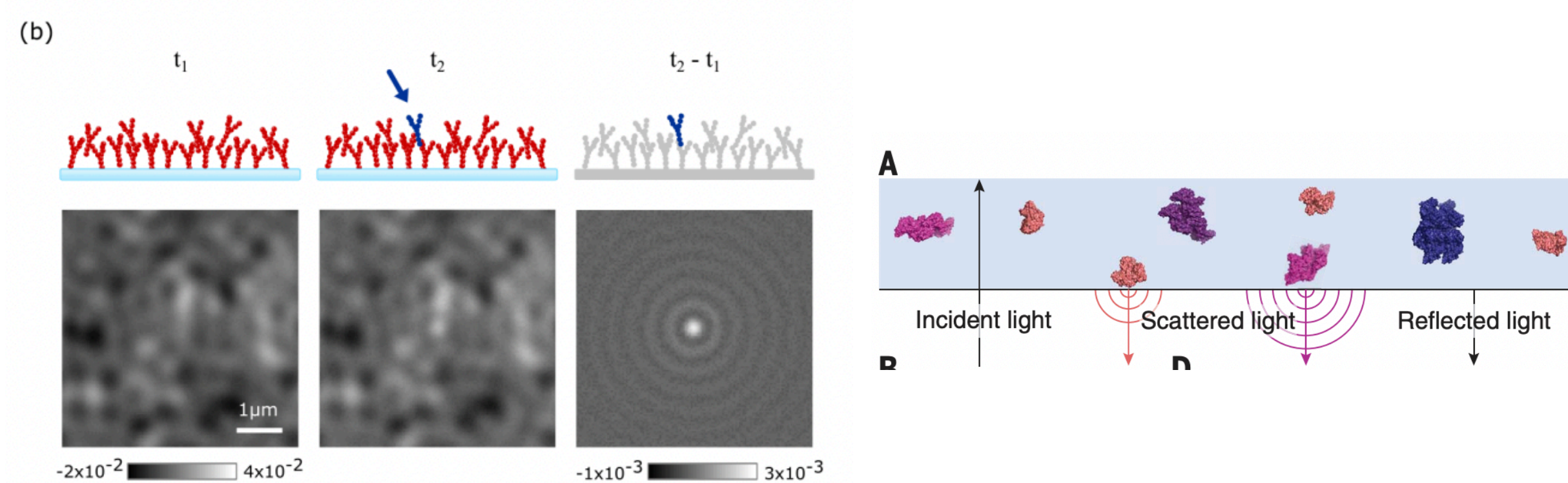


# Dynamic window sizes for contrast estimation in interferometric scattering

Ishan Taneja  
Scripps Research

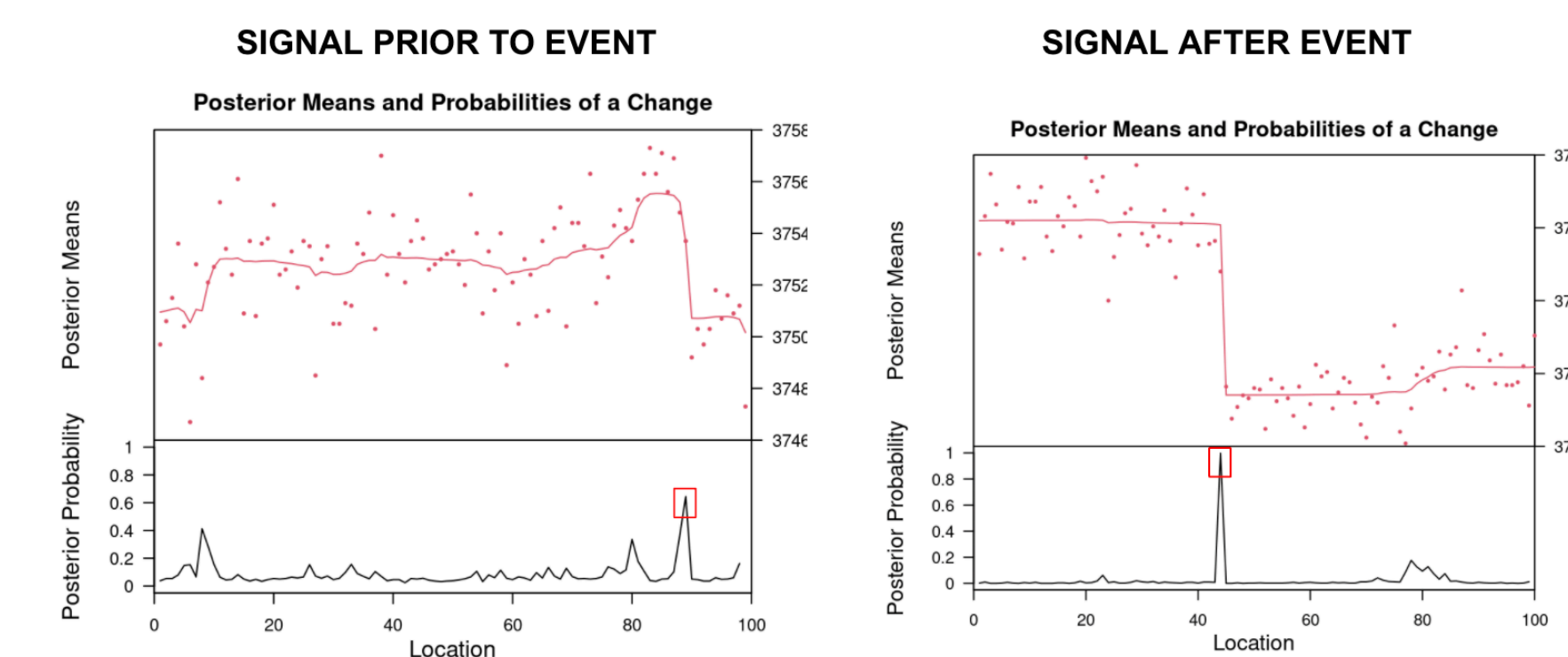
## Motivation

- Interferometric scattering microscopy is a label-free technique that can detect the mass of proteins by measuring their scattering contrast upon binding to a microscope coverslip
- This technique is very sensitive to background noise, and the ability to remove it is fundamental to making accurate measurements



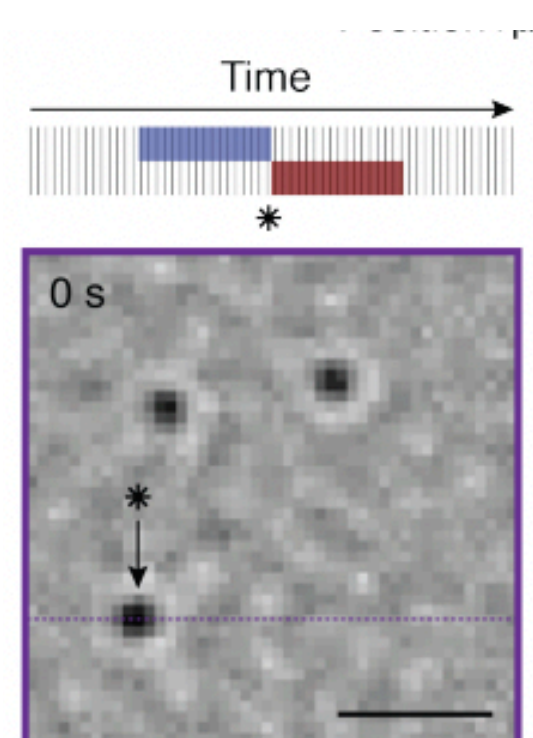
## New Technique

- A dynamic window size that is dependent on the data may be more effective than a fixed window size. We used changepoint detection to determine the optimal window sizes.



## Related Work

- Construction of a ratiometric image stack [2]

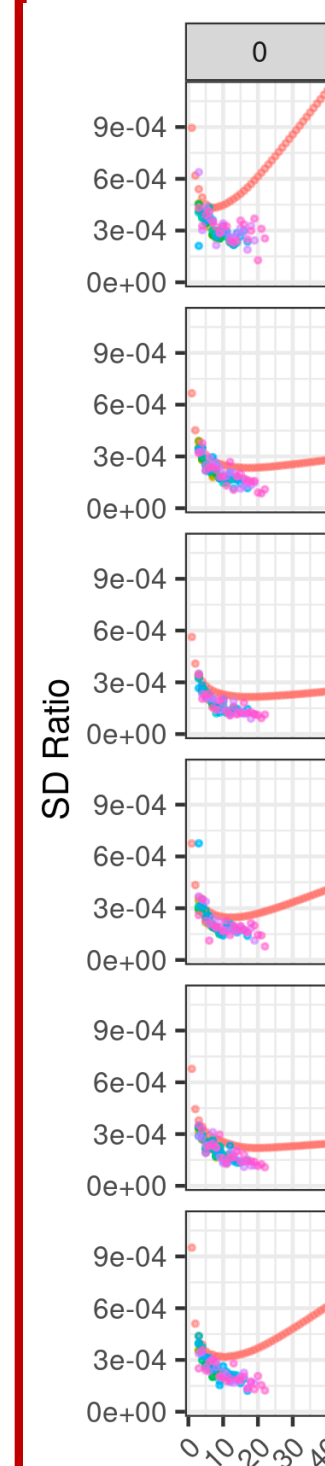


$$R_{w,i} = I_{i:i+w-1} / I_{i+w:i+2w-1} - 1$$

## References

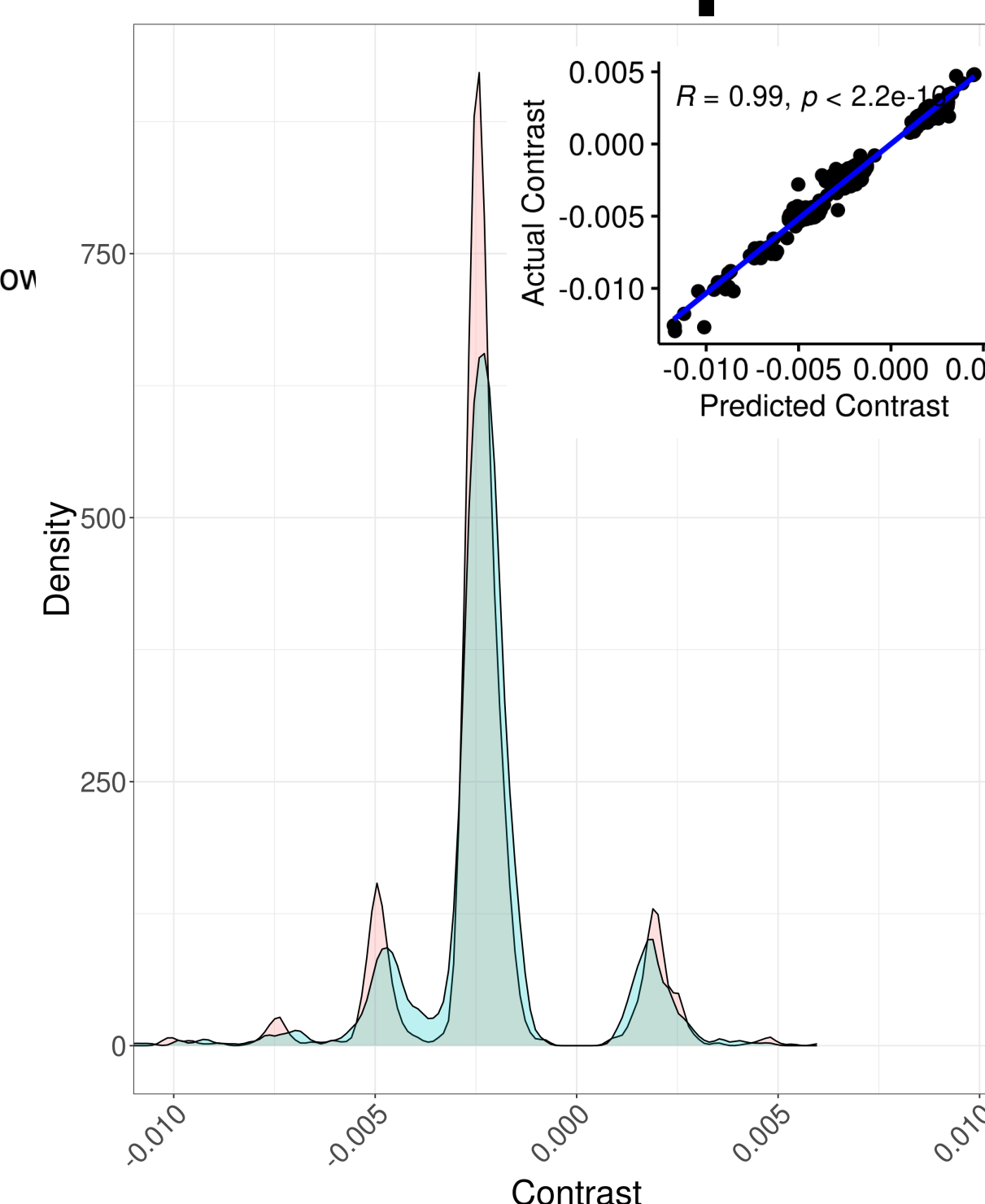
- Reza Gholami Mahmoodabadi et. al. 2020. "The Point Spread Function in Interferometric Scattering Microscopy (iSCAT). I. Aberrations in Defocusing and Axial Localization." *arXiv*.
- Gavin Young et al. 2018. "Quantitative Mass Imaging of Single Biological Macromolecules." *Science*.
- Richard W. Taylor et. al 2018. "Interferometric Scattering (iSCAT) Microscopy & Related Techniques." *arXiv*.

## Experimental Results



search\_window

- constant
- 20
- 25
- 30
- 35
- 40
- 45
- 50



method

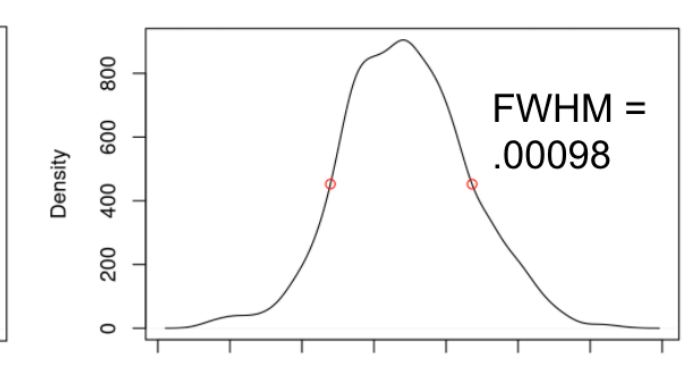
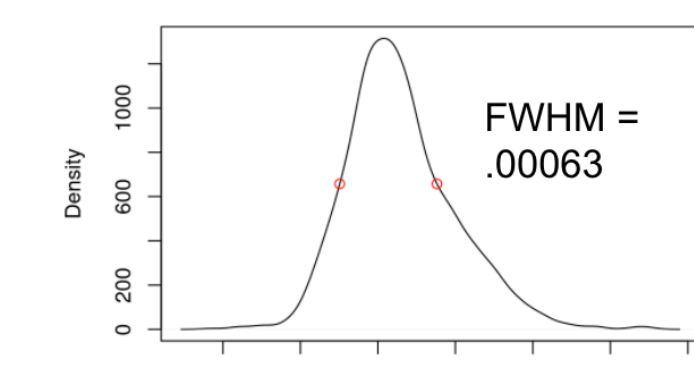
- Benchmark
- BayesCP

Method

Benchmark

Dynamic Window

Peak 1



Peak 2

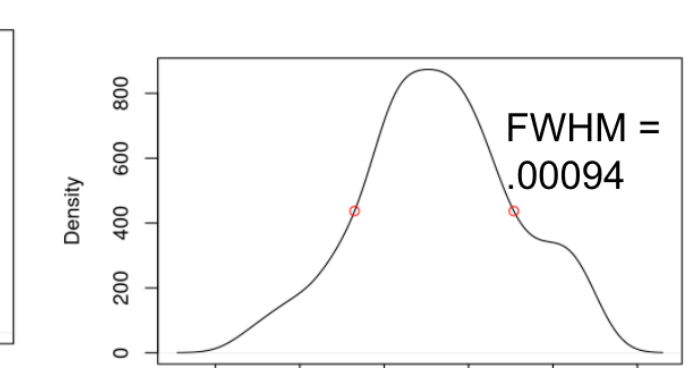
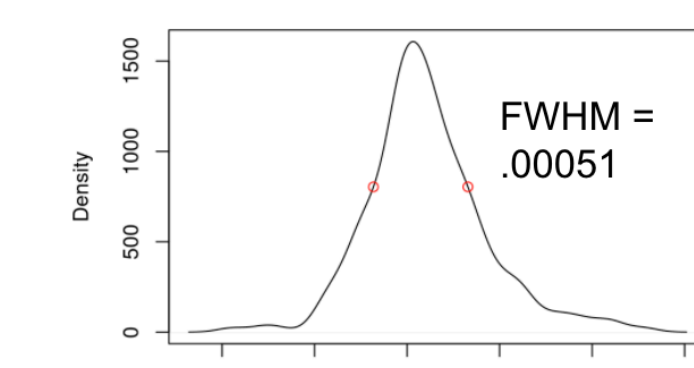


Fig. 1

Fig. 2

Fig. 3