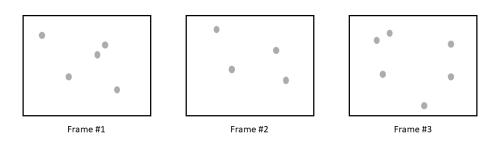
Tracking

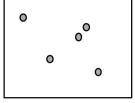
Thierry Pécot Research Engineer Biosit SFR UMS CNRS 3480 - Inserm 018 **CZI Imaging Scientist**



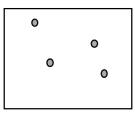




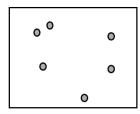
Segmentation





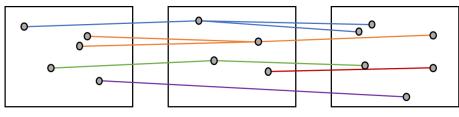


Frame #2



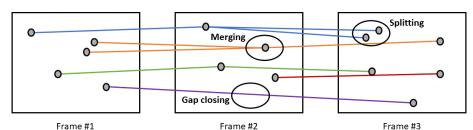
Frame #3

Data association, particle linking,...



Frame #1 Frame #2 Frame #3

Special events





Methods Volume 115, 15 February 2017, Pages 80-90



TrackMate: An open and extensible platform for single-particle tracking

Jean-Yves Tinevez ^a $\stackrel{\circ}{\sim}$ $\stackrel{\circ}{\sim}$, Nick Perry ^{a, 1}, Johannes Schindelin ^{b, 2}, Genevieve M. Hoopes ^c, Gregory D. Reynolds ^c, Emmanuel Laplantine ^d, Sebastian Y. Bednarek ^c, Spencer L. Shorte ^a, Kevin W. Eliceiri ^{b, a}

Bringing TrackMate into the era of machine-learning and deep-learning

- O Dmitry Ershov, Minh-Son Phan, O Joanna W. Pylvänäinen, O Széphane U. Rigaud, Laure Le Blanc, A Arbhur Charles-Orszag, O James R. W. Comway, O Romain F. Laine, O Nathan H. Roy, O Daria Bonazzi, Guillaume Duménii, O Guillaume Jacquemet, O Jean-Yves Tinevez doi: https://doi.org/10.1101/20.11.09.03.458852
- This article is a preprint and has not been certified by peer review [what does this mean?].

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Published: 20 July 2008

Robust single-particle tracking in live-cell time-lapse sequences

Khuloud Jagaman [™], Dinah Loerke, Marcel Mettlen, Hirotaka Kuwata, Sergio Grinstein, Sandra L Schmid & Gaudenz Danuser

1 - SEGMENTATION



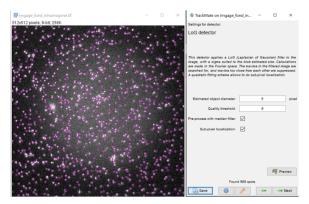
Classic detectors:

- Manual annotation
- Threshold detector
- DoG/LoG detector

1 - SEGMENTATION



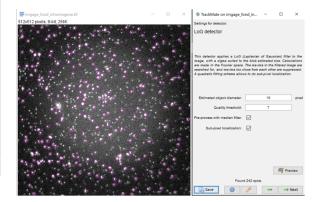
LoG detector: scale-based detector, powerful with **objects of the same size** such as intracellular particles



1 - SEGMENTATION

TrackMate on C1-CelegansEar... — Select a detector Label image detector This detector creates spots by importing regions from a label image. A label image is an image where the pixel values are integers. Each object in a label image is represented by a single common pixel value (the label) that is unique to the object. This detector reads such an image and create spots from each object. In 2D the contour of a label is imported. In 3D, spherical spots of the same volume that the label are created.

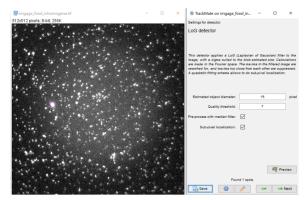
LoG detector: scale-based detector, powerful with **objects of the same size** such as intracellular particles



1 - SEGMENTATION



LoG detector: scale-based detector, powerful with **objects of the same size** such as intracellular particles



1 - SEGMENTATION



Classic detectors:

- Manual annotation
- Threshold detector
- DoG/LoG detector

New generation detectors:

- Stardist
- Cellpose
- Weka
- Ilastik
- MorpholibJ
- · Custom segmentation via label images

2 - SPOT FILTERING

Filter out spots based on spot features such as size, average intensity, ...

Set filters on spots

Possibility to visualize spot features with heat maps.

3 - DATA ASSOCIATION



Trackers:

- · Manual tracking
- Overlap tracker: based on intersection over union between two consective frames – only 2D
- Nearest-neighbor tracker: associate closest particles between two consecutive frames
- Kalman tracker: based on Kalman filtering, suited for objects with directed and constant motion, allows gap closing
- Lap tracker: implementation of utrack (Jaqaman et al.), based on the Minear Assignment Problem mathematical framework, allows to change linking cost between particles by penalizing features and to perform splitting, merging and gap closing
- Simple LAP tracker: same as LAP tracker, without cost penalties, splitting and merging

4 - TRACK FILTERING

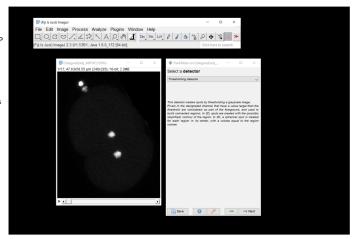
Filter out tracks based on track features such as number of spots, duration, ...

Set filters on tracks

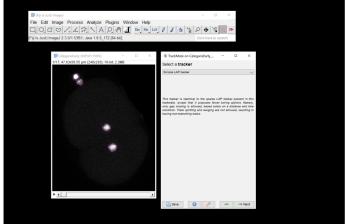
Possibility to visualize track features with heat maps.

Open CelegansEarly_MIP with TrackMate and segment cells with Thresholding detector

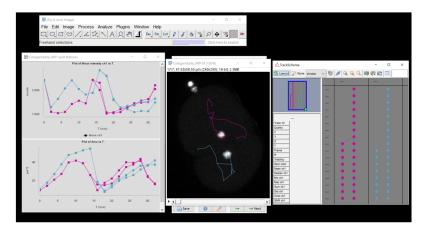
Filter out as many as possible non-cell objects but keep all cells detected



Use Simple LAP tracker, filter out track(s) corresponding to non-cell objects and manually split the tracks when cells divide with TrackScheme



Plot features, save results...



Use LAP tracker with splitting and filter out track(s) to directly obtain the 2 tracks for cells

