

BISET: Biological Image Super-resolution Enhanced with Tensor

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 SIGNET team - I3S Lab, (Gérad Favier, Henrique Goulard) Signal processing, Tensor

iBV Lab, (Sébastien Schaub, Guillaume Sandoz)
Optics, biology (ion channels)

Academy 1 of UCA, 4, dec. 2017

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Fluorescence microscopy

- ► Structure of interest in living cells can be imaged through the microscope using fluorescent proteins
- Gene of fluorescent molecule can be combined with gene of proteins of structure we want to study Nobel Prize of chemistry 2008
- Illumination with a laser causes the fluorophores to emit photons



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Resolution limits

- physical diffraction limit of optical systems
- ▶ 200nm lateral, 400 axial.



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 Study of ion channels, membrane proteins that allow the passage of ions through the cell membrane. Resolution needed less than 50 nm.

Multi-Angle TIRF \longrightarrow axial super-resolution





Figure: TIRF Acquisitions.

- ▶ TIRF *Total Internal Reflexion Fluorescence* Microscope is a commercial system
- ▶ Multiple Angle TIRF prototype at iBV, no more than 3 prototypes in France,
- Reconstruction algorithm for the depth information, 30nm of resolution over 400nm thick layer over the coverslip.
- ► Method of choice to visualize processes around the plasma membrane



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3D super-resolution by mixing MA-TIRF and SIF

Optical axial Fluctuation \longrightarrow *lateral super-resolution*

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- use axial SIF *Super-resolution Intensity Fluctuation* (blinking): Acquisition of 100 images in 1 second,
- ▶ each image is blurred and low level SNR (signal on noise ratio), but the temporal fluctuation of fluorescent molecule will be used to localize precisely the fluorophore
- ▶ analysis of the temporal image series: cumulant, source separation

Mixing MA-TIRF and SIF \longrightarrow Tensor analysis

- ▶ MA-TIRF + SIF gives both axial and lateral super-resolution but big volume of 3D + t data to process
- ▶ Two possible tensor-based approaches:
 - 1. Separation of independent sources based on decomposing high-order cumulants of temporal image series
 - 2. Direct decomposition of MA-TIRF + SIF data: joint exploitation of several diversities for deterministic (blind) source separation.

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