

## Looking Ahead

- 1) To solve technical issues on the reference library data. There is a drift along XY on some of the colloidal samples. Also, some images show uneven illumination over the field of view.
- 2) Begin tracking *E. coli* as a reference in the motility buffer.
- 3) Analyze the tracking performance and retrieve physical observables and statistical quantities related to bacterial motility (e.g., instantaneous speed distribution, turning angle distribution, etc).
- 4) If the data verifies a fully functional tracking, we can start experimenting on the wild-isolates.

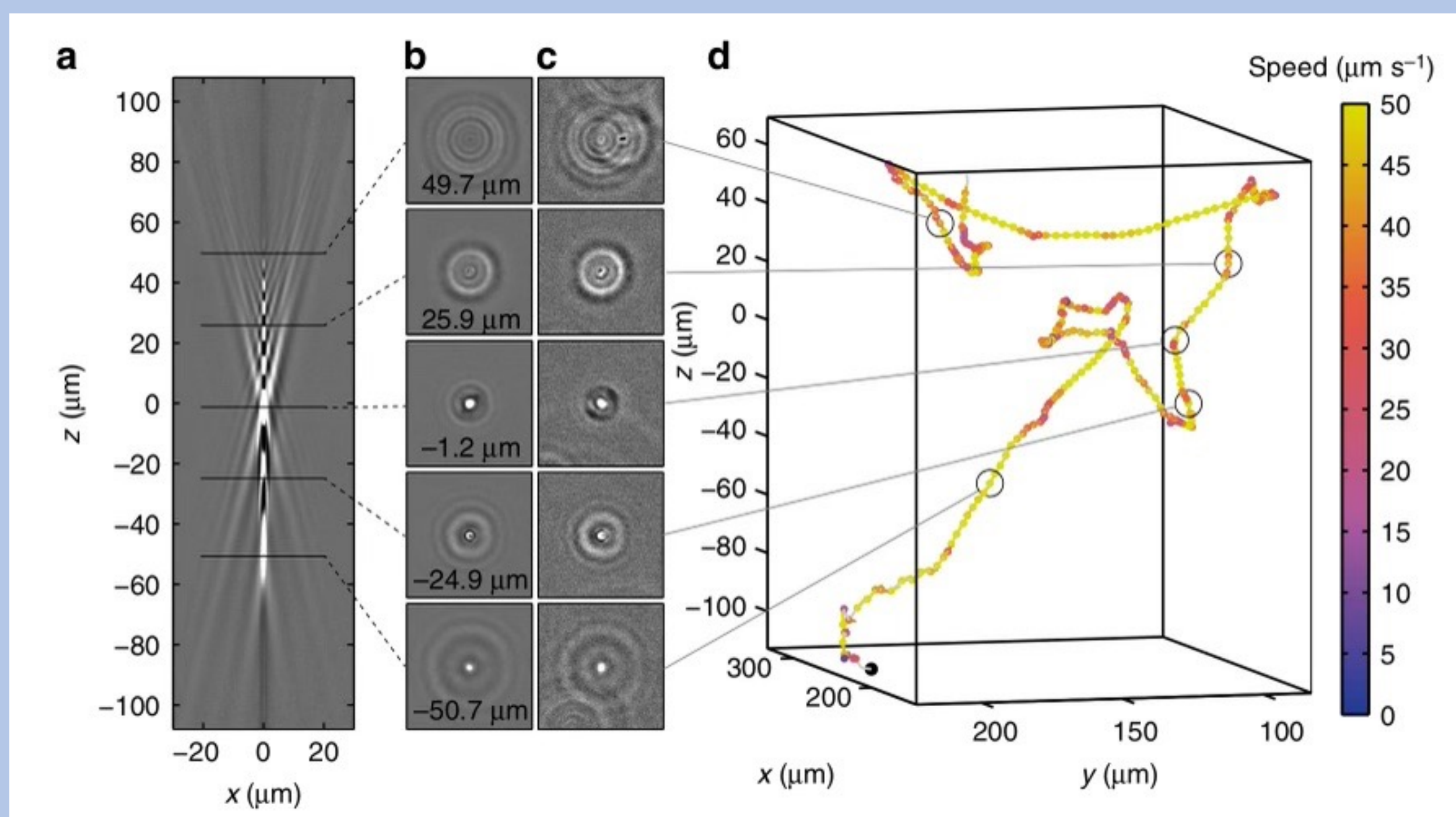


Figure #1: 3D bacterial tracking by comparing out-of-focus diffraction patterns to a reference library (Reference #1)

## Introduction

Bacteria display unique motility patterns when traveling in complex three-dimensional (3D) environments. Unfortunately, two-dimensional microscopy fails to detail critical features of 3D motile behavior. Today, there exists an easy and effective high-throughput 3D bacterial tracking method applicable in standard phase-contrast microscopy. We can localize bacteria (i.e., *E. coli*) at a micron-scale resolution by maximizing image cross-correlations amongst their phase rings and a reference library. Ultimately, we can create 3D trajectories of different microorganisms over a specific time span.

## Methods

- Bacteria culturing
- Bacterial sample chamber preparation
- Microscopy and data acquisition
- Image cross-correlation
- Reference stack generation
- Tracking algorithm



Image #1: Single colonies of *E. coli* in LB agar (left). Cultured falcon tubes with single colonies and a negative control (right).

## Progress

- This project is still in its fledgling stages of development, so research with this innovative technique has yet to begin.
- Building the reference library → silica particles of 1 micron in diameter were stuck inside 5% Polyacrylamide gel.
- The colloids at different heights help construct an aggregated set of images that constitute a reference library.
- Once complete, image cross-correlation between reference library frames with known z-distances and out-of-focus diffraction patterns will establish a trajectory from which we can analyze specific motility patterns.

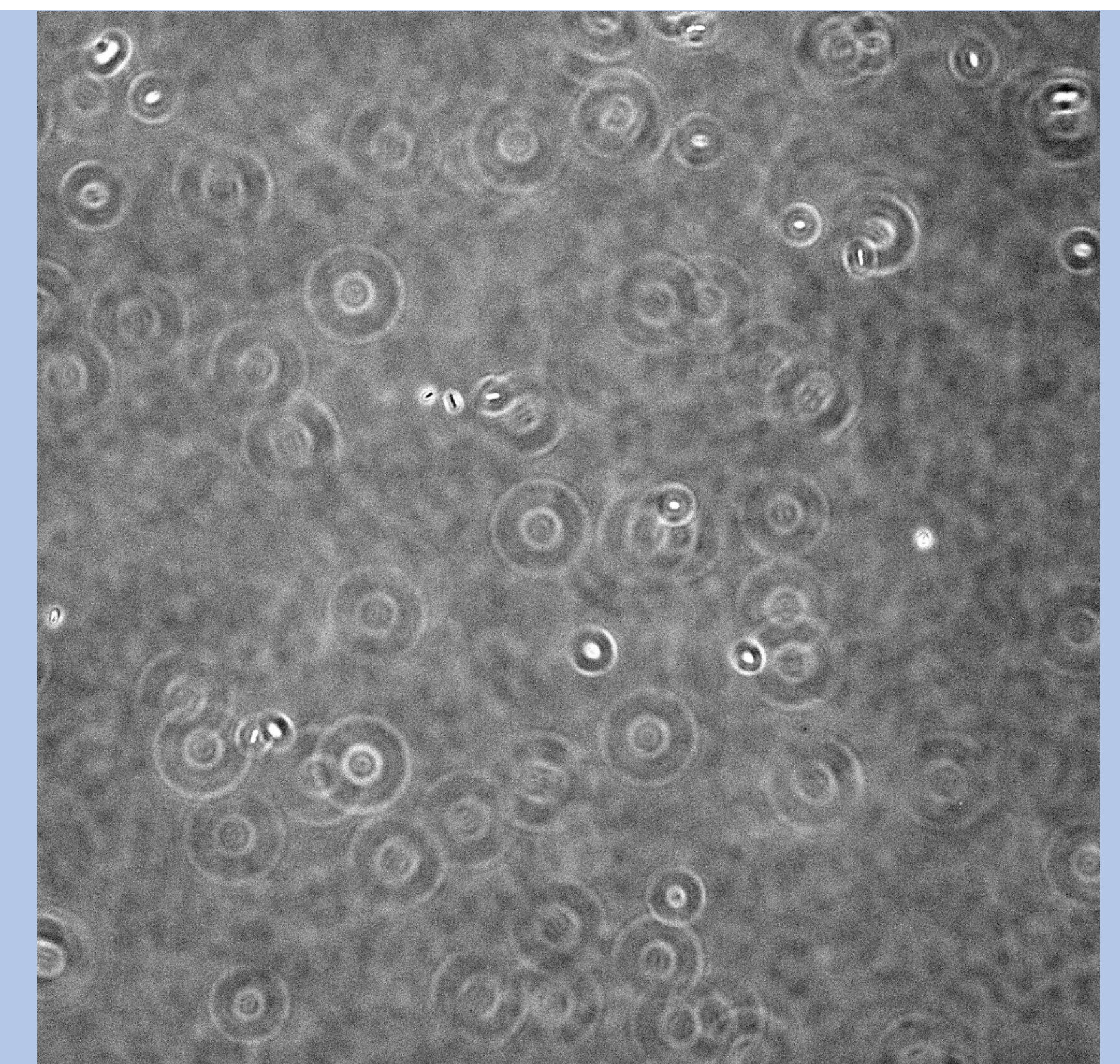


Image #2: Phase-contrasted images of swimming *E. coli* with noticeable phase rings

## Citations

Reference #1: Taute, K., Gude, S., Tans, S. *et al.* High-throughput 3D tracking of bacteria on a standard phase contrast microscope. *Nat Commun* **6**, 8776 (2015). <https://doi.org/10.1038/ncomms9776>

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