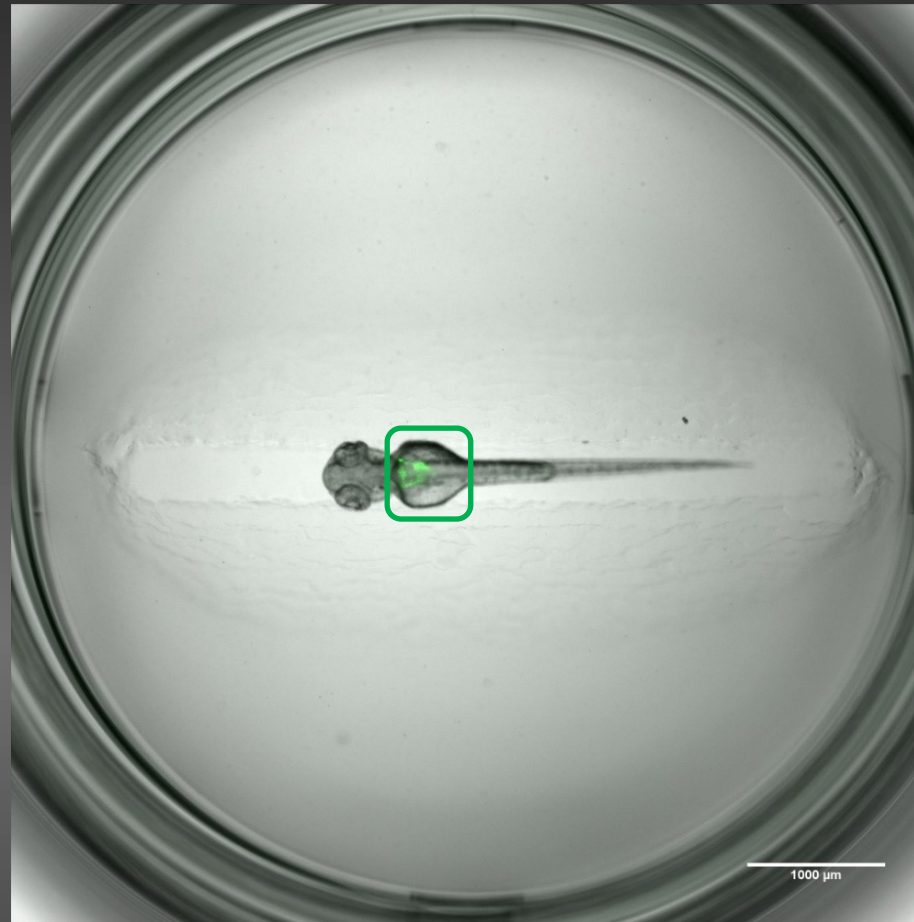


Multi-Template Matching: a versatile tool for object-localization in microscopy images

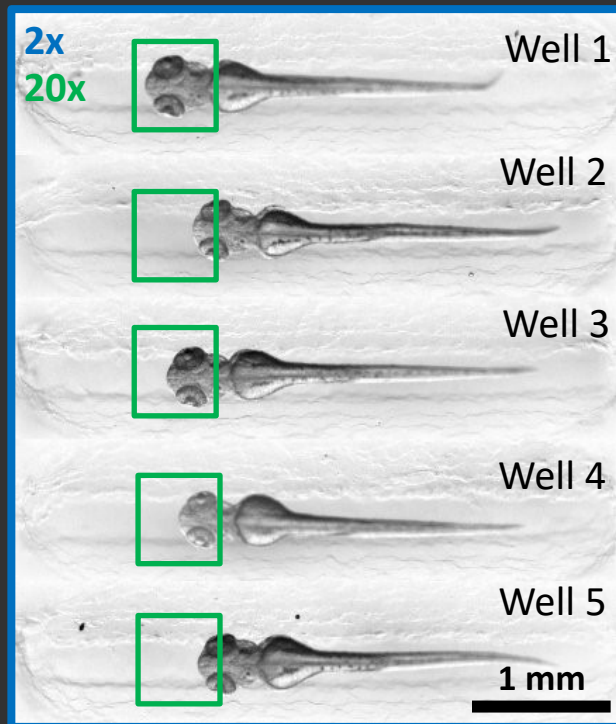


transgenic zebrafish (wt1b:egfp) in a well
2X - Overlay BF + GFP

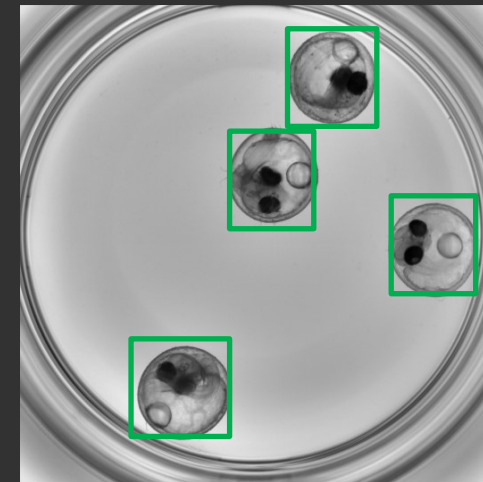
Laurent THOMAS
PhD student under supervision of Jochen Gehrig



Motivation : High resolution imaging of sample with variable positions



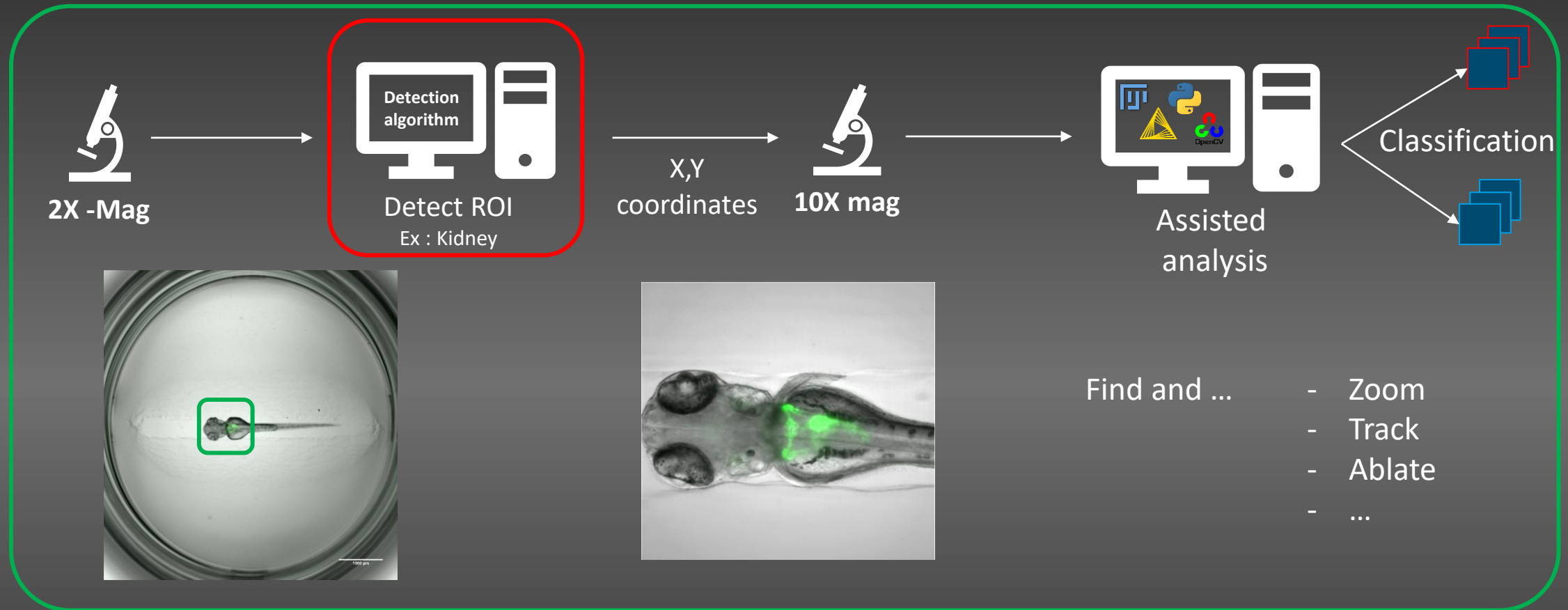
Slight position shift



Random positions
and
orientations

Using feedback microscopy for guided acquisition

Semi-automated imaging pipeline

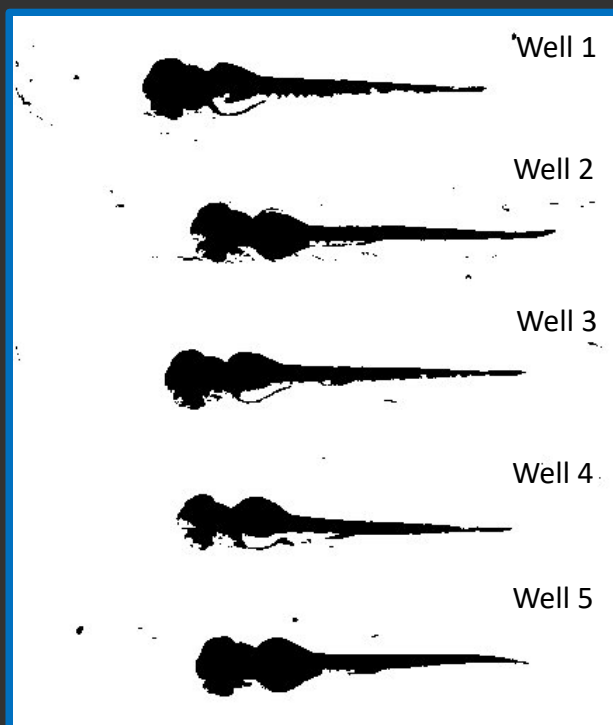


✓ Time-saving

✓ Reproducible

✓ Easy to use

ROI detection algorithm : Expectations



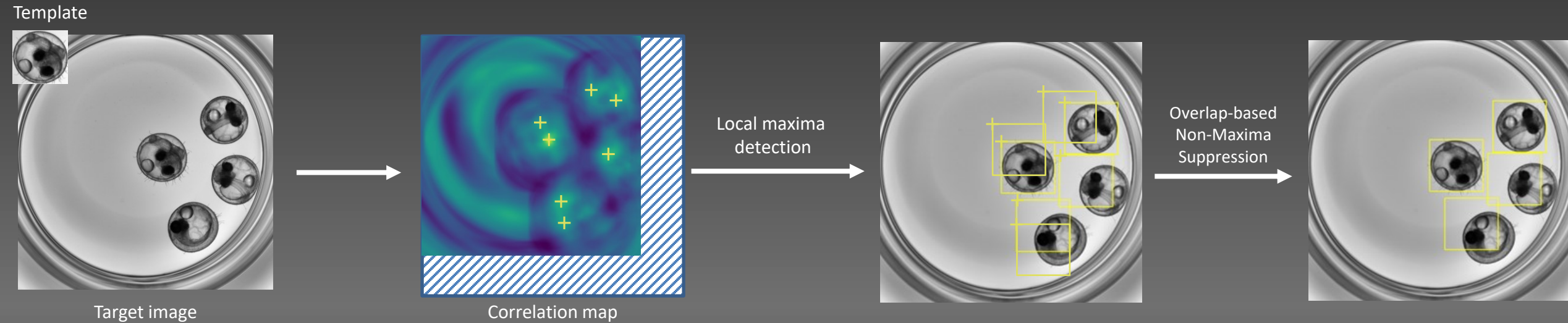
Requirements

- Generic (little tweaking)
- User-friendly (little parameters, simple to understand)

Challenges

- Diffuse fluorescent signal over specimen (not site-specific)
- Variable morphology and/or fluorescence intensity

Template Matching



- Simple (pixel-intensity comparison)

BUT

- Limited flexibility (rotation...)

Multi-template matching: a versatile tool for object-localization in microscopy images

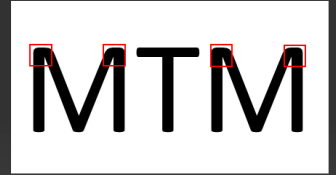
Laurent S. V. Thomas^{1,2*} and Jochen Gehrig^{1*}



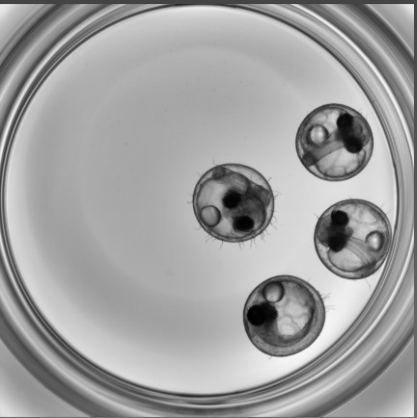
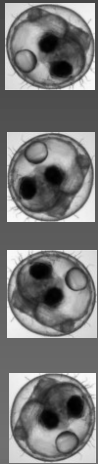
Check for updates

Multi-Template Matching

ACQUIFER

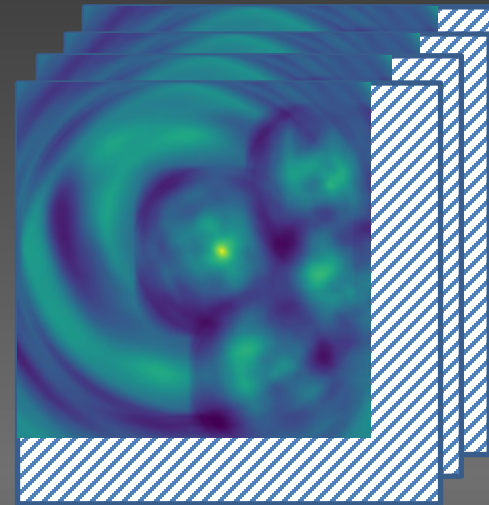


N Templates

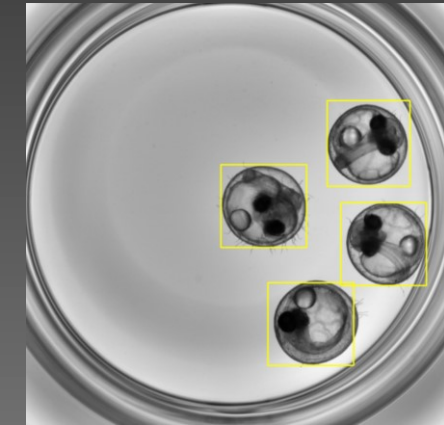


Target image

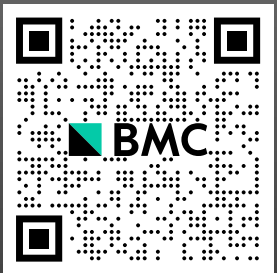
N correlation maps



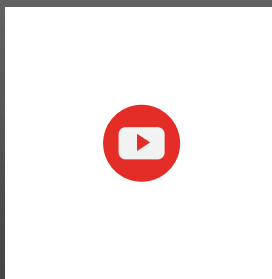
Local maxima
+
Overlap-based
Non-Maxima Suppression



Improved detection rate by supporting multiple detection patterns (templates)



Article in BMC
Bioinformatics



YouTube
tutorials



Source codes
and documentations

Available for :

- ✓ Fiji (dedicated update site)
- ✓ Python (via pip)
- ✓ KNIME (using python)

Multi-Template Matching results

Zebrafish head (single template)



N = 96

Image courtesy Gunjan Pandey

Spheroid (single template flipped and rotated)

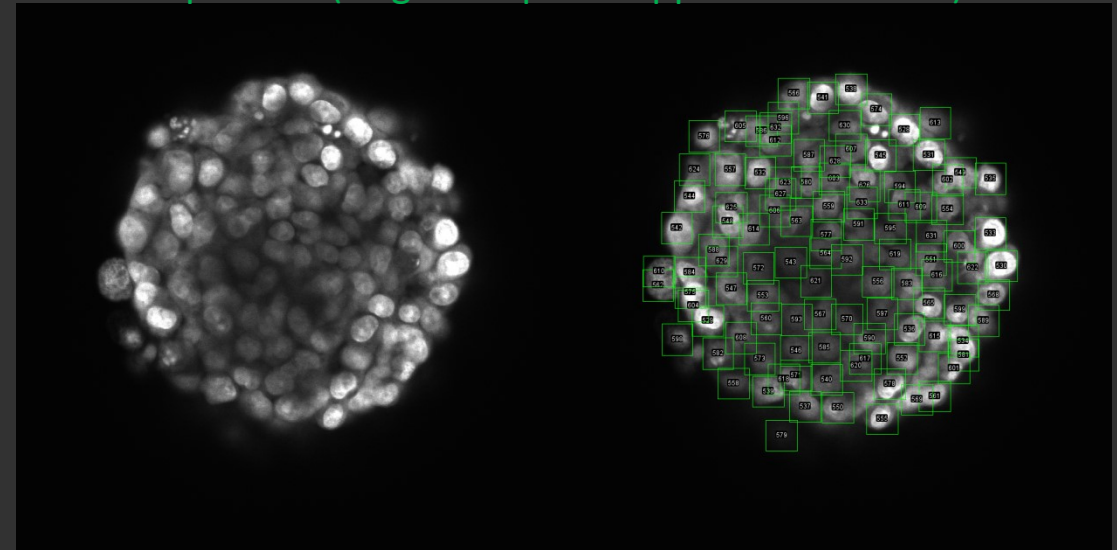
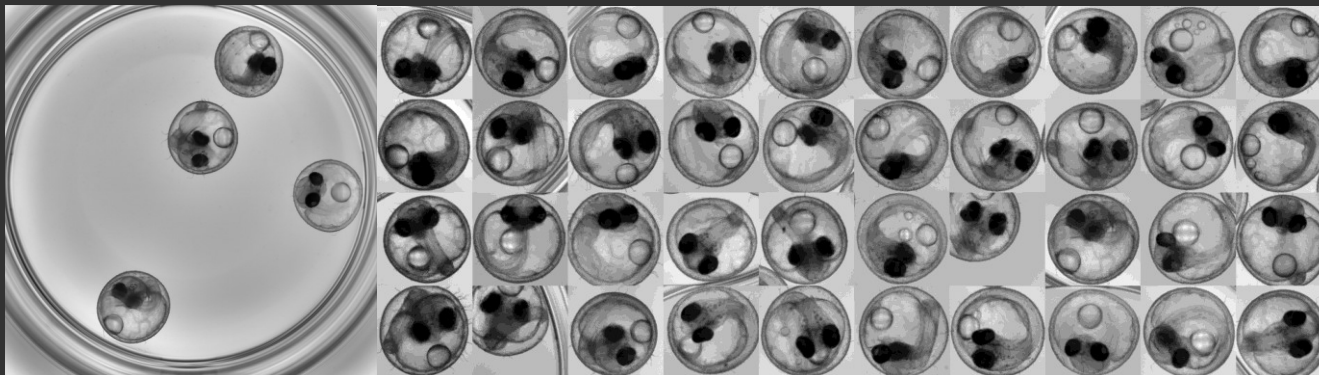


Image courtesy – Riviere C., Tricot V., Rousseau D., Rasti P., – Angers, France

Medaka larvae (single template flipped and rotated)



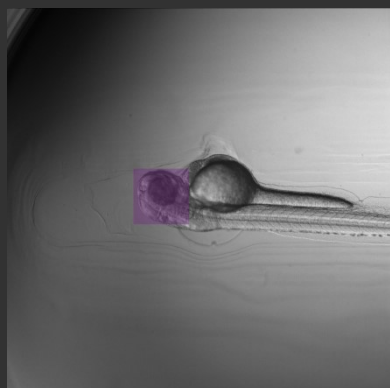
N = 10 images x 4 eggs

Image courtesy Jakob Gierten

- ✓ Simple (normalised pixel-pixel comparison)
- ✓ Robust to change of illumination, to mild morphology changes
- ✓ Easy to install and (re)-use
- ✗ Limited tolerance to variability (more templates = longer computation)

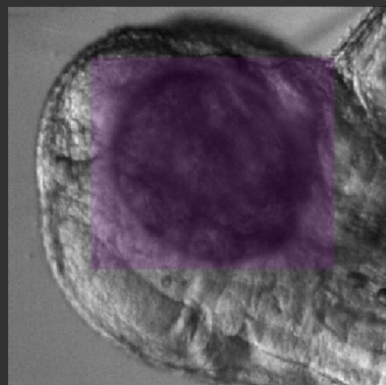
2-step template matching for more robust detection

Example : Eye and lens segmentation in Zebrafish larvae



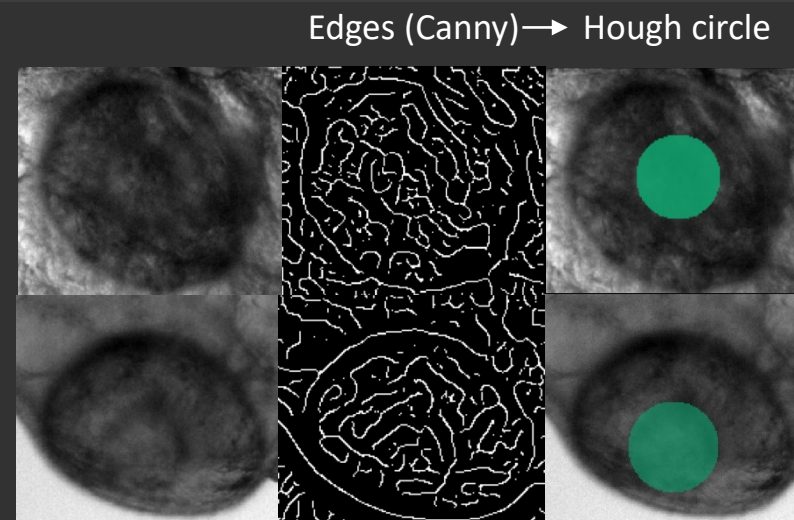
Head detection
Template matching

crop →



Eye detection
Template matching

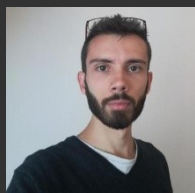
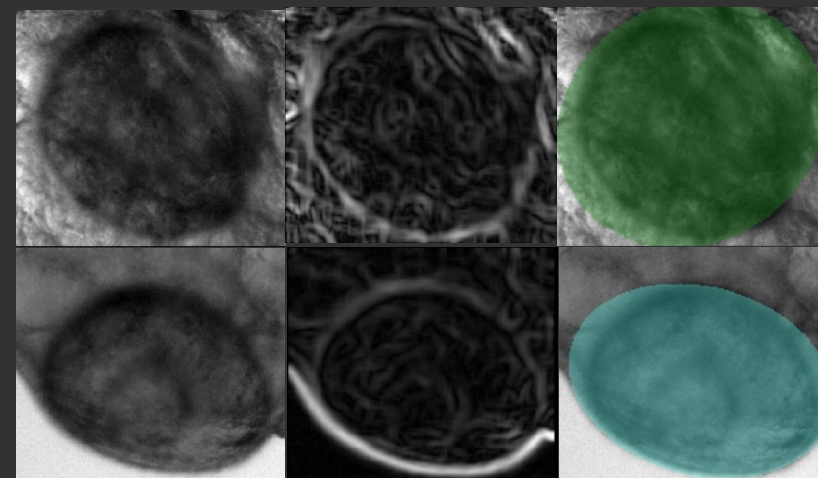
crop →



Lens

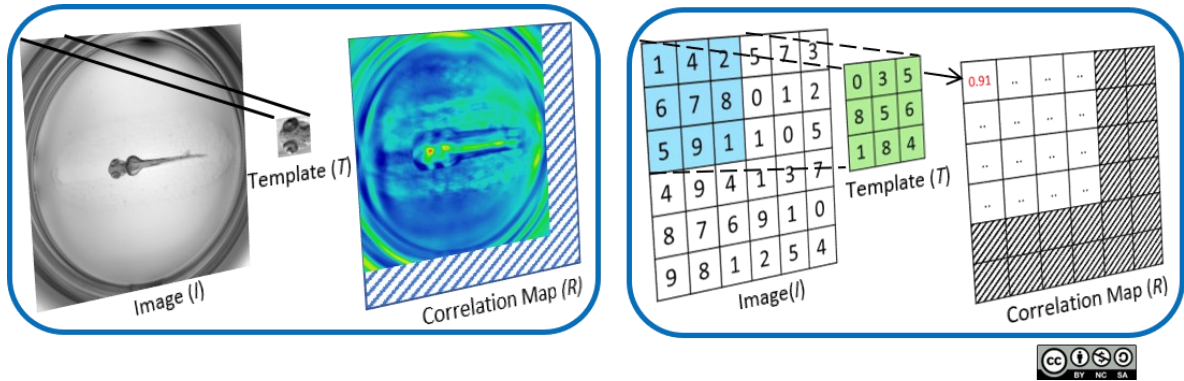
Eye

Edges (Sobel) → Ellipse fitting



Alexandre
Jeanne

A bit of theory: the correlation map



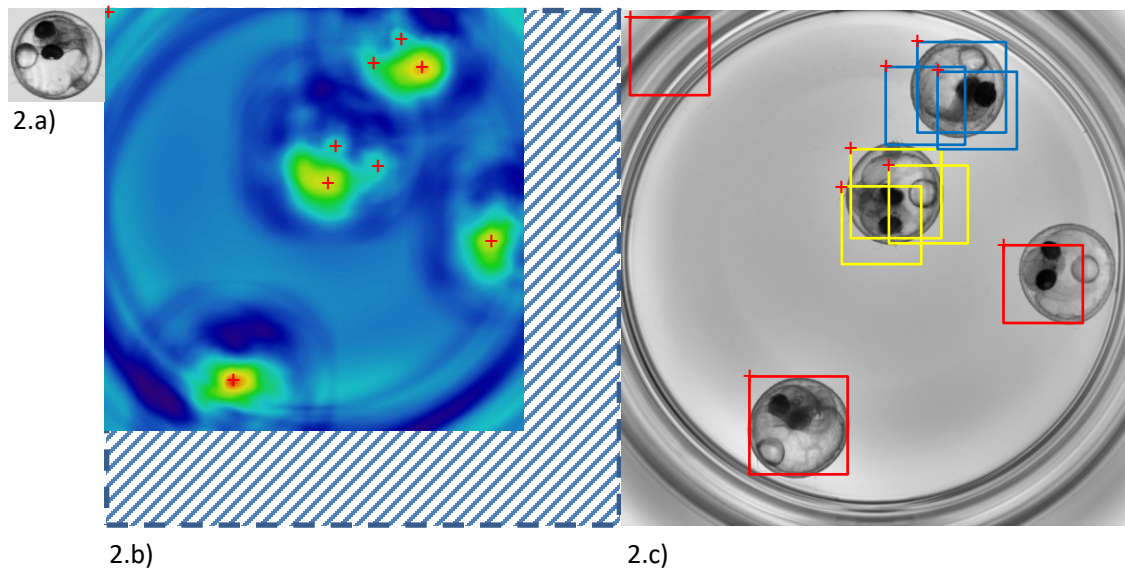
$$CorrMap(x, y) = \frac{Template * ImagePatch\ at\ (x, y)}{\sqrt{\sum_{x', y'} Template^2 * \sum_{x', y'} ImagePatch\ at\ (x, y)^2}}$$

$$CorrMap(x, y) = \frac{\sum_{x', y'} T(x', y') * I(x + x', y + y')}{\sqrt{\sum_{x', y'} T(x', y')^2 * \sum_{x', y'} I(x + x', y + y')^2}}$$

- 1 correlation map/template

$$R(0,0) = \frac{1 * 0 + 4 * 3 + 2 * 5 + 6 * 8 + 7 * 5 + 8 * 6 + 5 * 1 + 9 * 8 + 1 * 4}{\sqrt{(1^2 + 4^2 + 2^2 + 6^2 + 7^2 + 8^2 + 5^2 + 9^2 + 1^2) * (0^2 + 3^2 + 5^2 + 8^2 + 5^2 + 6^2 + 1^2 + 8^2 + 4^2)}} = 0.91$$

A bit of theory: maxima detection

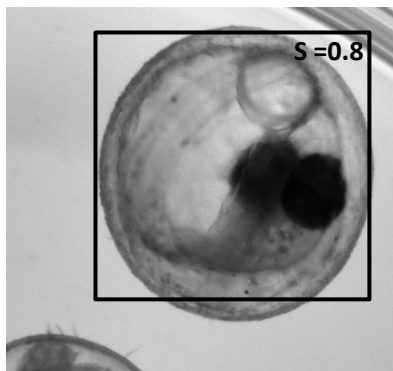
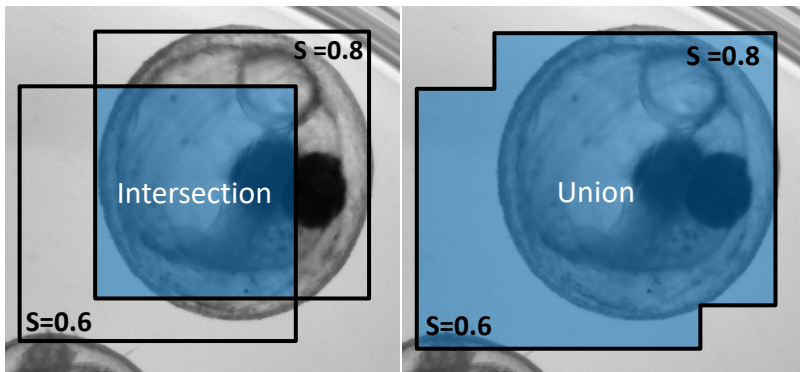


Probable object-location
-> maxima of the correlation map

- Problem -> Multiple hits per object
- Apply a score-threshold, still redundant detections

Non-Maxima Suppression (NMS)

Removing overlapping detection with NMS



$$\text{Normalized overlap} = \frac{\text{Intersection}}{\text{Union}}$$

Overlap > threshold

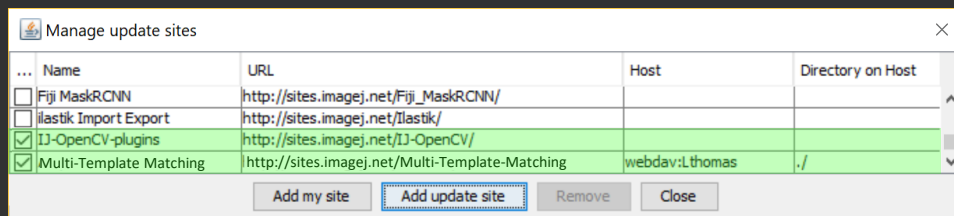
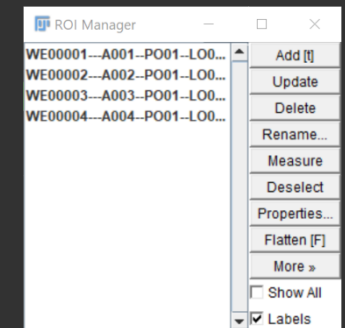
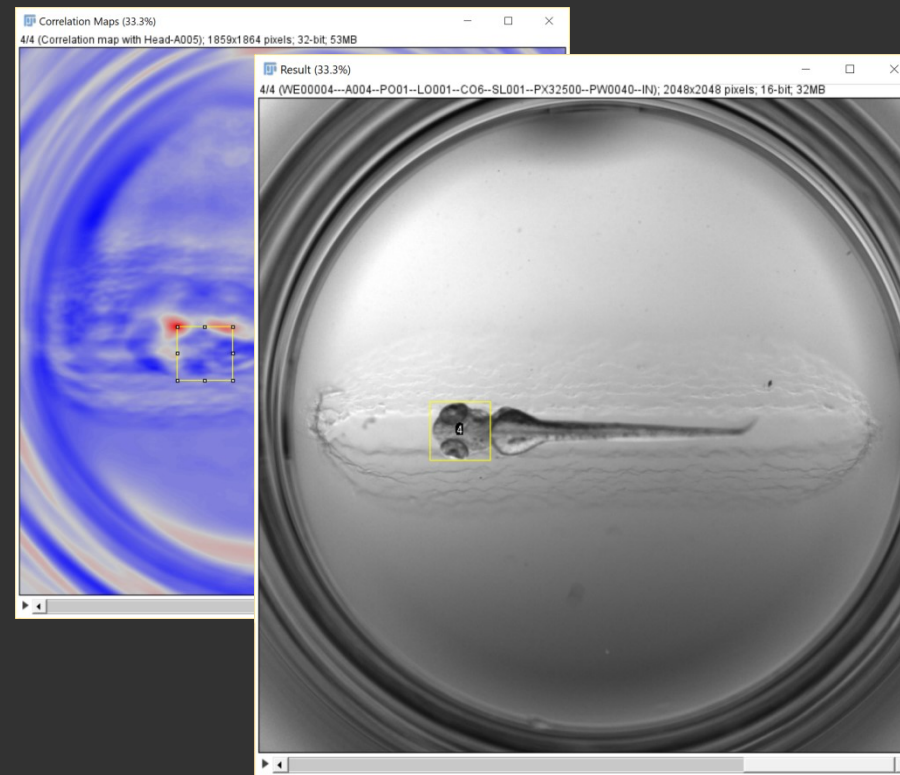
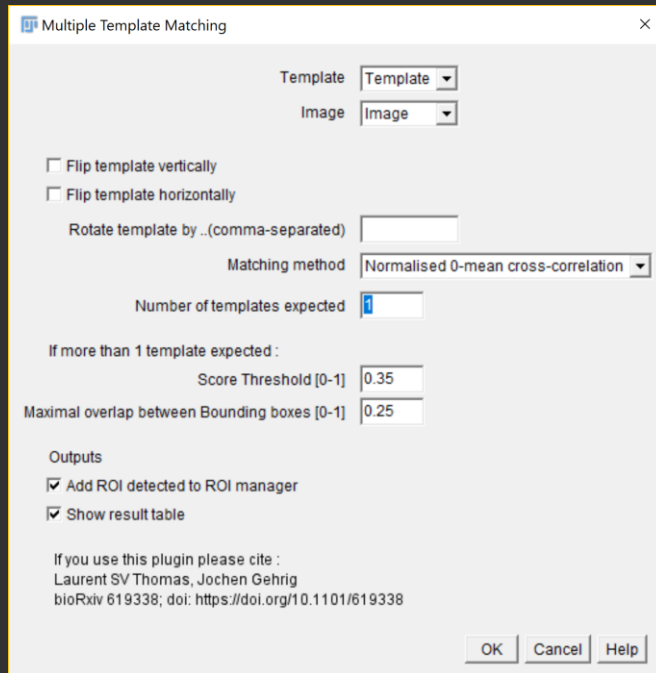
- Redundant detections
- Discard the lower-score detection

Overlap < threshold

- Distinct detections
- Keep both detections

Multi-Template matching

As a Fiji plugin



- Macro recordable
- Available via an update site

Faster detection

1) Down resolution

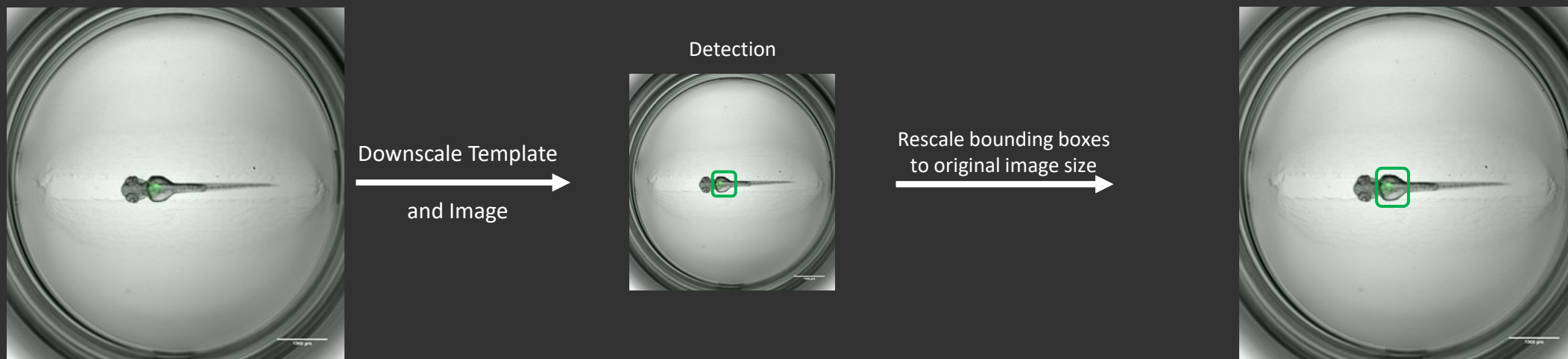
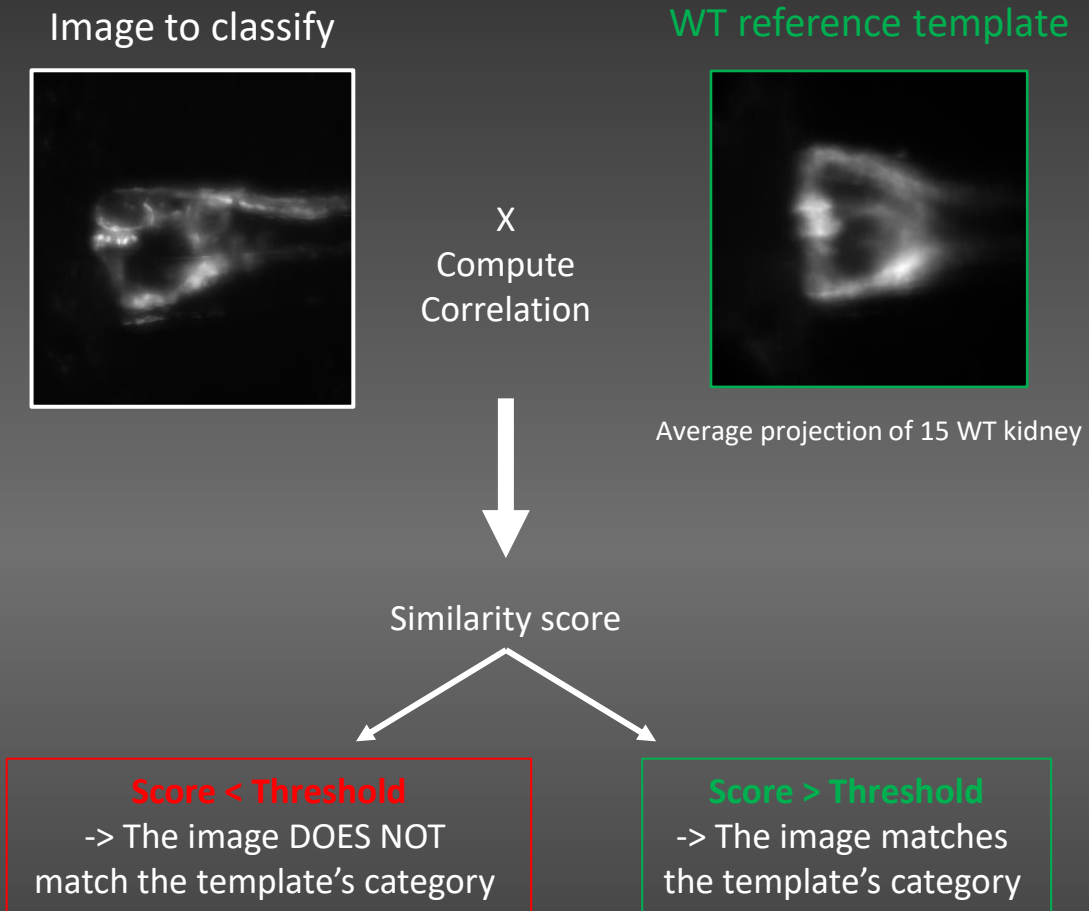
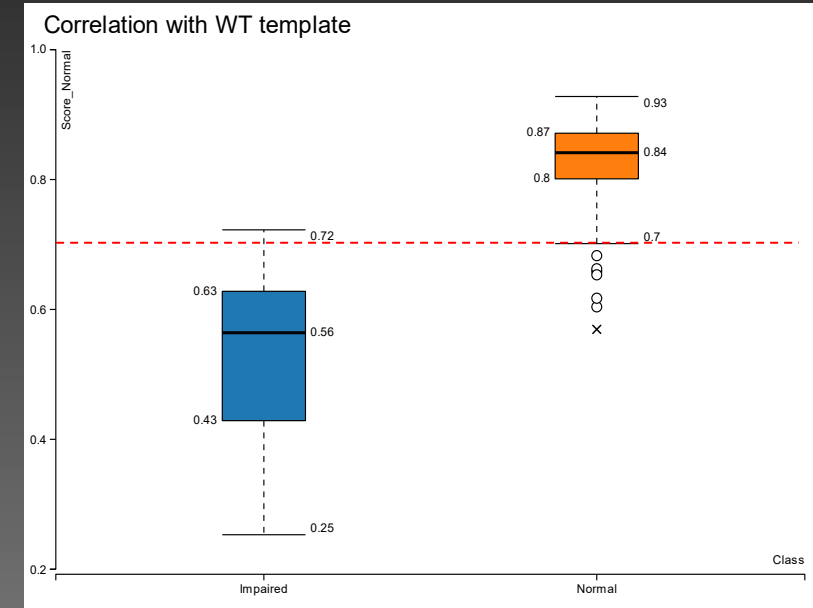


Image-classification with template matching



Score distribution for a fraction of the annotated dataset ("Training set": 310 images)



Using 0.7 as a threshold between the normal and impaired category ("Test set": 930 other images)

Class \ Predicted_Class	Normal	Impaired
Normal	603	23
Impaired	16	288

Correct classified: 891 Wrong classified: 39

Accuracy: 95.806 % Error: 4.194 %

Cohen's kappa (κ) 0.905