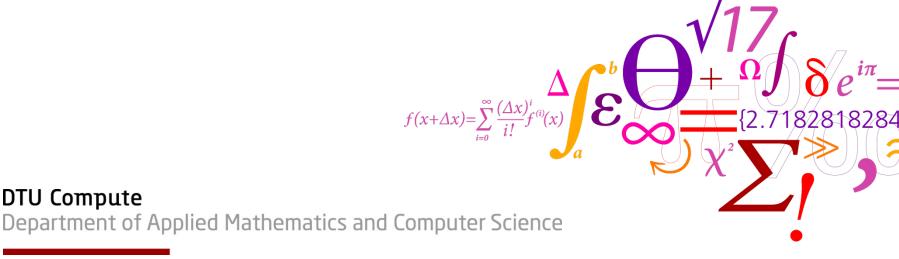
CAMELYON 2017 TEAM DTU

Jeppe Thagaard, MSc student Biomedical Engineering, Technical University of Denmark

Thesis supervisors:

DTU Compute

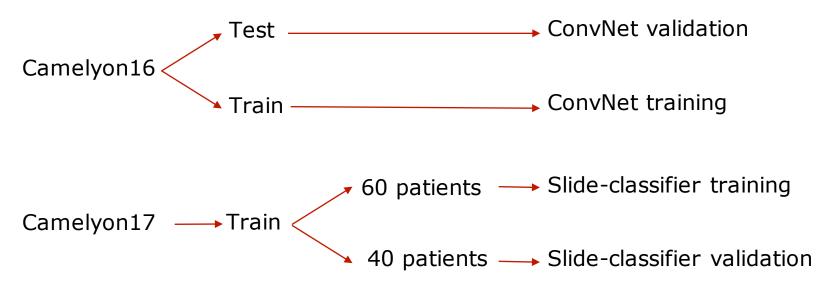
Anders B. Dahl PhD, Section for Image Analysis & Computer Graphics Søren Hauberg PhD, Section for Cognitive Systems





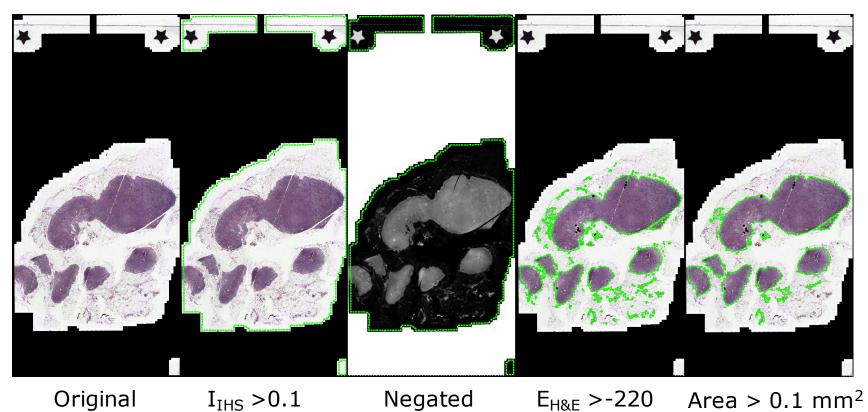
Overview

- Tissue detection using Visiopharm Integrator System (VIS)
- Patch-classification with convolutional neural networks
- Post-processing and high-level features with VIS
- Slide-classification with Random Forest classifier
- Data split:





Tissue detection



Eosin-channel

 $R_{RGB} > 50$

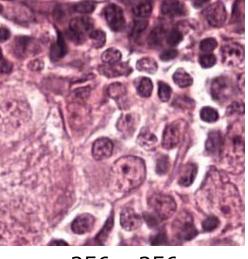
Model



Patch size: How much detail?

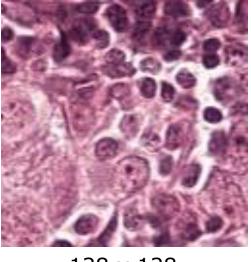
- Same receptive field in tissue, same performance
- 128 x 128 pixels at 20X magnification
- Smaller input size less computational

40X



256 x 256





128 x 128

Pre-processing

- Sampling & model iteration
- Sampling in 2 iterations on Cam16 training

Sampling

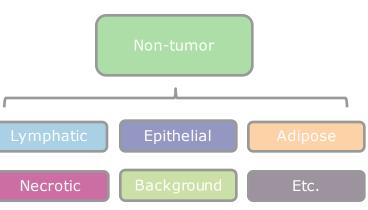
Model

- Uniform random sampling (URS)
- 2000 tumor per slide
- 1000 normal per slide
- 2^{nd} :

1st:

Overview

- Run inference and obtain errors (N = 347000)
- Resample N tumor patches on tumor slides with URS
- Resample $\frac{N}{2}$ normal patches on normal slides and and $\frac{N}{2}$ normal patches on tumor slides with URS
- More difficult dataset: 350K tumor, 700K normal



Post-processing

Flip

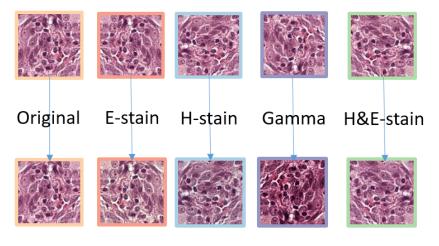


Pre-processing

Motivation

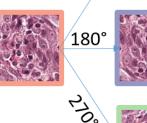
Overview

- Rotational invariant
- Stain/color invariant
- 4 spatial transformation
- Augment more stain variation digitally than available in the training data
 - H&E de-convolution
 - Randomly vary stains with $\pm 10\%$
 - Convolve back to RGB
 - Stain variation
- Gamma-correction
 - Random γ -value [0.4-1.6]
 - Scanner setting variations
- 5x amount of data: 5.2M patches







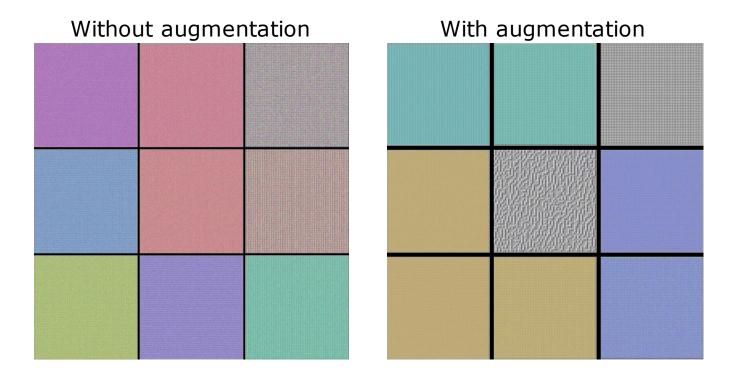




Model



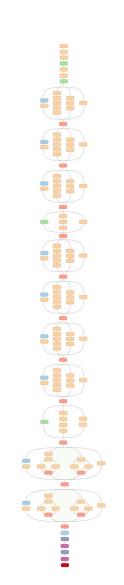
Filters of first convolutional layer



Visualization: Weight activation maximization using backpropagation

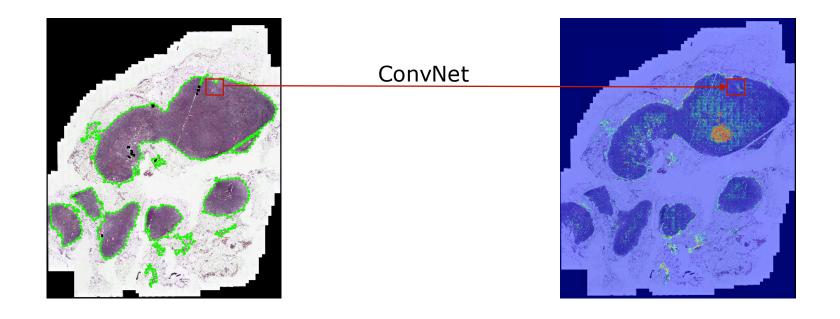
ConvNet details

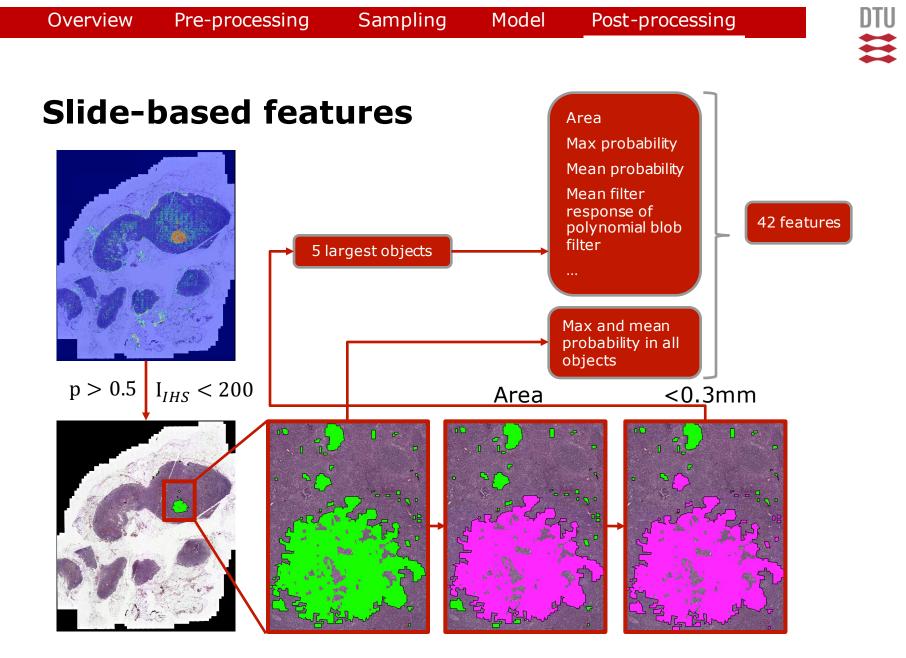
- Slightly changed Inception-V3 (C. Szegedy 2015)
- Extra fully-connected layer
- Dropout (p = 0.2) to prevent overfitting
- Optimizer
 - SGD with Nesterov's momentum (0.9)
- Mini-batch size: 32
- Learning rate schedule:
 - Initial: 0.1
 - Dropped 50% every 125K gradient update
- Single NVIDIA GTX 1080 GPU
- Patch-based performance (full sampling)
 - Camelyon16 test: 96.6 % accuracy



WSI inference

- Sliding field-of-view (FOV)
- Step-length: 128 pixels
- Each FOV is assigned a probability $p \in [0,1]$
- No ensemble average due to limited time

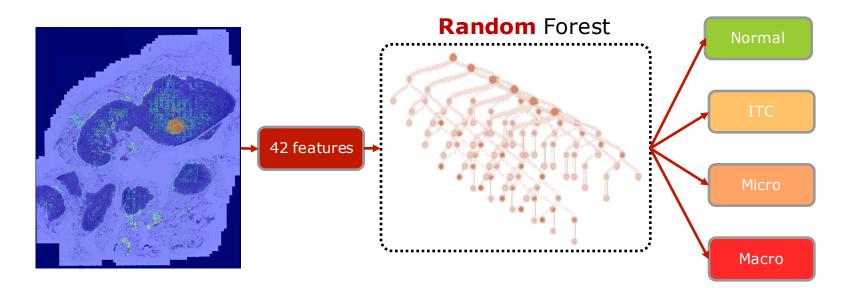




CAMELYON17 - TEAM DTU 18/04/2017



Slide classification



- Hyperparameters tuned on Camelyon17 training spilt using 3-fold cross-validation
- Tested on Camelyon17 validation split
- Features with highest feature importance:
 - Maximum probability and area of largest object



Patient classification

- Rule-based scheme only
- Official pN-staging system in Camelyon17

pN-stage	Rule
pN0	No micro-metastases or macro-metastases or ITCs found.
pN0(i+)	Only ITCs found.
pN1mi	Micro-metastases found, but no macro- metastases found.
pN1	Metastases found in $1-3$ lymph nodes, of which at least one is a macro-metastasis.
pN2	Metastases found in 4–9 lymph nodes, of which at least one is a macro-metastasis.



2.7182818284

THANK YOU

Contact

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- LinkedIn: www.linkedin.com/in/jeppethagaard

DTU Compute

Department of Applied Mathematics and Computer Science

 $f(x+\Delta x)=\sum_{i=1}^{\infty}$