

Comparison of deep transfer learning strategies for digital pathology

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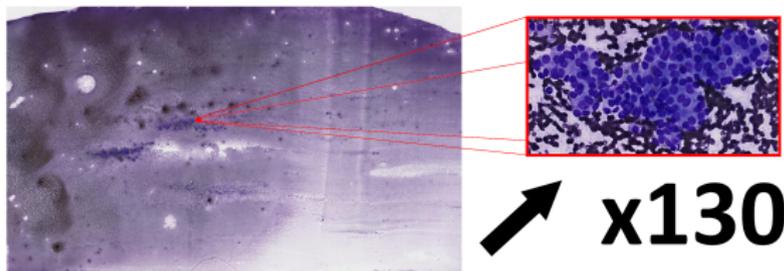
CVMI 2018, poster **FP249**

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**: up to millions of biological objects per multi-gigapixel image
- **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

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

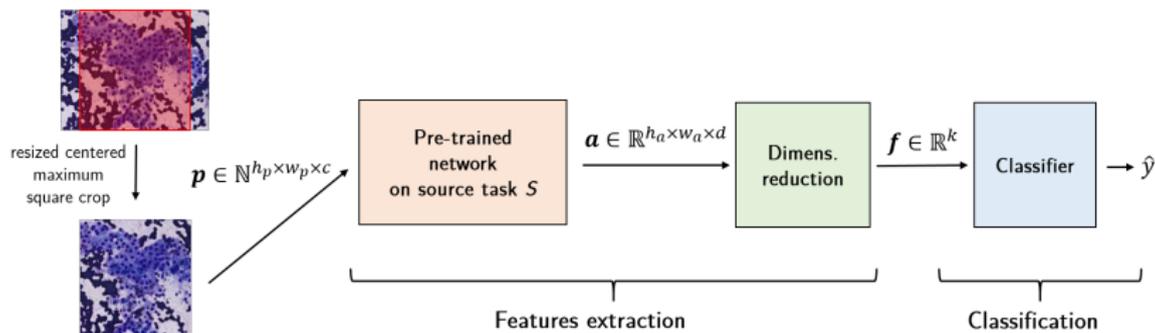
Deep transfer learning alleviates **deep learning** requirements:

- requires **less data**
- requires **less computing resources** and **time**

Using pre-trained networks

There are mainly two ways of using pre-trained networks:

1. using pre-trained features off-the-shelf (OTS)



2. fine-tuning the networks

Deep transfer learning: how to ?

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 ?
- ...

Deep transfer learning: how to ?

We have carried out **several experiments** with **ImageNet** as source task:

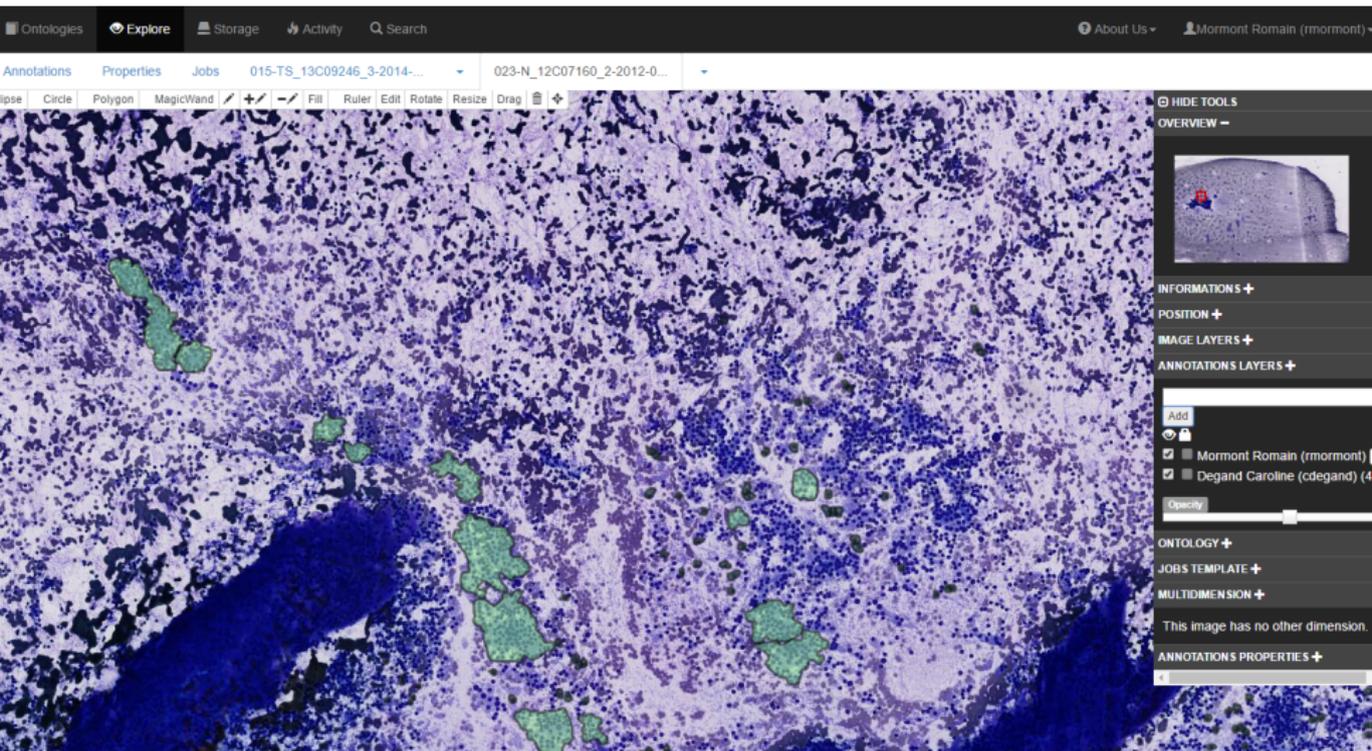
- **OTS** vs. **fine-tuning**
- **Networks**: ResNet50, DenseNet201, VGG16/19, InceptionResNetV2,...
- **Features classifiers**: SVM , extra-trees (ET),...
- OTS features extraction at **increasing depth**
- ...

Cytomine

cytomine is an open-source web-based environment enabling collaborative multi-gigapixel image analysis

(uliege.cytomine.org)

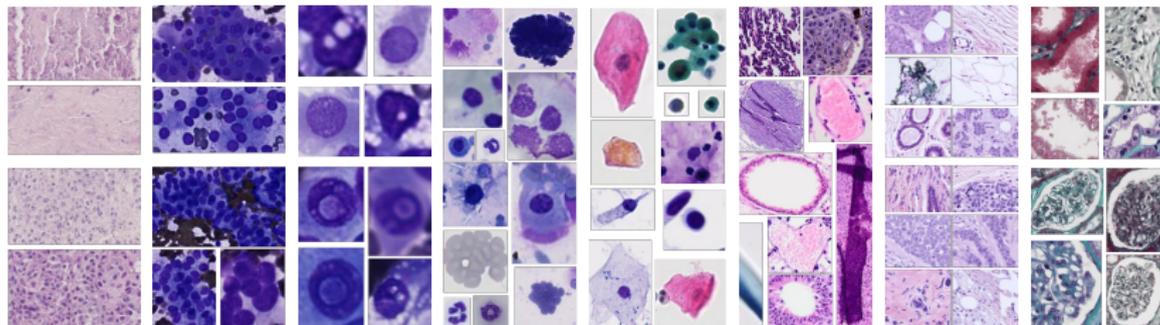
Marée & al., Bioinformatics; 2016



Datasets

8 image classification datasets.

Dataset	Domain	Cls	Total	
			Images	Slides
Necrosis (N)	Histo	2	882	13
ProliferativePattern (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



Results

Fine-tuning is the best performing strategy

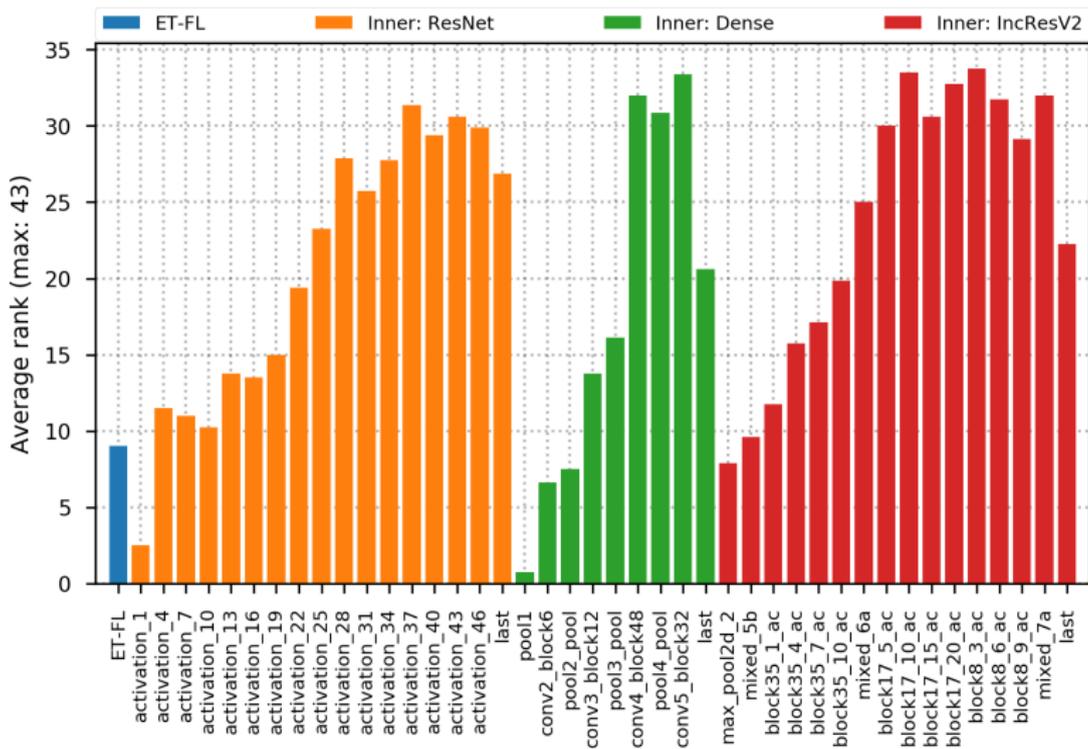
Fine-tuning is often the best performing method
... but OTS features are close on most problems and less computationally expensive !

Strategy	Datasets							
	Cell	Prolif	Glom	Necro	Breast	Mouse	Lung	Human
Baseline (ET-FL)	0.9250	0.8268	0.9551	0.9805	0.9345	0.7568	0.8547	0.6960
Last layer	0.9822	0.8893	0.9938	0.9982	0.9603	0.7996	0.9133	0.7820
Feat. select.	0.9676	0.8861	0.9843	0.9994	0.9597	0.7438	0.8941	0.7703
Merg. networks	0.9897	0.8984	0.9948	0.9864	0.9549	0.8169	0.9155	0.7928
Merg. layers	0.9808	0.8906	0.9944	0.9964	0.9639	0.7941	0.9268	0.7977
Inner ResNet	0.9748	0.8959	0.9949	0.9964	0.9664	0.8131	0.9291	0.8113
Inner DenseNet	0.9862	0.8984	0.9962	0.9917	0.9699	0.8012	0.9268	0.7967
Inner IncResV2	0.9873	0.8948	0.9962	0.9982	0.9720	0.8137	0.9234	0.7713
Fine-tuning	0.9926	0.8797	0.9977	0.9970	0.9873	0.8727	0.9405	0.8641
Metric	Roc AUC				Accuracy (multi-class)			

Best in **bold**, second best in *italic*

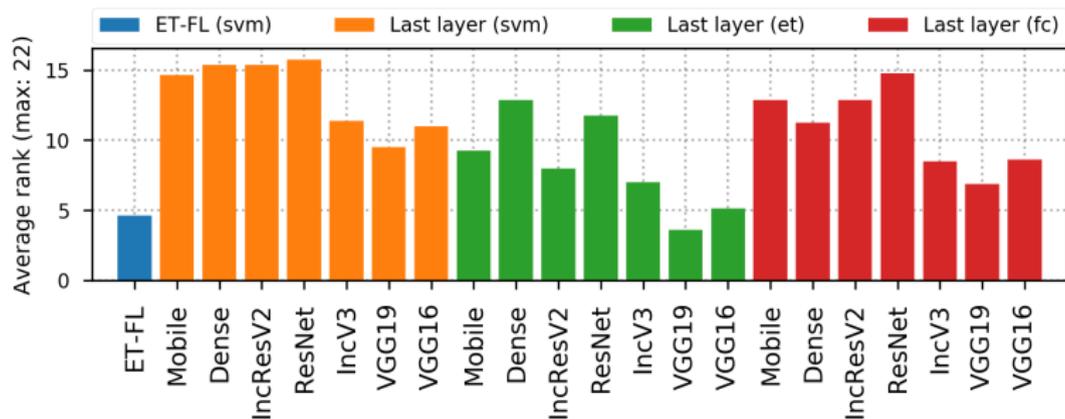
Results

When working with OTS features, use some inner layer features



Results

More recent networks like DenseNet or ResNet work better



See also: Kornblith, S., Shlens, J., & Le, Q. V. (2018). *Do Better ImageNet Models Transfer Better?*. arXiv preprint arXiv:1805.08974.

Conclusion

Main takeaways:

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

Thank you !

Meet me at poster **FP249** for more information !