

An approach for glomeruli detection in multisite digital pathology

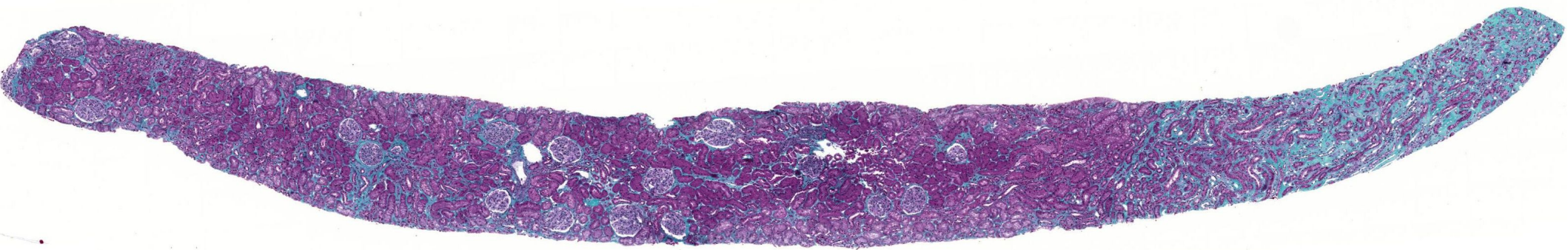
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**ISBI
2016
Prague**



Large biomedical studies have to be multicentric

Benefits :

- Large number of participants
- Wider range of population groups (<> genetic, environmental, demographic factors)

→ Increase the generalizability of the study



Challenges of *multisite* studies in digital pathology

Big data management

- High-resolution images (100 K x 100 K)
- Large number of whole-slide images (hundreds...thousands...)
- Very large number of « objects » (e.g. cells,...) to detect, quantify, and store
- Distributed and poorly structured

Image variabilities require robust pattern recognition

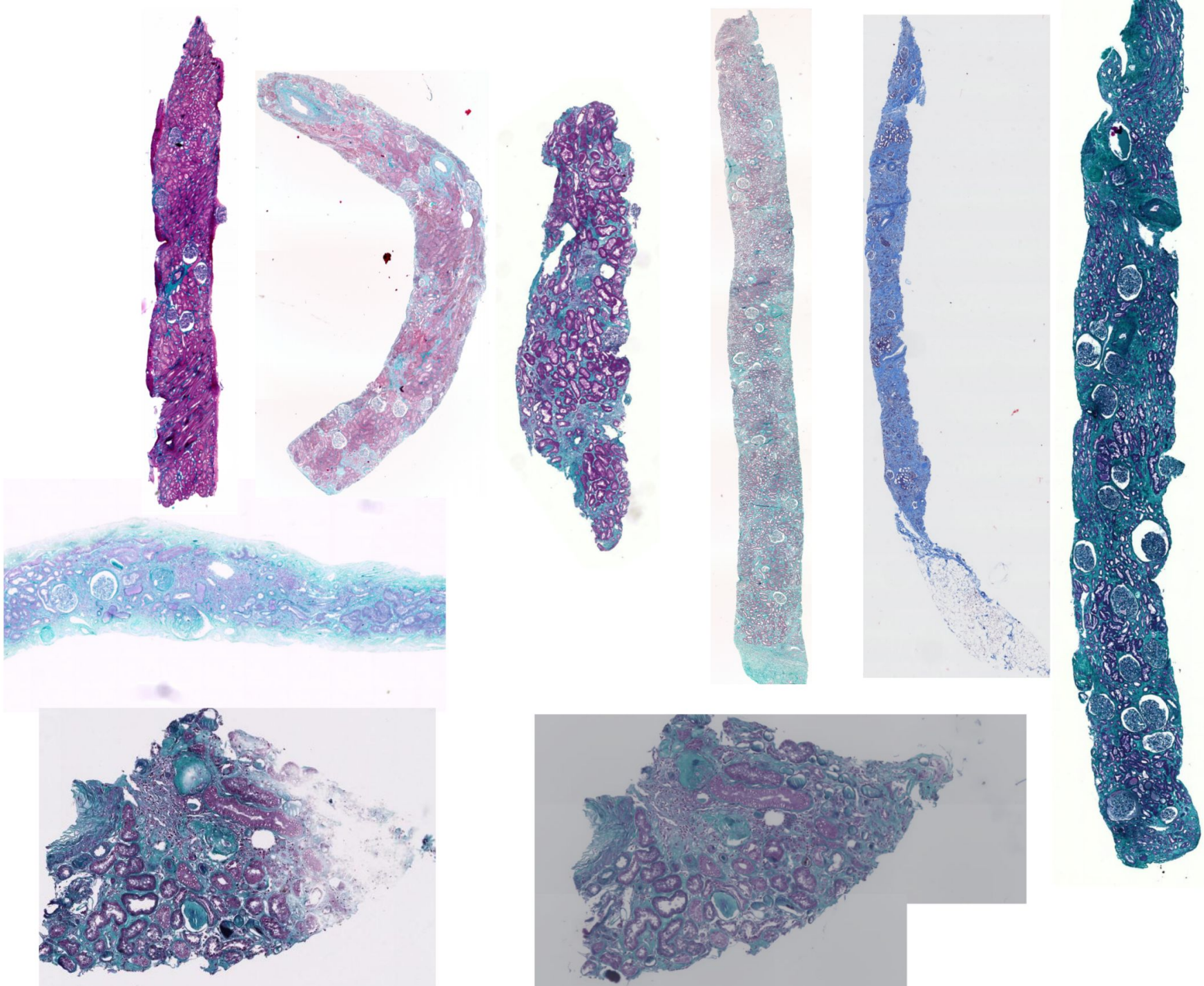
- Tissue (healthy or not, age, ...)
- Preparation protocols (staining, ...)
- Acquisition setups (slide scanners, ...)

Collaborative software platforms:

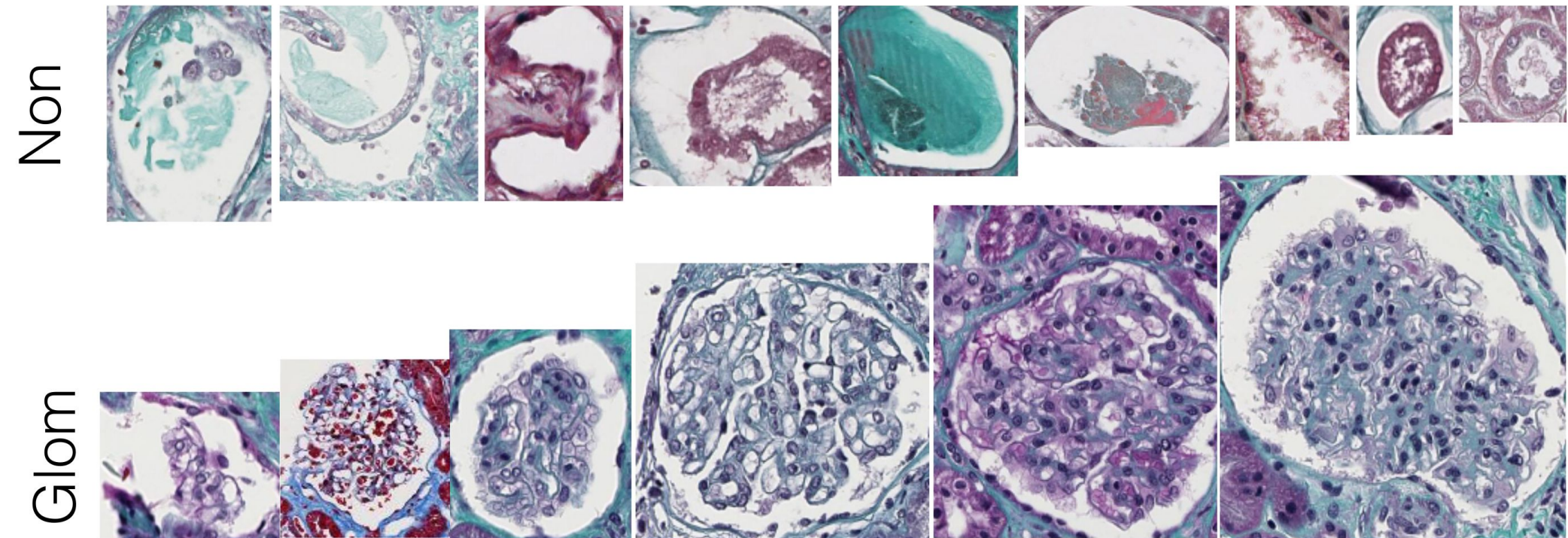
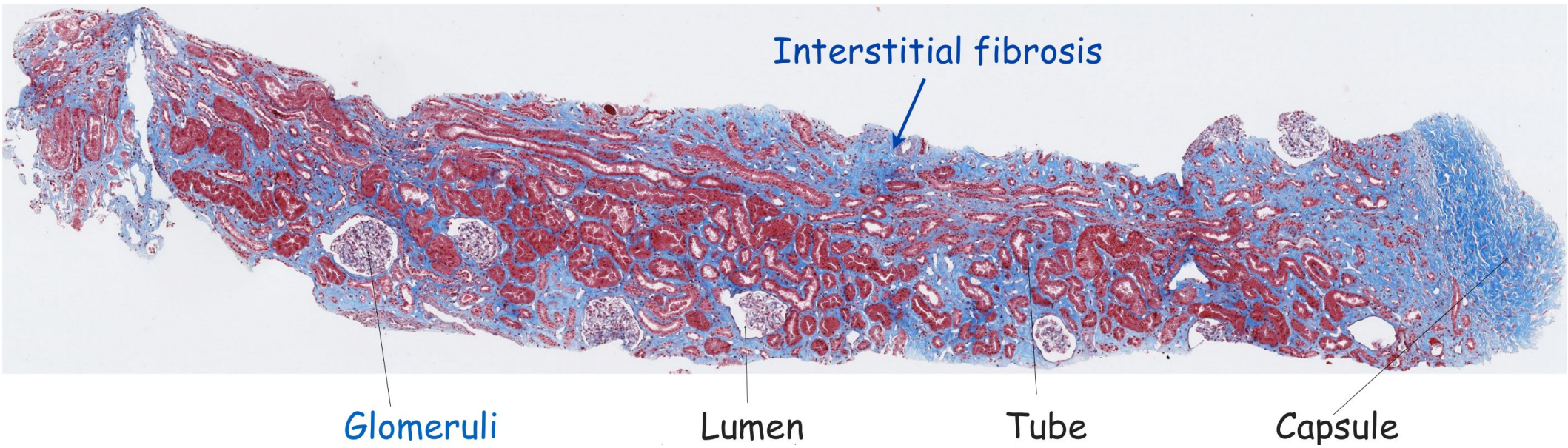
- Groups of pathologists
 - Ground-truth creation
 - Validation of algorithms
- Groups of image analysts
 - Image processing/analysis
 - Machine learning

...

Challenges of glomeruli detection (1/2)



Challenges of glomeruli detection (2/2)

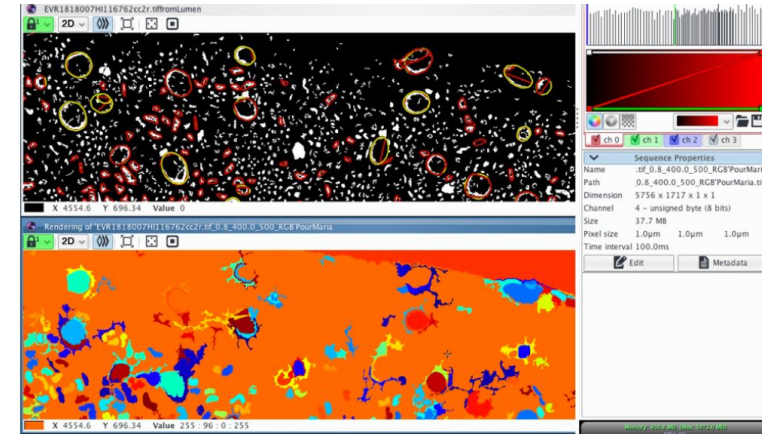


Our approach overview



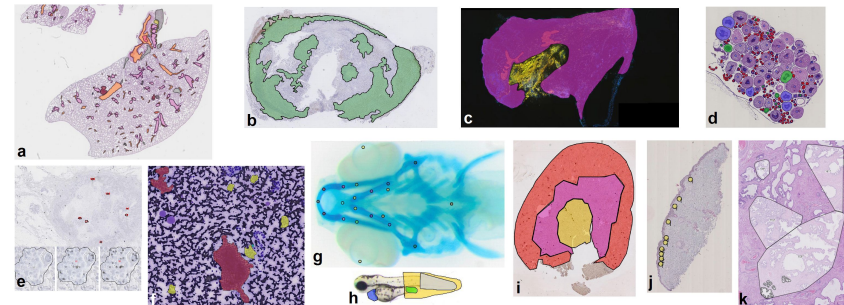
(icy.bioimageanalysis.org ; Nature Methods, 2012) :

- Image analysis
 - Glomeruli candidate detection (Transplantation, 2011)
 - Color normalization/transfer



(www.cytomine.be ; Bioinformatics, 2016) :

- Database to organize whole-slide images (various formats) and semantic annotations
- A rich internet application : data sharing and collaborative modes (proofreading)
- Supervised machine learning algorithms for object recognition
- Open API and software templating mechanisms to add algorithms



Our approach overview



(icy.bioimageanalysis.org)

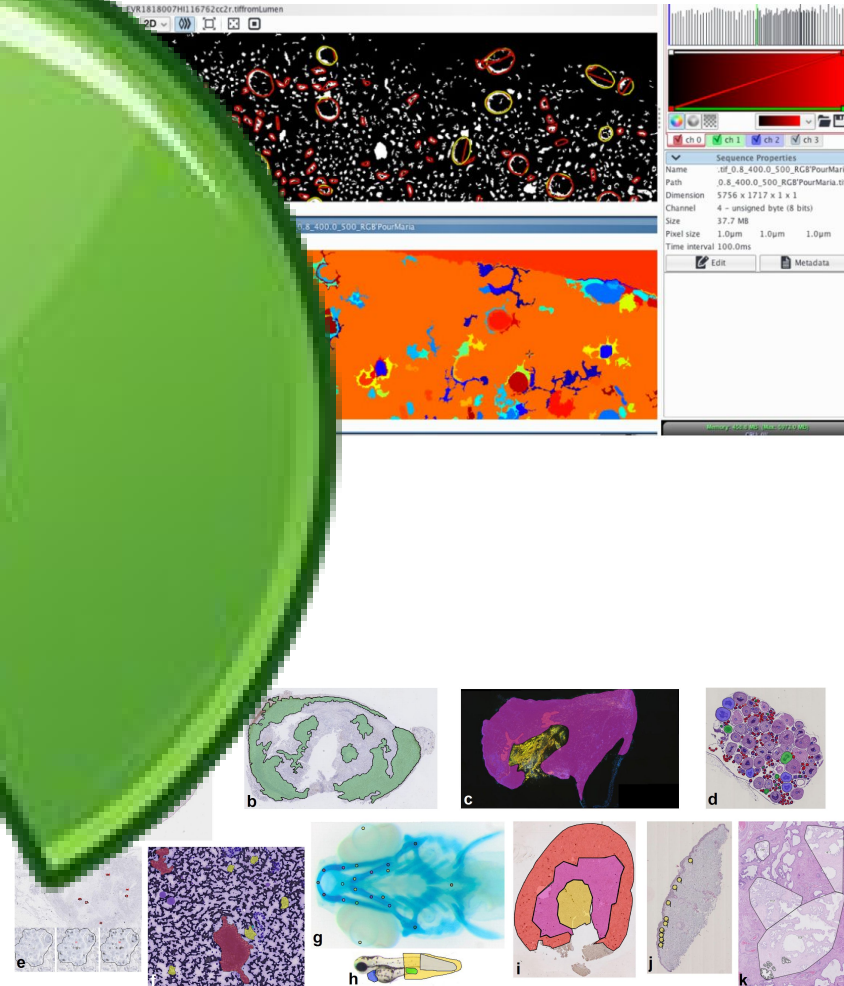
- Image analysis
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cytomine

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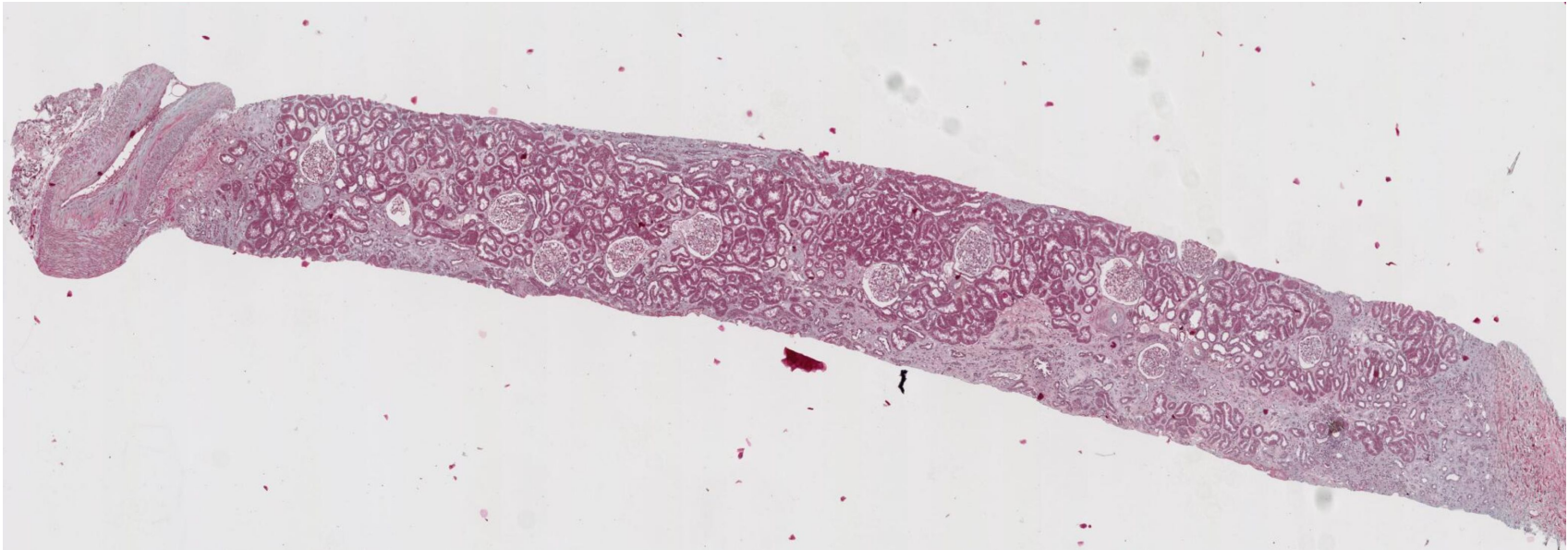
open source



Proposed approach :

1. Segmentation of candidates within whole slides using image analysis

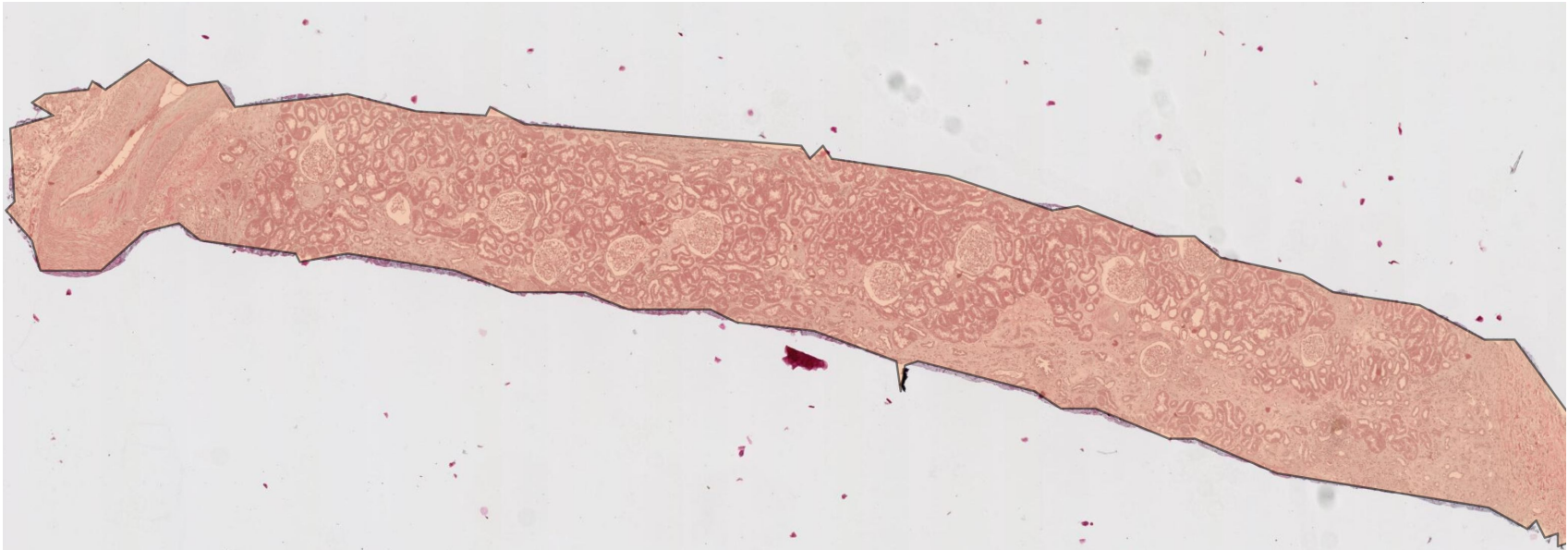
Section detection : Otsu thresholding



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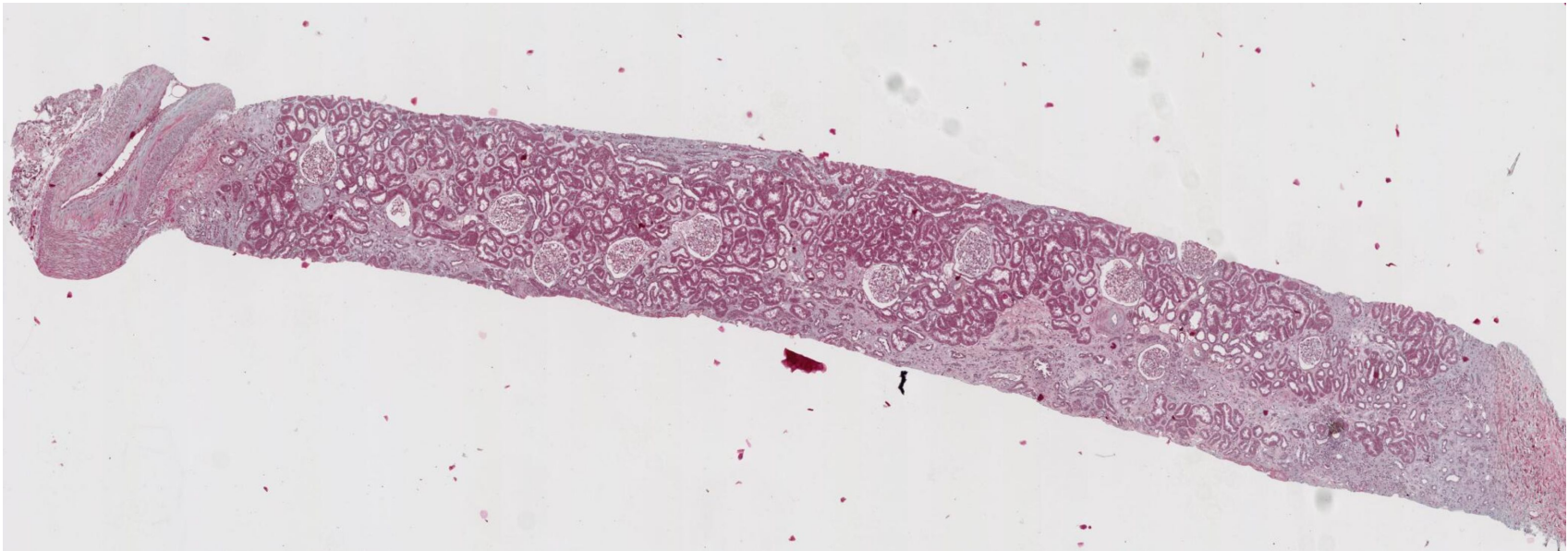
Section detection : Otsu thresholding



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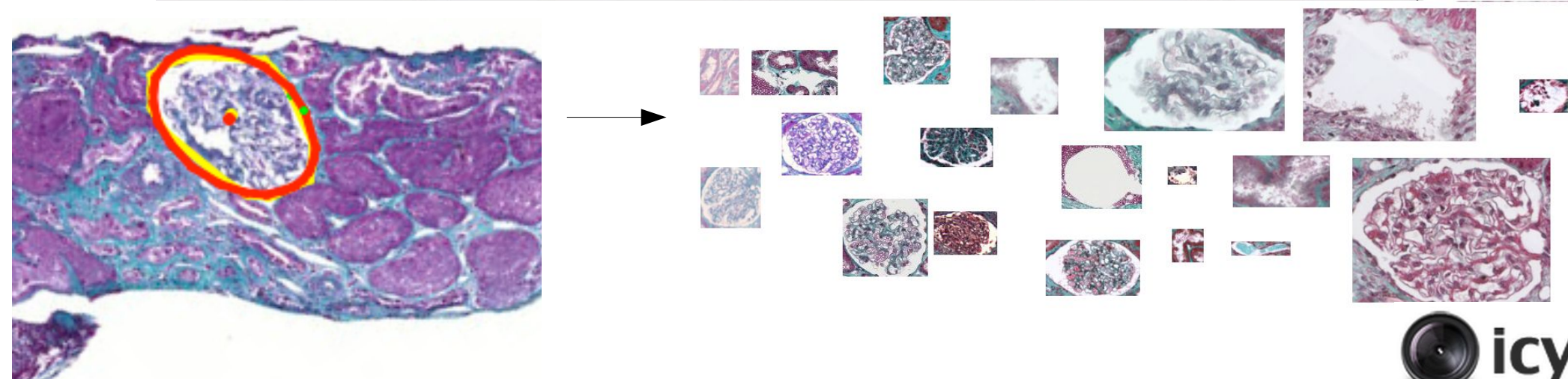
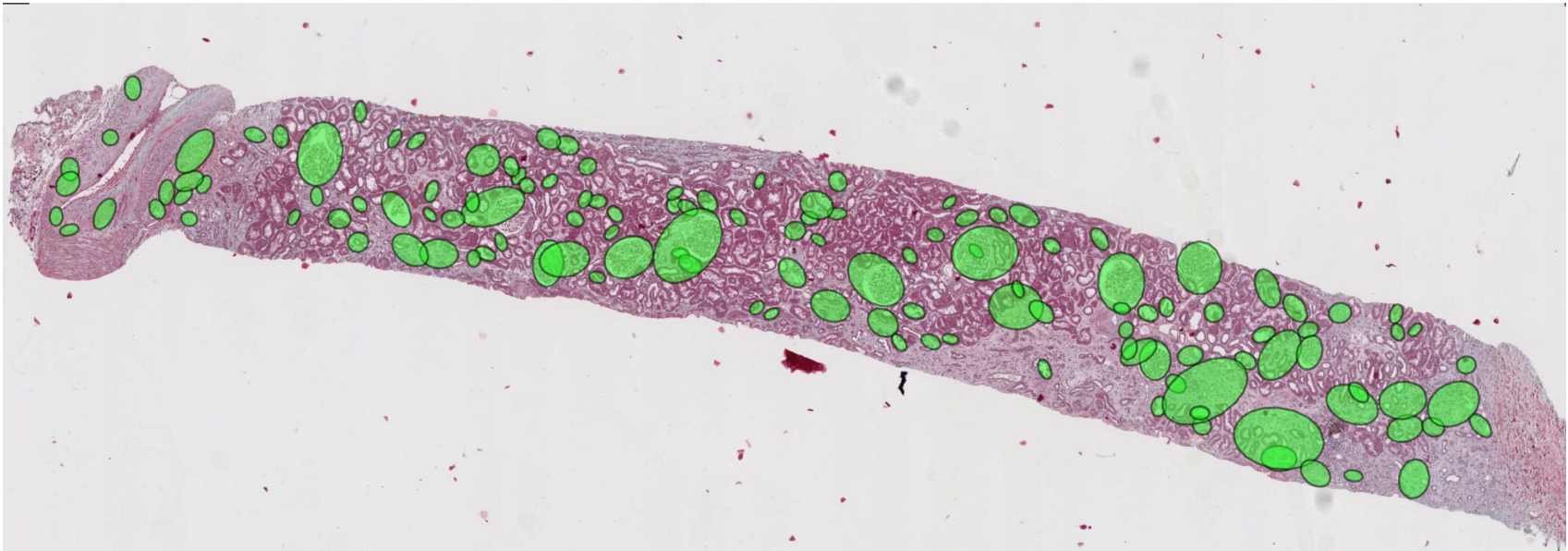
Candidate glomeruli detection : Lumen extraction (contour + convex Hull)
+ Ellipse detection (Fitzgibbon, 1999)
+ Superpixel segmentation (Felzenszwalb, 2004)



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Proposed approach :

2. Classification of candidates using supervised algorithms

Semi-automated ground-truth creation: candidates generated by Step 1 are classified manually using Cytomine web proofreading UI → training set

Cytomine Dashboard Projects Explore Storage Activity

PASTEUR-LAIQ-EVEROLD Images Annotations Properties Jobs Configuration Review

Review for project PASTEUR-LAIQ-EVEROLD

You are reviewing image EVR01013V0 - 2015-01-05 19.44.58.ndpi.

User: Icy_Glomeruli_Finder 2015-06-19 16h52 Term: All

Explore

Icy_Glomeruli_Finder 2015-06-19 16h52: 99 / 240 reviewed
Icy_Section_Finder 2015-06-19 16h52: 0 / 6 reviewed

Annotation details

Created by Icy_Glomeruli_Finder
Date 2015-06-19 16h53
Term associated Icy_Glomeruli_Finder has associated Glomerule

Glomerule

Section

NON

GLOM

cytomine

Proposed approach :

2. Classification of candidates using supervised algorithms

Generic image classification algorithm

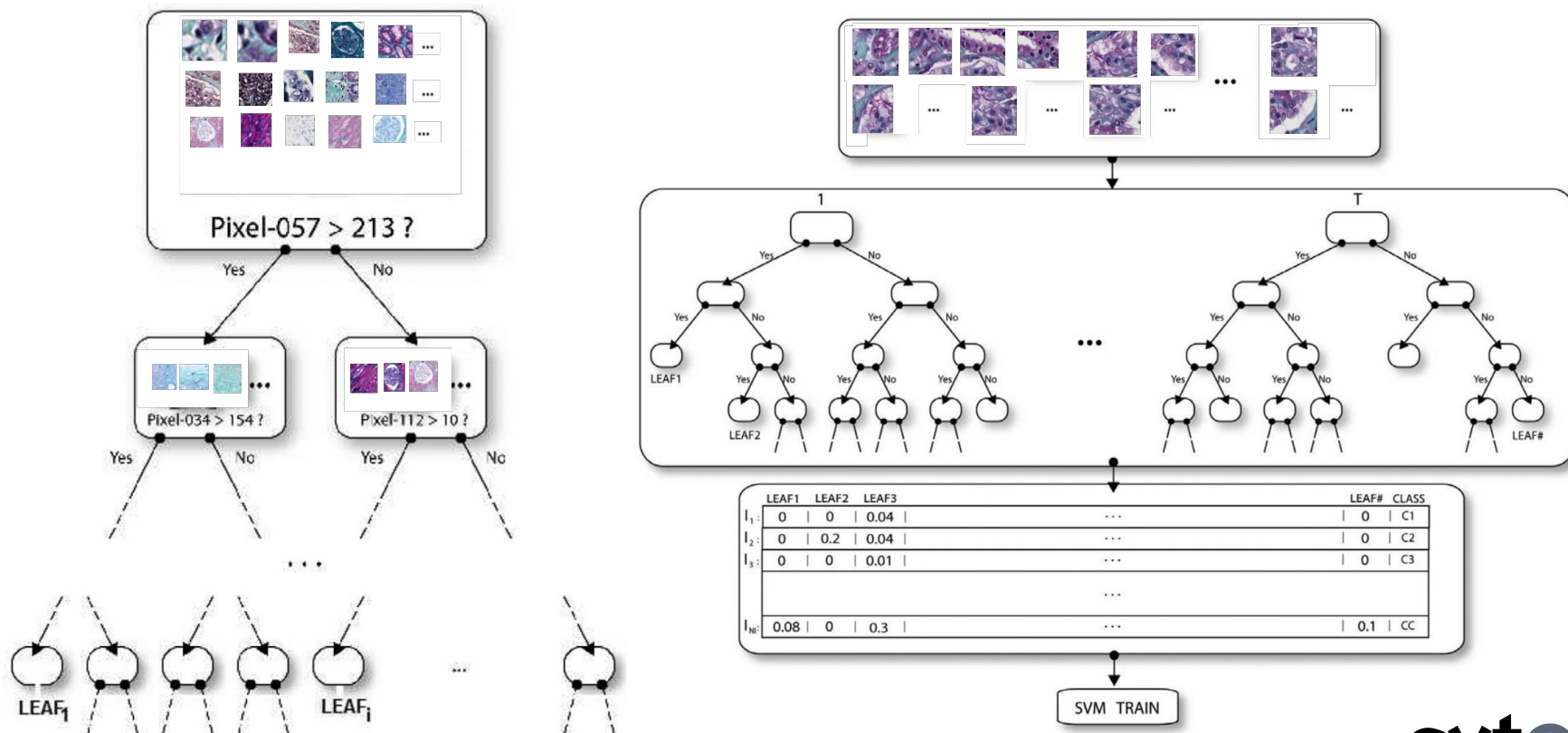
(Marée et al., Pattern Recognition Letters 2016):

Random subwindows (patches) extracted in glom/non-glom images

Described by raw pixel values

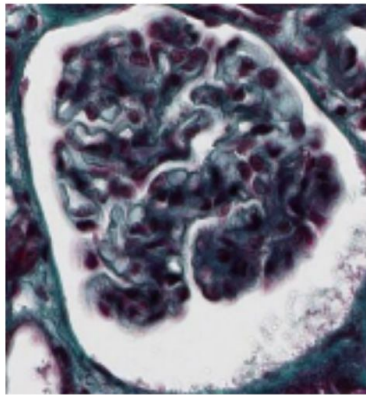
Extremely Randomized Trees for feature learning

Final linear SVM classifier

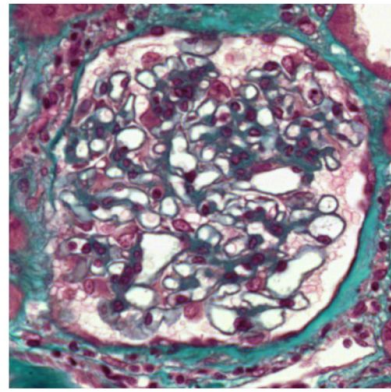


Proposed approach :

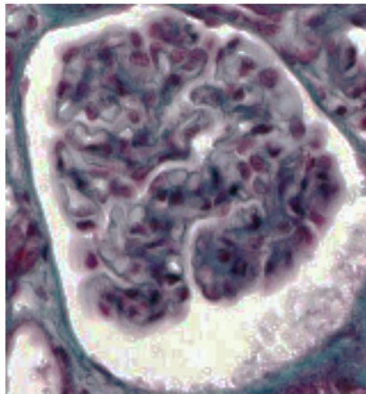
2. Classification ... + color normalization



Source

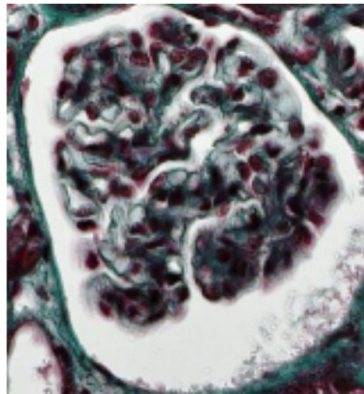


Target

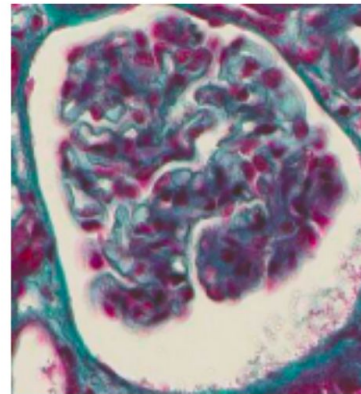


HS

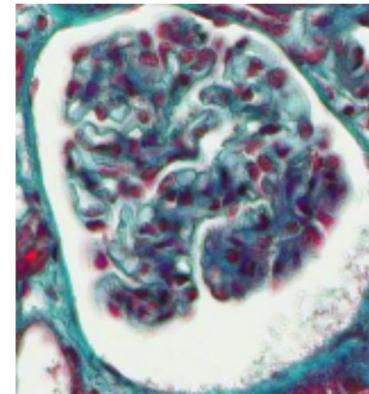
HS : Histogram equalization



ACE



RH



MM

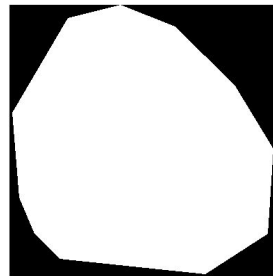
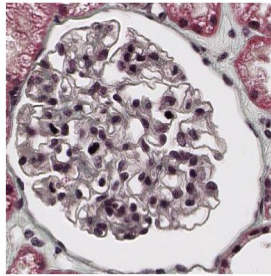
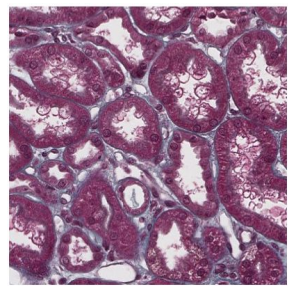
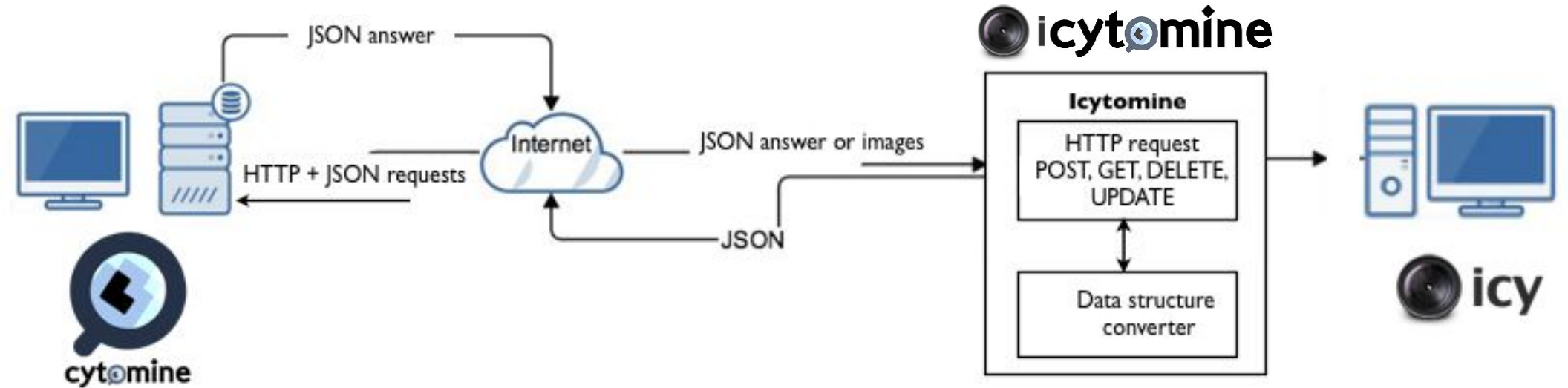
ACE : Automatic color enhancement (Getreuer, IPOL 2012)

RH: Color transfer (per-pixel basis, equalization of μ and σ in the perceptual color space)
(Reinhard et al., IEEE Computer Graphics and Applications 2011)

MM : SVD based approach to estimate color deconvolution matrices (Macenko et al., ISBI 2009)



Implementation based on web services



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{"class": "be.cytomine.ontology.UserAnnotation", "id": 95189319, "created": "1390233213025", "location": "POLYGON ((83409.75 96490.375, 83412.75 96492.375, 83412.75 96481.375, 83411.75 96476.375, 83408.75 96473.375 ... 96489.375, 83409.75 96490.375))", "image": 95062074, "project": 94152784, "container": 94152784, "user": 94115159, "area": 8953.0, "perimeterUnit": "mm", "areaUnit": "micron\u00b2", "perimeter": 1.0, "centroid": {"x": 83426.59663300458, "y": 96382.22057123308}, "term": [95062790], "cropURL": "http://beta.cytomine.be/api/userannotation/95189319/crop.jpg", "reviewed": true}
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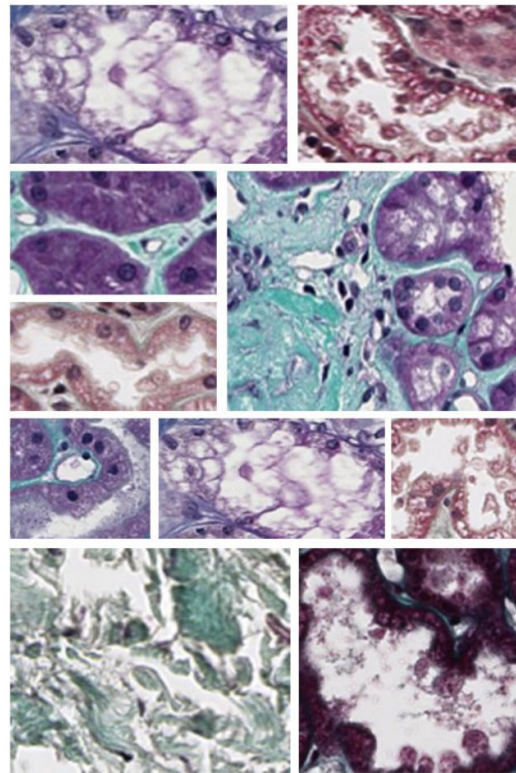
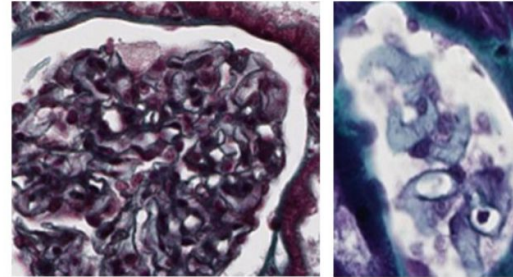
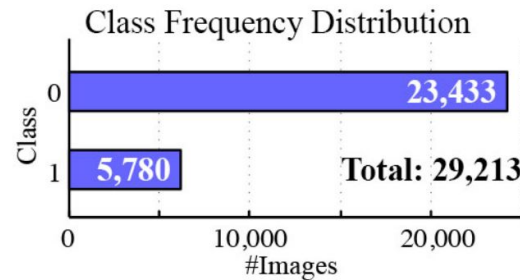
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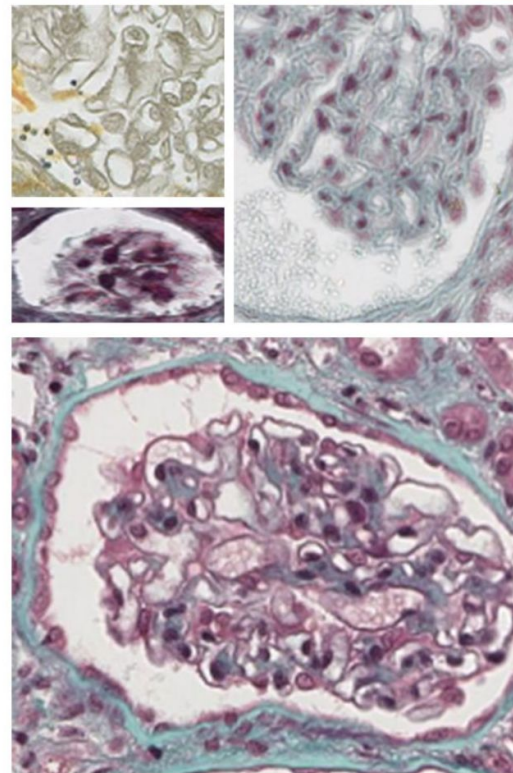
[http://\\$HOST/api/userannotation/95189319.json](http://$HOST/api/userannotation/95189319.json)

Dataset and evaluation protocol

Dataset : 200 whole-slide images, 29213 annotated objects (ground-truth)



Non-glomerular Structures (Class 0)



Glomeruli (Class 1)

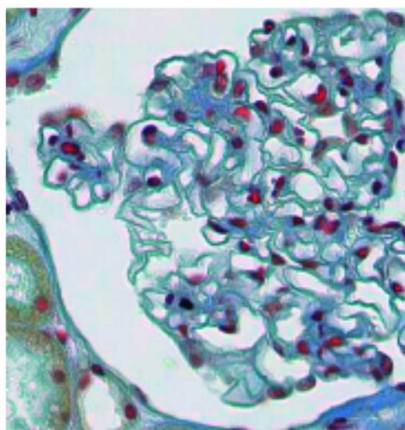
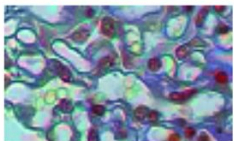
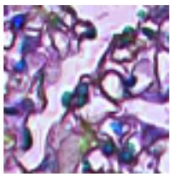
Training set: Objects from 100 slides (2927 glomeruli + 13648 non-glom)

Testing set: Objects from 100 unseen slides (2853 glomeruli + 9785 non-glom)

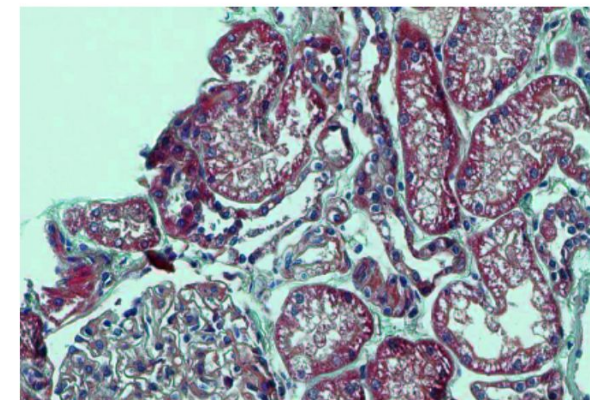
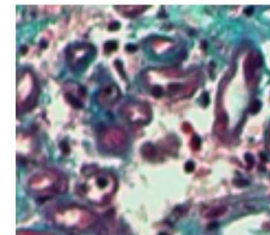
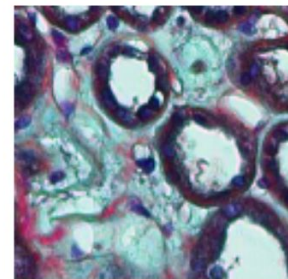
Glomeruli classification results

Method	Avg. acc.	NonGlom acc.	Glomeruli acc.
WND-CHARM GRAY	73.71	81.38	47.39
ET-FL GRAY	87.17	96.87	53.91
ET-FL HS	89.01	98.85	55.29
ET-FL COLOR	91.67	98.61	67.89
ET-FL ACE	91.04	98.71	64.73
ET-FL RH	92.62	98.95	70.93
ET-FL MM	93.78	98.97	75.98

Misclassification



NonGlom



Glom

Summary

- A novel approach combining Icy and Cytomine algorithms (open-source)
- It enables multisite and collaborative digital pathology
- Application for glomeruli detection in kidney human biopsies on a unprecedented dataset
 - > 90 % object recognition rate but...
 - Current results are not yet satisfactory for fully automated routine use
- Ongoing and future work :
 - Improving results
 - Other color normalization techniques
 - Data augmentation and balancing
 - Other classification approaches (convnets...)
 - Consider other evaluation/optimization criteria more related to the final application
 - Comprehensive analysis
 - Recognition of other kidney tissue components and their spatial relations

Acknowledgments

- Unité d'Analyse d'Images Biologiques (Pasteur) :
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Emmanuel Morelon (Hospices Civils de Lyon)

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