

Authors

Wael GHARBI
Guillaume FIDANZA

Supervision

TELECOM SudParis
Nicolas ROUGON

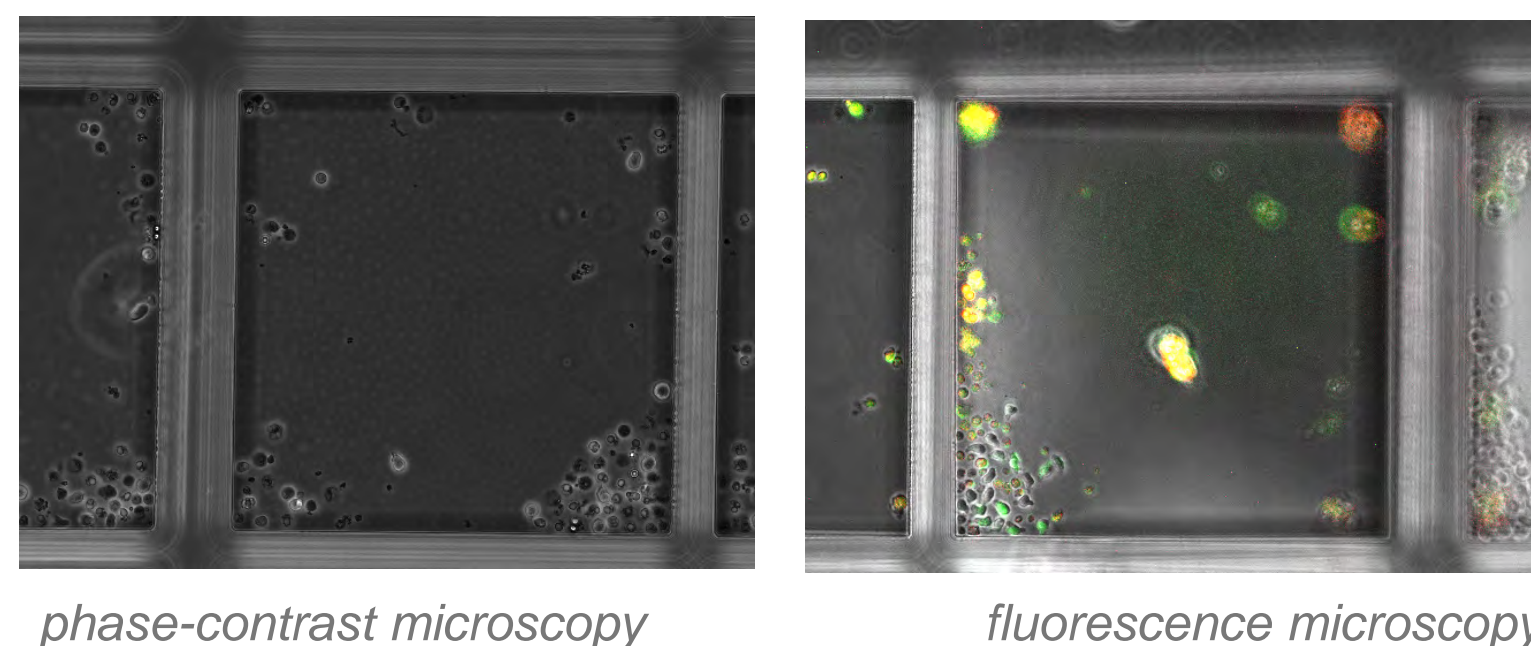
GENETHON
Daniel STOCKHOLM



MOTIVATION & OBJECTIVES

Single-cell microscopic imaging

- Single-cell imaging in **phase-contrast** and **fluorescence microscopy** allows for monitoring *in vivo* human T cells confined in μ -trap arrays
- Analyzing **time-lapse video sequences** could bring new insights on the way T cells evolve into **T-reg** cells which govern immune tolerance
- This requires image processing methods for automated cell counting, segmentation and tracking, coupled to data processing techniques for understanding cell interactions



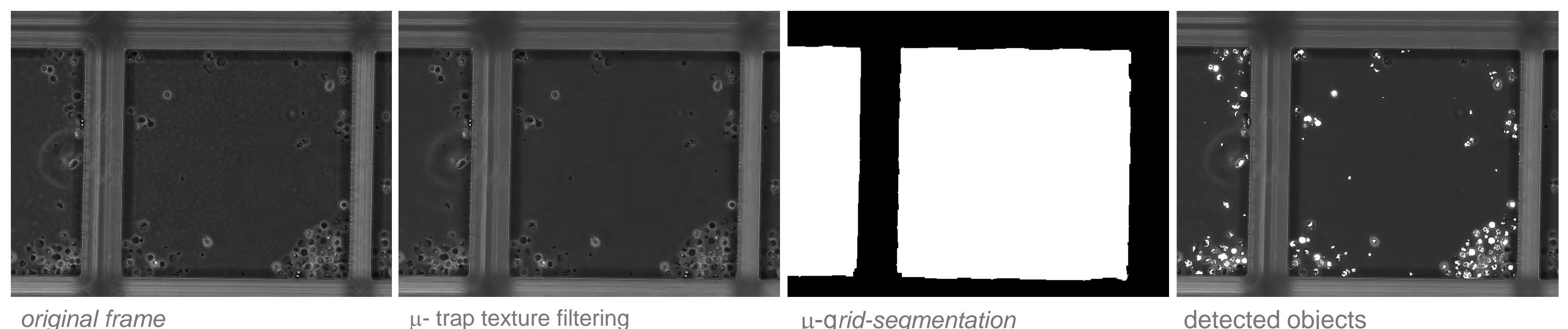
Challenges

- Detect all the objects in the field of view
- Distinguish between impurities, dead and alive cells
- Track individual cell to reconstruct trajectories

MATERIAL & METHODS

- **Setup:** Human T cell lines in μ -trap arrays with macrophages and beads connected to specific molecules
- **Database:** 12 time-lapse video sequences = phase-contrast images of μ -grid sub-regions at regular time intervals during 24 hours + fluorescence images acquired 2-3 minutes after phase-contrast data
- **μ -grid segmentation** using **temporal mathematical morphology** on frames groups
- **Object segmentation** in each frame:
 - > **Preprocessing:** μ -trap background texture homogenization via **PDE filtering** (anisotropic diffusion with Tukey biweight-based conduction function)
 - > **Detection** using **geodesic morphology** using μ -trap background and object dark sub-region markers

RESULTS



- Sound performances showing **high sensitivity** / **low specificity**: all objects are detected, including beads and impurities
- Spurious over-segmentation artefacts can occur due to possibly non simply connected object markers

CONCLUSION & PERSPECTIVES

- Using **temporal features** would improve robustness and allow for distinguishing between dead/alive cells
- Specificity could be enhanced by relaxing the requirement to detect impurities
- Future work:
 - > implement a **cell tracker** based on frame-by-frame object segmentations
 - > use **fluorescence data** to discriminate T-reg cells

