



## Abstract

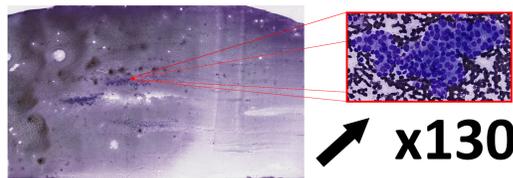
In this paper, we study deep transfer learning as a way of overcoming object recognition challenges encountered in the field of digital pathology. Through several experiments, we investigate various uses of pre-trained neural network architectures and different combination schemes with random forests for feature selection. Our experiments on eight classification datasets show that densely connected and residual networks consistently yield best performances across strategies. It also appears that network fine-tuning and using inner layers features are the best performing strategies, with the former yielding slightly superior results.

## Digital pathology

"Digital pathology incorporates the acquisition, management, sharing and interpretation of pathology information — including slides and data — in a digital environment"

Digital slides:

- **big data:** several multi-gigapixel slides per patient and case
- **high variability:** content, staining, acquisition,...
- **data scarcity:** annotating data is expensive and tedious



Need for **efficient and versatile computer vision methods** that can cope with **data scarcity** !

## Deep transfer learning

Deep learning has a lot of potential for digital pathology but requires...

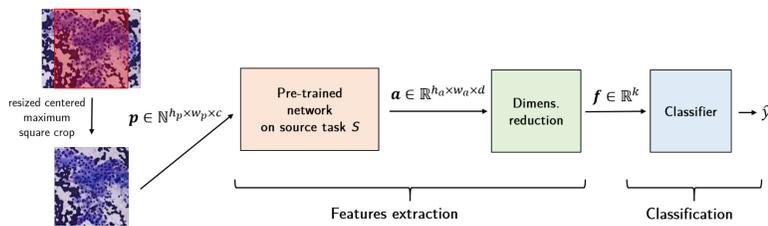
- lots of data
- expensive computing resources (i.e. GPUs)

Deep transfer learning alleviates those requirements:

- Either no network training or training converges faster
- Need less data and fewer resources

There are mainly two ways of using pre-trained networks [7]

- using pre-trained **features off-the-shelf** (OTS)
  - with another classifier such as SVM or random forests
  - can run on CPU
  - requires less data than training from scratch
- **fine-tuning** the networks
  - predicting with the fine-tuned network or using other classifiers on the fine-tuned features
  - need for a GPU but models converge faster
  - requires less data than training from scratch



Because of **data scarcity**, **deep transfer learning** is a **promising approach** for digital pathology.

## Experiments

**Goal:** devising **guidelines and best practices** for deep transfer learning in digital pathology:

- **Fine-tuning vs. OTS features:** which one works better ?
- Which **network** works better ?
- **Where to extract** OTS features ?
- ...

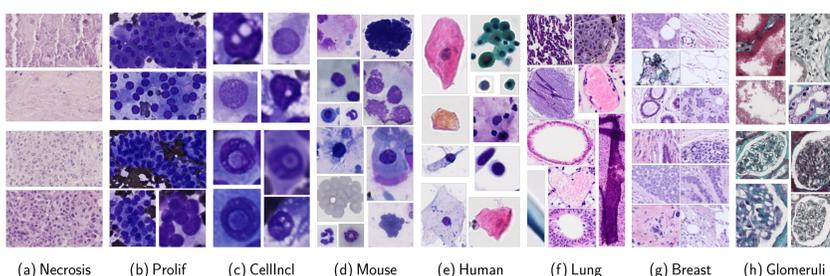
To answer those questions, we have carried out **several experiments**:

- Comparing **OTS features with fine-tuning**
- Comparing **networks**: ResNet50 [4], DenseNet201 [5], VGG16/19 [9], InceptionResNetV2 [10],...
- Several **classifiers for OTS features**: SVM [1], extra-trees (ET) [2],...
- Evaluating features extracted at **different depth** of different networks

## Datasets

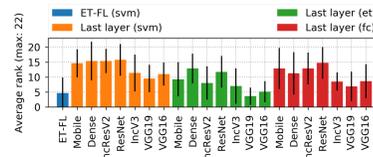
| Dataset           | Domain | Cls | Total  |        |
|-------------------|--------|-----|--------|--------|
|                   |        |     | Images | Slides |
| Necrosis (N)      | Histo  | 2   | 882    | 13     |
| ProlifPattern (P) | Cyto   | 2   | 1857   | 36     |
| CellInclusion (C) | Cyto   | 2   | 3638   | 45     |
| MouseLba (M)      | Cyto   | 8   | 4284   | 20     |
| HumanLba (H)      | Cyto   | 9   | 5420   | 64     |
| Lung (L)          | Histo  | 10  | 6331   | 882    |
| Breast (B)        | Histo  | 2   | 23032  | 34     |
| Glomeruli (G)     | Histo  | 2   | 29213  | 205    |

Our study uses **eight image classification datasets** collected over the years by biomedical researchers and pathologists using the **Cytomine** [8] web application. These contain tissues and cells from human or animal organs (thyroid, kidney, breast, lung,....).



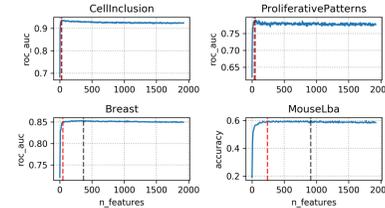
## Results

### Last layer features



Classify **last layer OTS features** with **SVM**, **single layer perceptron** and **extra-trees**. Older networks like VGGs and InceptionV3 are not competitive with more recent networks. SVM is better at classifying features. **Best results with ResNet and SVM**.

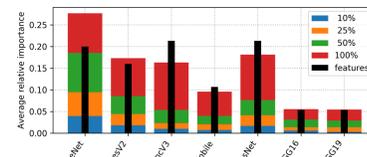
### Feature selection with extra-trees



Feature selection with **recursive feature elimination (RFE)** [3] and **extra-trees** as features rankers. This experiment shows that **most features are actually uninformative or redundant** as RFE excluded on average 92.5% of the features. Our analysis also suggests that the best features are task-dependent.

### Merging features across networks

Merging **last layer OTS features** from all studied networks. An analysis of the **features importances** given by ET shows that **DenseNet features bring 25% of the information** on average. The next most informative networks features are from IncResV2, ResNet, IncV3.



### Inner layers features

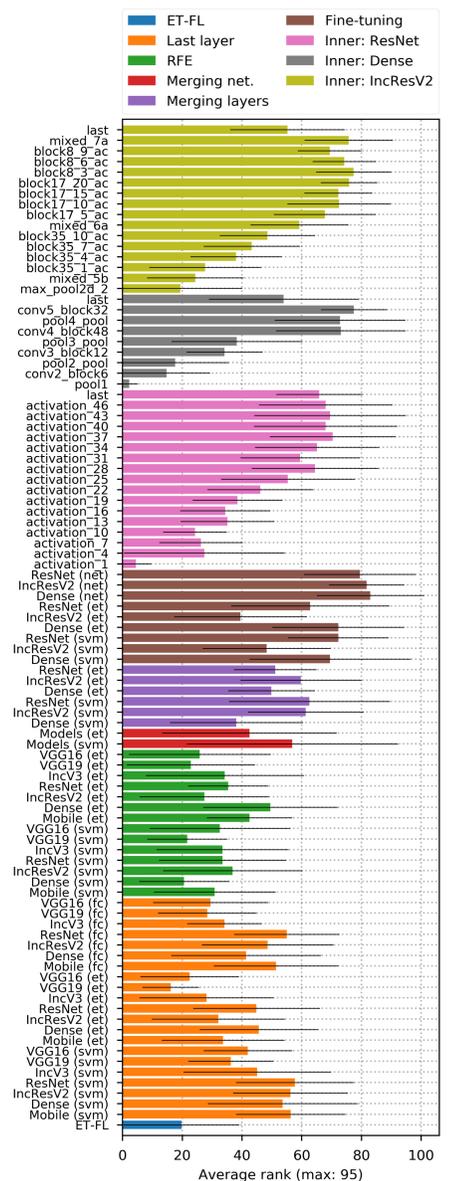
Classify **inner layers features** with **SVM** and **extra-trees**. In all cases, the **last layer features are always outperformed** by features taken from an inner layer of the network. However, the **optimal layer is always located rather at the end** of the network, while the first layers are clearly never competitive.

### Fine-tuning

**Fine-tuning networks**, then either **predicting with the fine-tuned networks** directly or using **fine-tuned features with SVM or ET**. Best performances are obtained by **predicting with the network**. **Fine-tuning usually outperforms all other methods** whatever the network.

### About networks

**ResNet and DenseNet often yield the best performing models** whatever the way they are exploited. Performances obtained with the VGG networks are below those of the others. This is corroborated by the recent findings of [6].



## Conclusion and future works

We have carried out several experiments with 8 histopathology image datasets to **devise guidelines** for **deep transfer learning** in digital pathology.

Main takeaways:

- **Fine tuning** is the best performing method
- **OTS features** often close to fine-tuning and less computationally intensive
- Prefer **inner layers OTS features** to last layers OTS features
- Use **more recent networks** such as DenseNet and ResNet

In the future, we want to study further deep transfer learning and to collect and merge larger annotated biomedical datasets to train networks using a source dataset closer to our target tasks.

## Bibliography

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