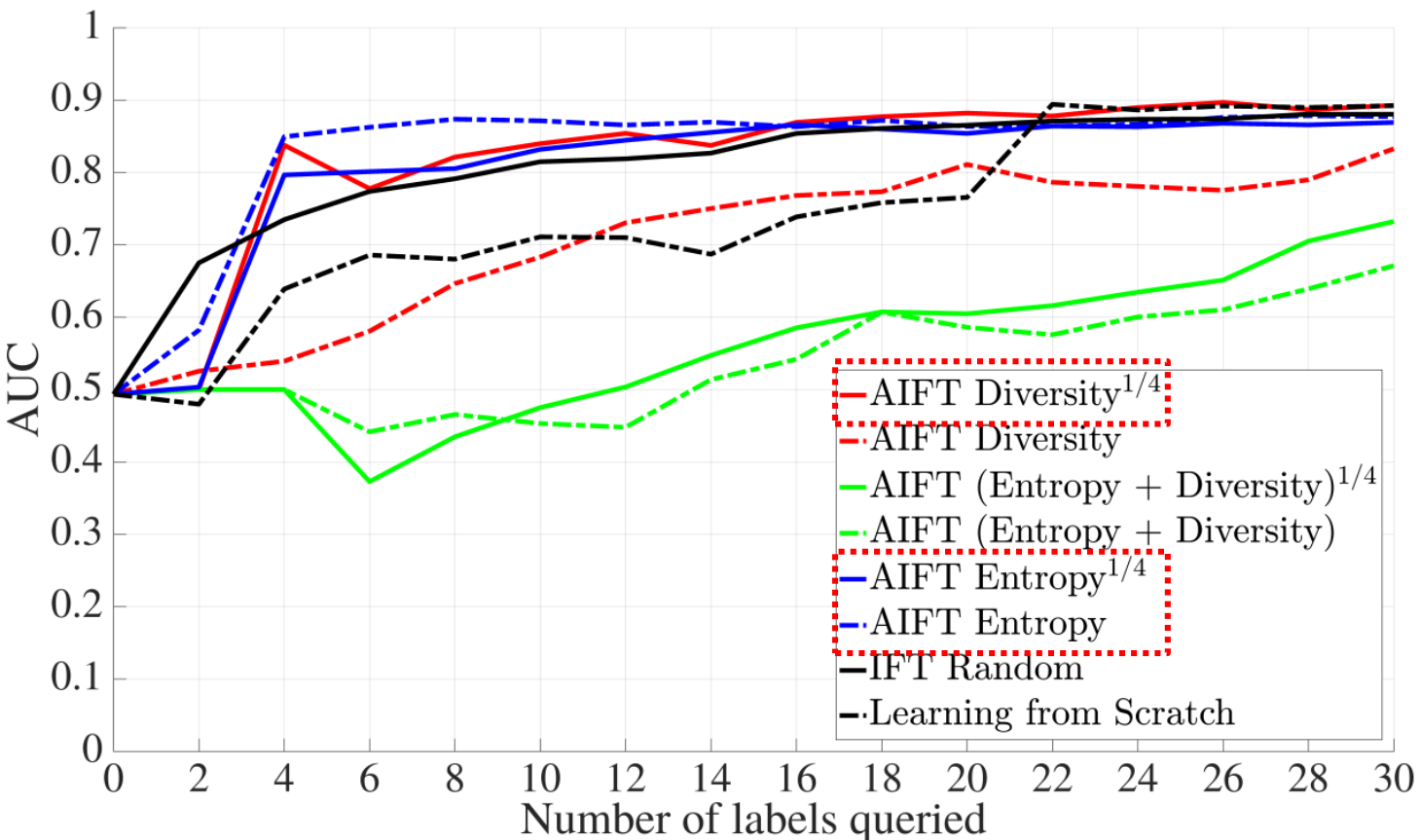
➤ Two compared methods

- IFT Random
- Learning from Scratch

➤ Three applications

- Colonoscopy Frame Classification
- Polyp Detection
- Pulmonary Embolism Detection

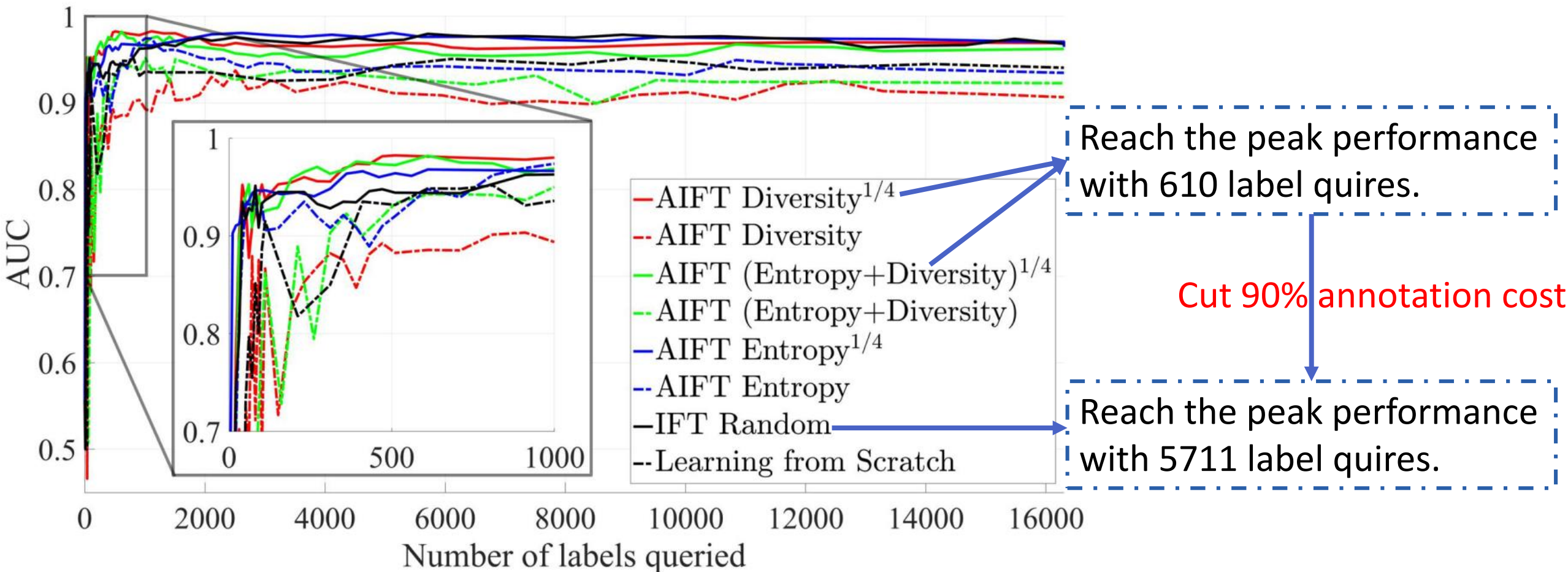
Colonoscopy Frame Classification



- Data: 4000 frames from 6 videos
- Training set: 2000 frames
- Test set: 2000 frames
- Patch: 21 patches for each frame
- Label: informative or non-informative

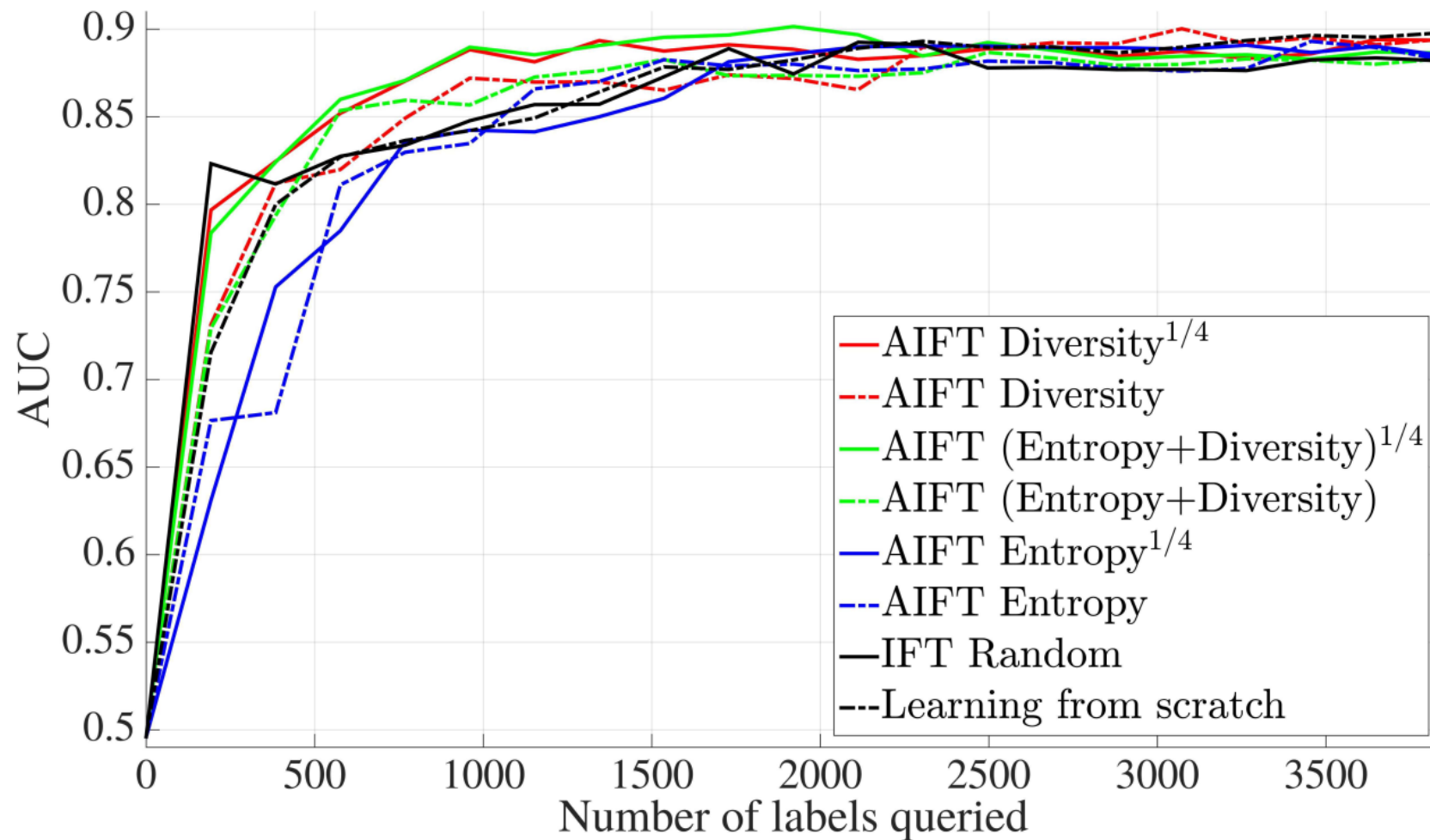
Comparing 8 methods in colonoscopy frame classification

Polyp Detection



It's contributed to the majority selection, which can efficiently select the informative and representative candidates while excluding those with noisy labels.

Pulmonary Embolism Detection



THANKS