

# **SLDC: an open-source workflow for object detection in multi-gigapixel images**

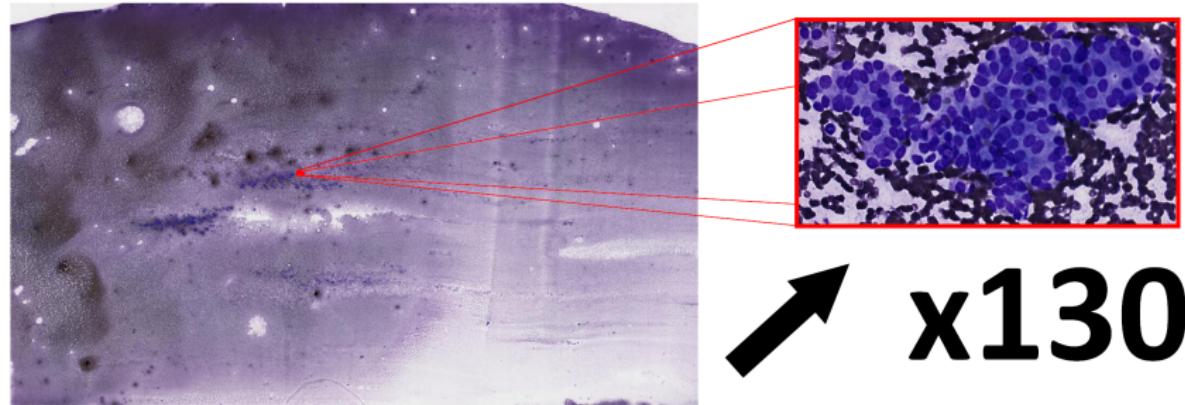
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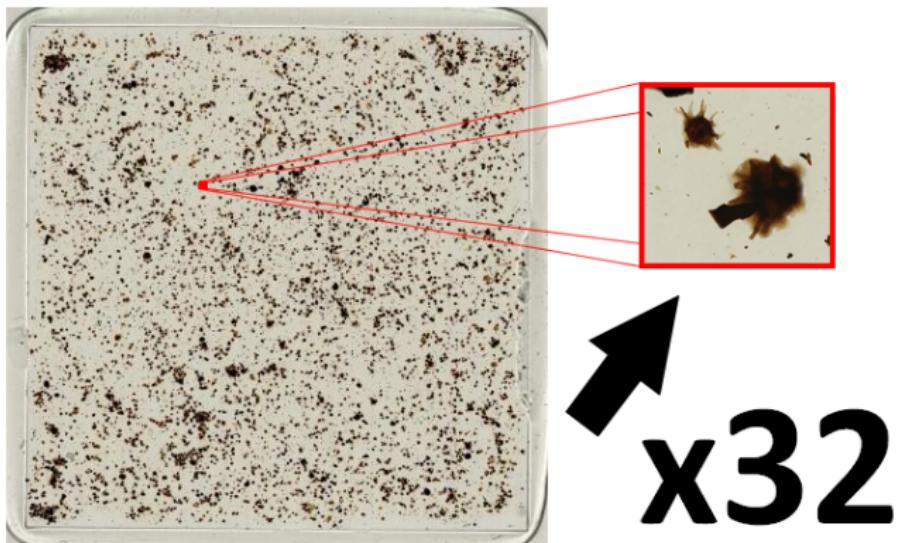
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# Context



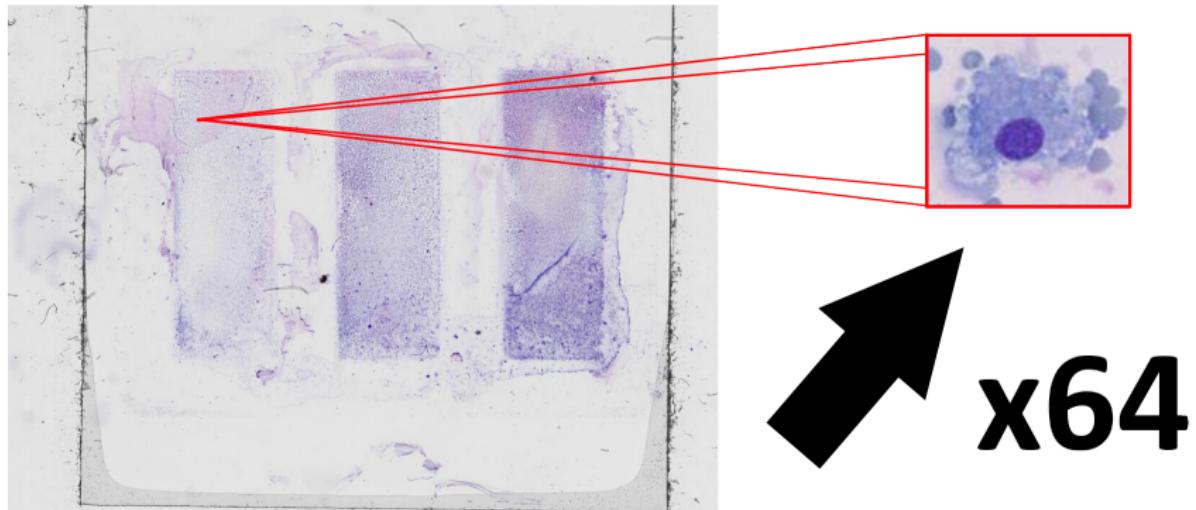
Microscope slide smeared with thyroid cell samples (15 gigapixels).

# Context



Microscope slide smeared with core samples (11 gigapixels).

# Context



Microscope slide smeared with lung cell samples (3 gigapixels).

- Huge slides usually **analysed manually** !
- Machine learning (ML) and image processing (IP) could be used to assist humans
- Problems of **object detection and classification**

*SLDC* is an **open-source Python framework** created for accelerating development of large image analysis workflows.

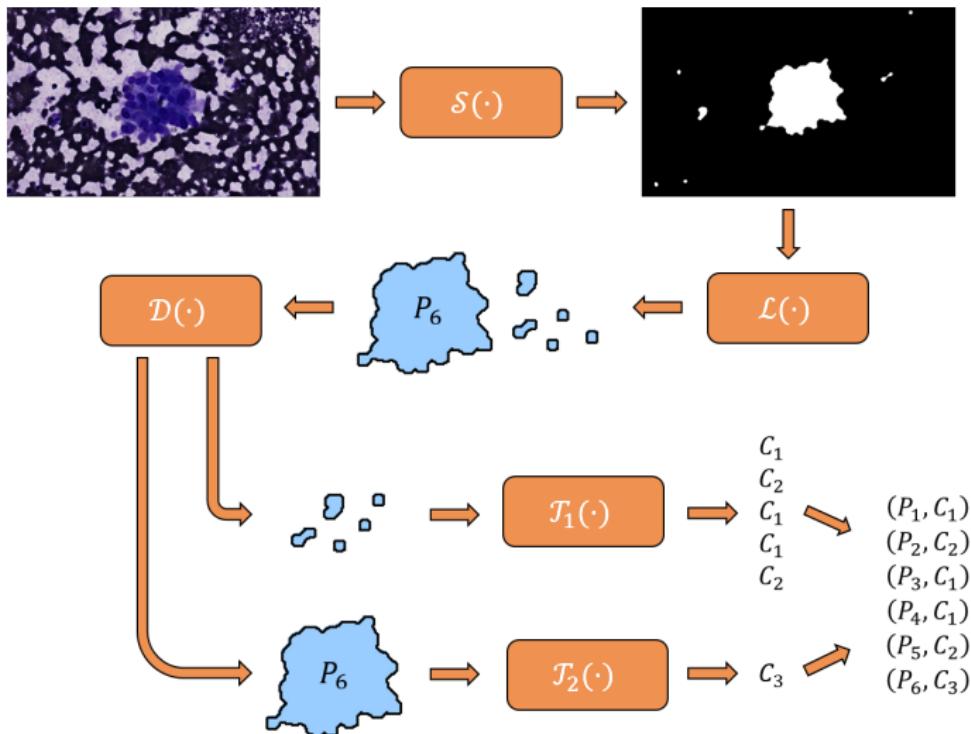
## How ?

- It encapsulates problem-independent logic (parallelism, memory limitation due to large images handling,... )
- It provides a concise way of declaring problem dependant components (segmentation, object classification,... )

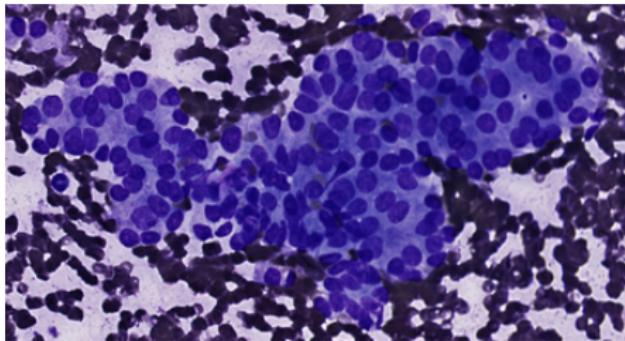
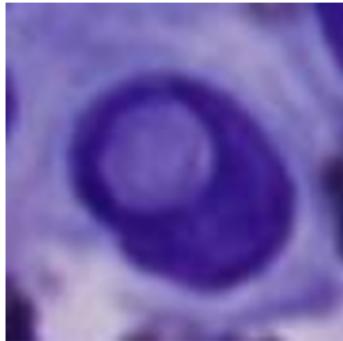
**Where ?** On GitHub at <https://github.com/waliens/sldc>

- **Tile-based processing** to avoid loading a full image into memory
- Several level of **parallelism**: tiles, objects, images,...
- A **customizable logging system** providing a rich feedback about the execution
- **Effortless integration** with other Python libraries: scikit-learn (ML), open-cv (IP), PyCuda (GPU),...
- **Builder components** providing an easy way of constructing complex workflows

# SLDC: how it works



Aim: detect **cells with inclusion** and **proliferative architectural patterns**

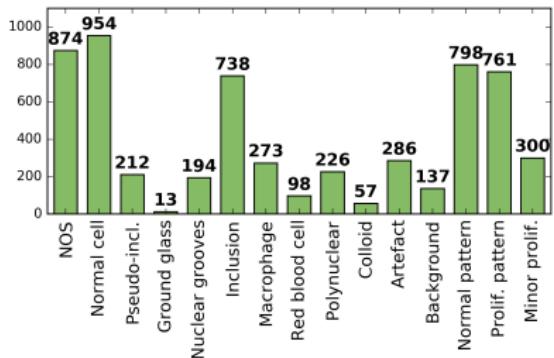


# SLDC at work: Cytomine

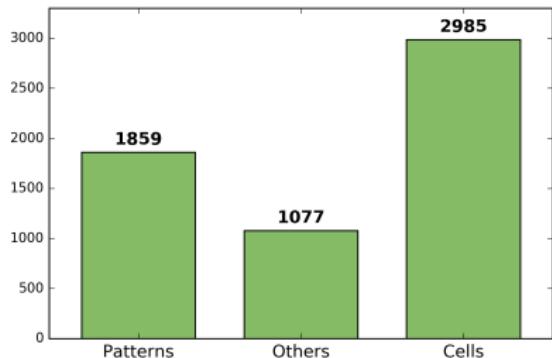
**cytomine** is a web-based environment enabling collaborative multi-gigapixel images analysis. (Marée & al., Bioinformatics; 2016)

The screenshot shows the Cytomine interface for analyzing a histology image. The main view displays a blue-stained tissue section with several green polygonal regions outlined by a user. The top navigation bar includes links for Ontologies, Explore, Storage, Activity, and Search. Below the navigation is a toolbar with various drawing tools like Line, Circle, Polygon, MagicWand, and selection tools. The main image area has two dropdown menus showing file paths: "Annotations" and "Properties". On the right side, there's a sidebar titled "OVERVIEW" containing sections for "INFORMATIONS", "POSITION", "IMAGE LAYERS", and "ANNOTATIONS LAYERS". Under "ANNOTATIONS LAYERS", a list shows "Add" and two users with checked checkboxes: "Mormont Romain (rmormont)" and "Degand Caroline (cddegand) (4)". An "Opacity" slider is also present. At the bottom of the sidebar, it says "This image has no other dimension." and "ANNOTATION'S PROPERTIES".

- **84 images** with size ranging from 4 to 18 gigapixels
- **68 annotated images**
- **5921 labelled annotations** made by cytopathologists<sup>1</sup>



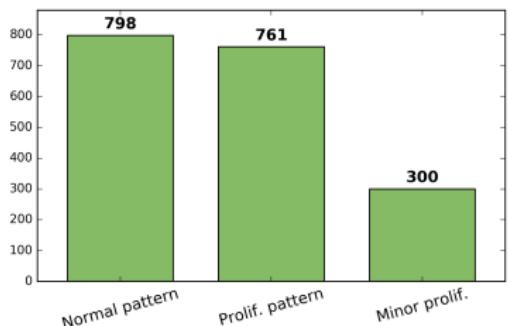
(a) Annot. per term



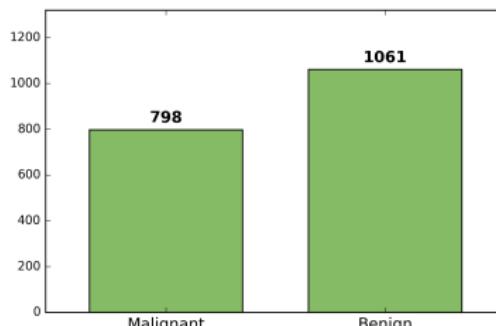
(b) Annot. per group

<sup>1</sup>Team of Pr. Isabelle Salmon, Department of Pathology, Faculty of Medicine, ULB

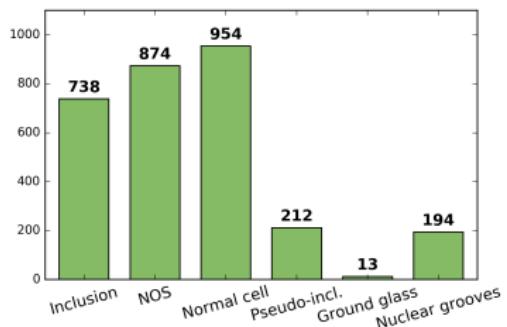
# SLDC at work: data



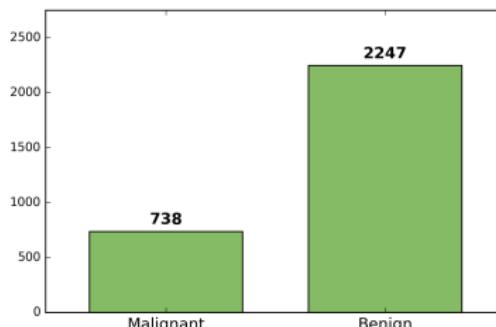
(c) Pattern annot. per term



(d) Pattern annot. per group

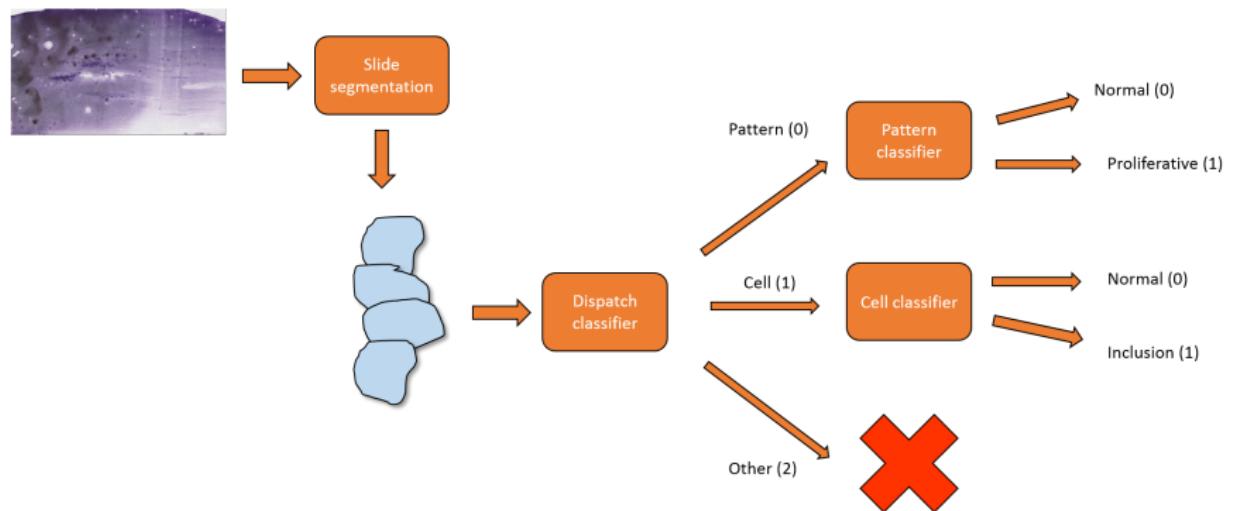


(e) Cell annot. per term

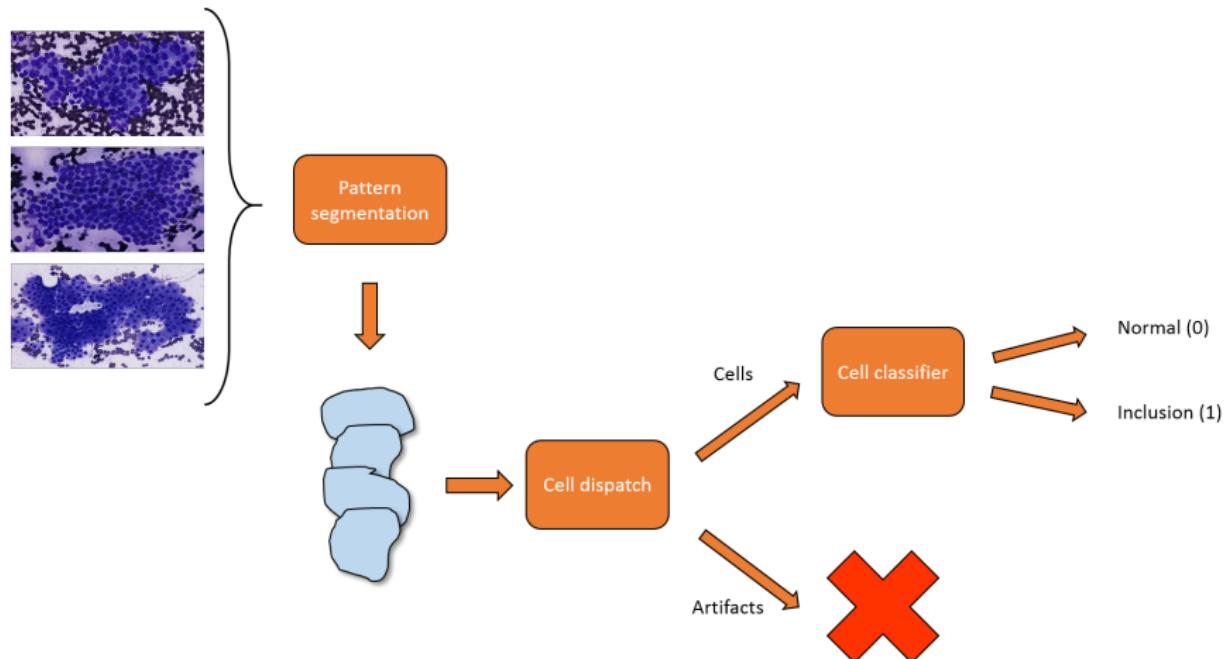


(f) Cell annot. per group

# SLDC at work: workflow



# SLDC at work: workflow (cont'd)



## SLDC at work: workflow (cont'd)

Classification is performed based on the detected object's crop image using **random subwindows** and **extremely randomized trees**<sup>2</sup>.

### Cell with inclusion vs. normal cells:

Accuracy: 0.8523

Precision: 0.6310

Recall: **0.4930**

### Proliferative vs. normal patterns:

Accuracy: 0.8625

Precision: 0.8363

Recall: 0.9493

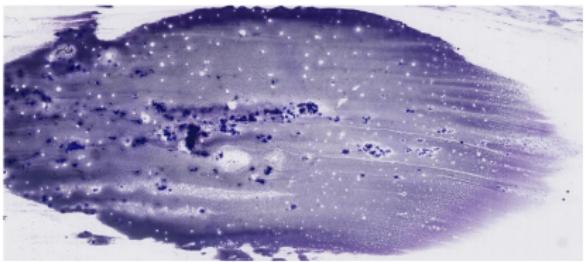
	Normal	Inclusion
Normal	881	62
Inclusion	109	106

	Normal	Prolif.
Normal	158	55
Prolif.	15	281

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<sup>2</sup>Marée et al., Pattern Recognition Letters ; 2016

# SLDC at work: results



Size: 131072 × 57856

Objects found: **20046**

Cells found: 18966

Patterns found: 1080

Time (1st pass): **7 min 30 sec**

Time (2nd pass): 1 h 10 min



Size: 163840 × 95744

Objects found: **79063**

Cells found: 72740

Patterns found: 6323

Time (1st pass): **18 min 20 sec**

Time (2nd pass): 4 h 50 min

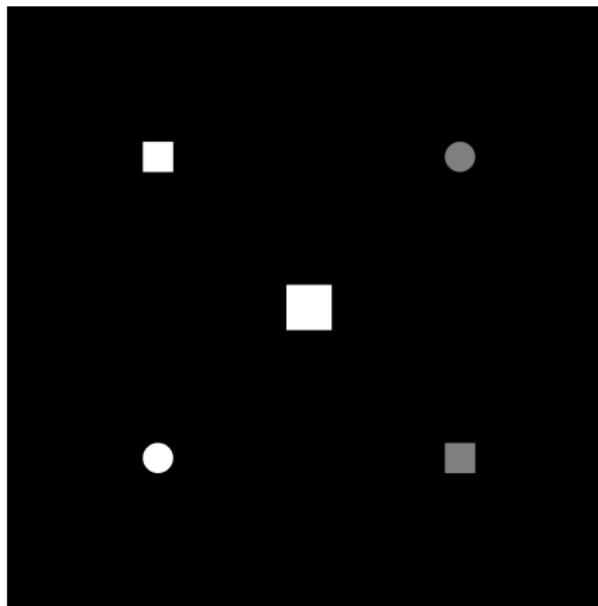
## Future works

???

Thank you for your attention !  
Any question ?

## SLDC: toy example

The aim is to detect circles in the following image. As a bonus, we want to know their center color.



## SLDC: toy example (cont'd)

```
# Defining a segmenter
class CustomSegmenter(Segmenter):
    """All non-black pixels are in an object of interest"""
    def segment(self, image):
        return (image > 0).astype(np.uint8)

# Defining a dispatching rule
class CircleRule(DispatchingRule):
    """A rule which matches circle polygons"""
    def evaluate_batch(self, image, polygons):
        return [circularity(p) > 0.85 for p in polygons]

# Defining a polygon classifier
class ColorClassifier(PolygonClassifier):
    """
    A classifier which returns the color (greyscale)
    of the center pixel of the object
    """
    def predict_batch(self, image, polygons):
        classes = [center_pxl_color(image, p) for p in polygons]
        probas = [1.0] * len(polygons)
        return classes, probas
```

## SLDC: toy example (cont'd)

```
# Build the workflow
builder = WorkflowBuilder()
builder.set_n_jobs(100)
builder.set_segmenter(CustomSegementer())
builder.add_classifier(CircleRule(), ColorClassifier(), disp_label="circle")
workflow = builder.get()

# Process an image
results = workflow.process(image)

# Go through the detected objects
for polygon, dispatch, label, proba in results:
    print "Detected polygon {}".format(polygon)
    print "Dispatched by {}".format(dispatch)
    print "Predicted class {}".format(label)
    print "Probability {}".format(proba)
    print ""
```

## SLDC: toy example (cont'd)

Detected polygon POLYGON ((...))

Dispatched by 'circle'

Predicted class 128

Probability 1.0

Detected polygon POLYGON ((...))

Dispatched by 'circle'

Predicted class 255

Probability 1.0