Superresolution fluorescence microscopy

Leonid Keselman, Daniel Fernandes

Overview

What is "super-resolution"

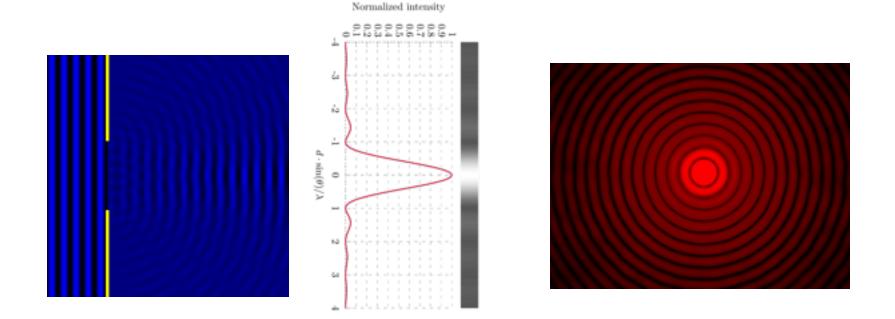
 a. Diffraction
 b. STORM

 Compressed Sensing

 a. Applied to STORM

 Light Sheet Imaging

 a. Lattice-Light Sheets



sources: Wikipedia (6wavelength=slitwidthblue.gif, Single_Slit_Diffraction_(english).svg, Beugungsscheibchen.k.720.jpg)



sources: font-awesome fa-bicycle

For typical cameras

$$d = 1.22 * \lambda * f #$$

Raleigh Criterion

iPhone 7: =1.22 * 650nm * f/1.8 =1.4 μm pixels are only 1.22 μm!

For microscopes

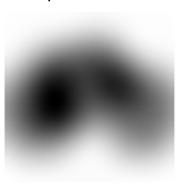
$$d = \frac{\lambda}{2n\sin\theta} = \frac{\lambda}{2NA}$$

Abbe diffraction limit

Typical Limit: = 500nm/(2 * 1.25) = 0.2 µm = 200nm Microtubules are ~24nm

NA is typically 0.1-0.4 for common lenses in air, up to 1.0-1.5 for oil lenses.

Rust, Bates, Zhuang. "Stochastic optical reconstruction microscopy (STORM) provides sub-diffraction-limit image resolution." *Nature Methods* 3.10 (2006)

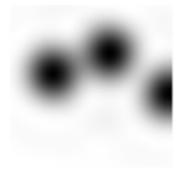


All pixels "on":

Random 1% of pixels "on"

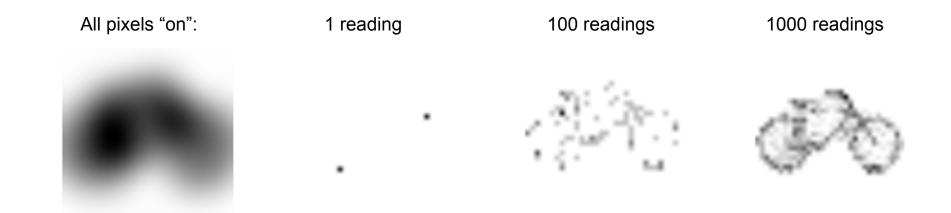
Random 1% of pixels "on"

Random 1% of pixels "on"

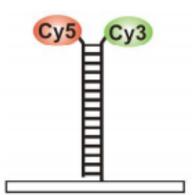


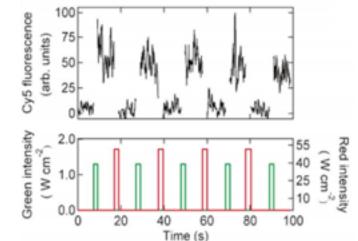


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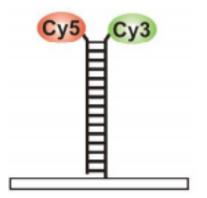


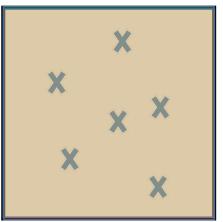
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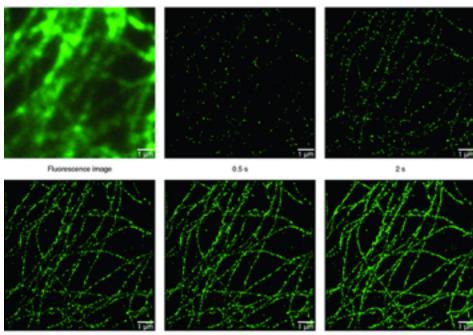
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(2010)







20 s

If your data is "compressible", you can take just a handful of random measurements, and, using "simple" math, you can reconstruct your data (with minimal error and high probability)

Emmanuel Candes and Terence Tao. "Near-optimal signal recovery from random projections: Universal encoding strategies?." *arXiv:math/0410542* (2004)

min
$$||x||_{\ell_1}$$
 subject to $||Ax - y||_{\ell_2} \le \epsilon$.

	1	2	3	4	5	6
1	Х	Х		Х		
0		Х	Х	Х		
2	Х			Х		Х

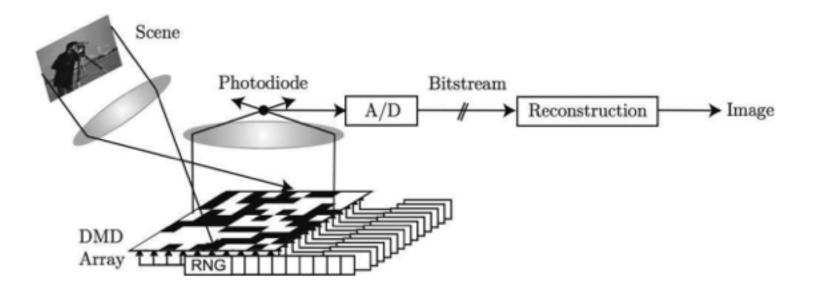
	1	2	3	4	5	6
1	Х	0		0		
0		0	0	0		
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Davenport, Duarte, Eldar, Kutynoik, Introduction to Compressed Sensing

Compressed Sensing

Duarte, et al. Single-Pixel Imaging via Compressive Sampling. (2008)



Compressed Sensing

Real Picture (65,536





CS Reconstruction (3 300 samples)



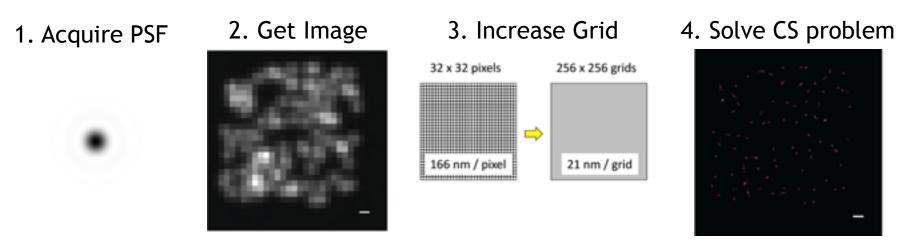
CS Reconstruction (<u>1,300 samples</u>)



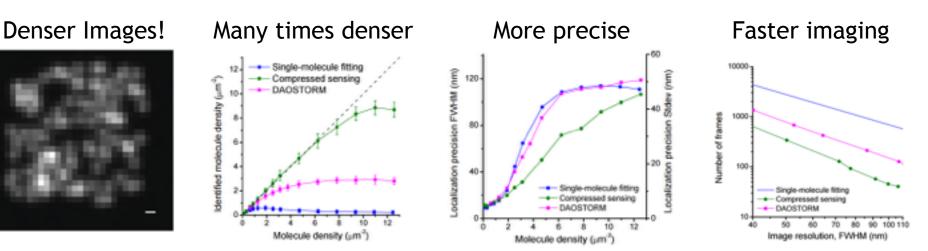
CS Reconstruction (6.500 samples)

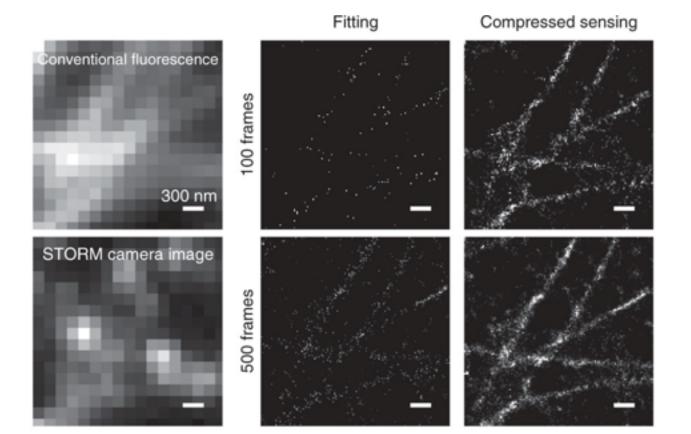


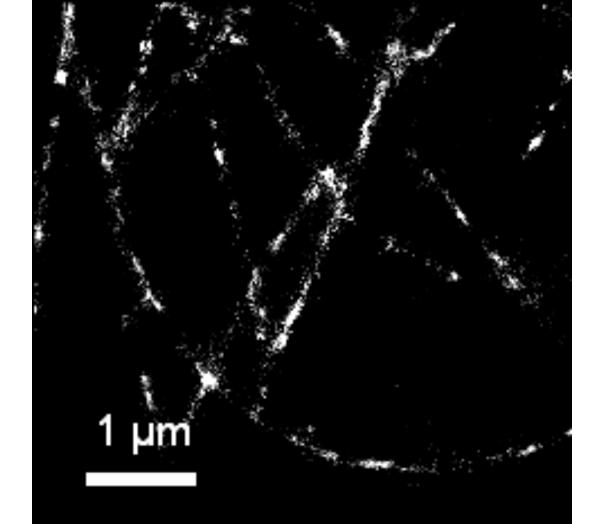
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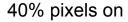


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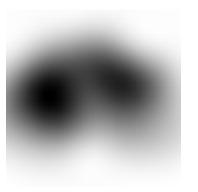


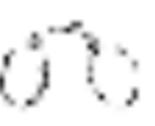


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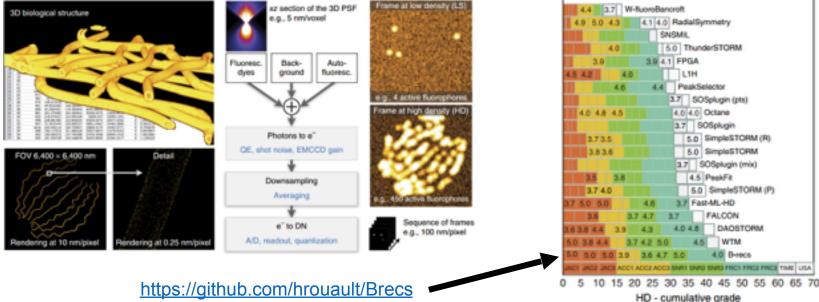




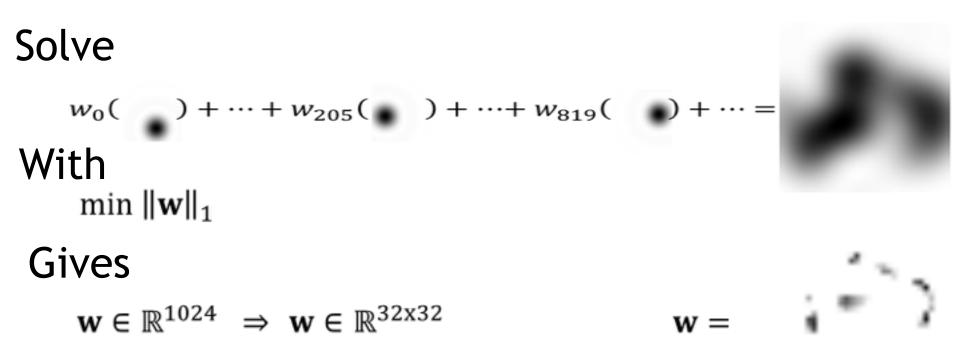
https://github.com/leonidk/cs371

Quantitative Comparison

Sage, Daniel, et al. "Quantitative evaluation of software packages for single-molecule localization microscopy." Nature Methods 12.8 (2015)



Extra Slides



DAOSTORM

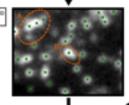
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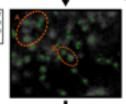
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DA0570RM schemati

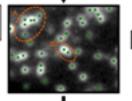
as candidate molecules



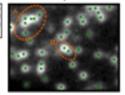
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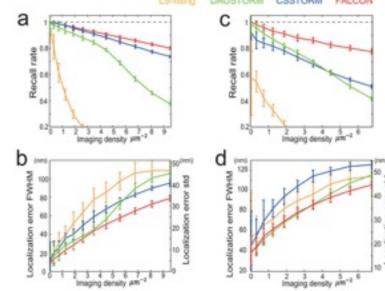


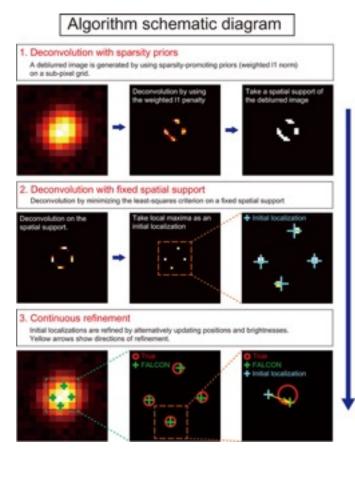
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FALCON

Min, Junhong, et al. "FALCON: fast and unbiased reconstruction of high-density super-resolution microscopy data " Scientific reports 4 (2014)





Supplementary Figure: Switching kinetics

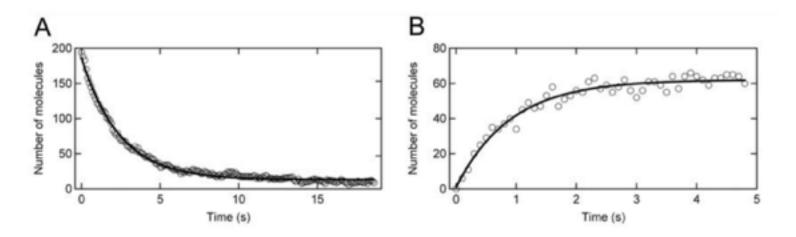


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Abstract

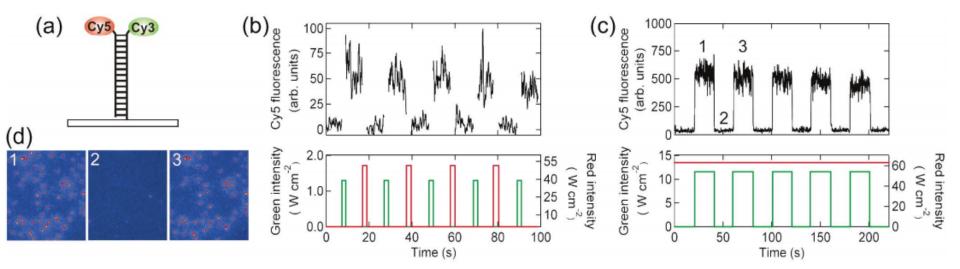
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This paper shows that if the objects of interest are sparse in a fixed basis or compressible, then it is possible to reconstruct f to within very high accuracy from a small number of random measurements by solving a simple linear program. More precisely, suppose that the *n*th largest entry of the vector |f| (or of its coefficients in a fixed basis) obeys $|f|_{(n)} \leq R \cdot n^{-1/p}$, where R > 0 and p > 0. Suppose that we take measurements $y_k = \langle f, X_k \rangle$, $k = 1, \ldots, K$, where the X_k are *N*-dimensional Gaussian vectors with independent standard normal entries. Then for each f obeying the decay estimate above for some $0 and with overwhelming probability, our reconstruction <math>f^{\sharp}$, defined as the solution to the constraints $y_k = \langle f^{\sharp}, X_k \rangle$ with minimal ℓ_1 norm, obeys

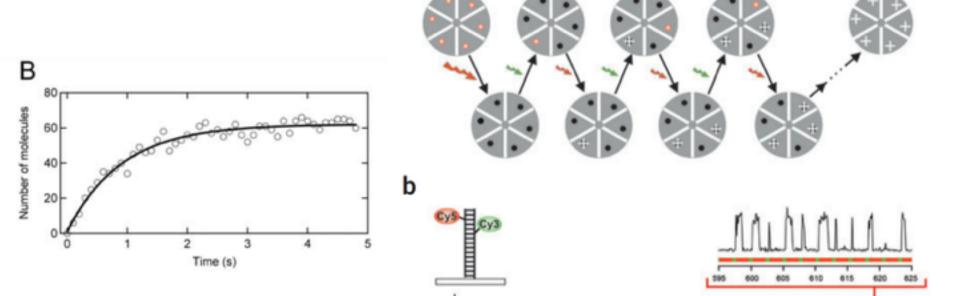
$$||f - f^{\sharp}||_{\ell_2} \le C_p \cdot R \cdot (K/\log N)^{-r}, \quad r = 1/p - 1/2.$$

There is a sense in which this result is optimal; it is generally impossible to obtain a higher accuracy from any set of K measurements whatsoever. The methodology extends to various other random measurement ensembles; for example, we show that similar results hold if one observes few randomly sampled Fourier coefficients of f. In fact, the results are quite general and require only two hypotheses on the measurement ensemble which are detailed.

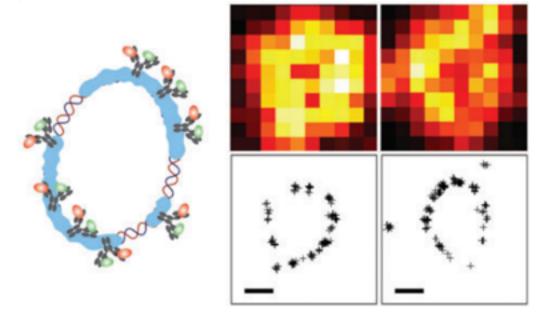
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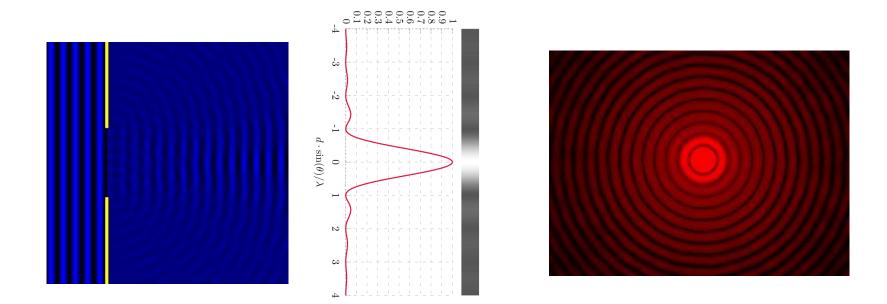
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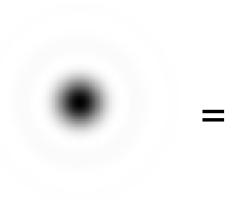
Normalized intensity

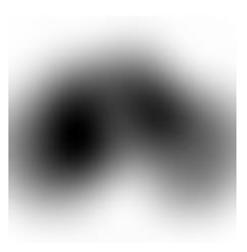


sources: Wikipedia (6wavelength=slitwidthblue.gif, Single_Slit_Diffraction_(english).svg, Beugungsscheibchen.k.720.jpg)



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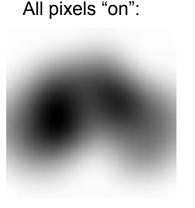
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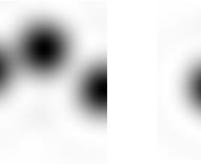
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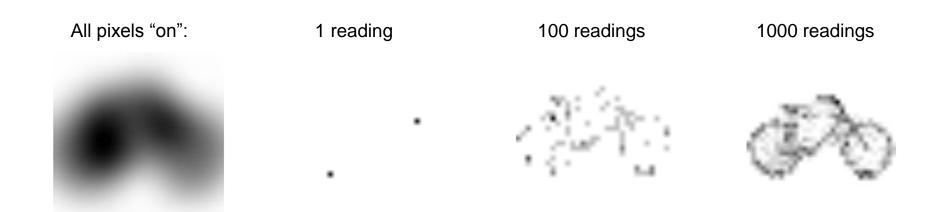
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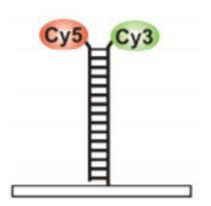
Random 1% of pixels "on" Random 1% of pixels "on" Random 1% of pixels "on"

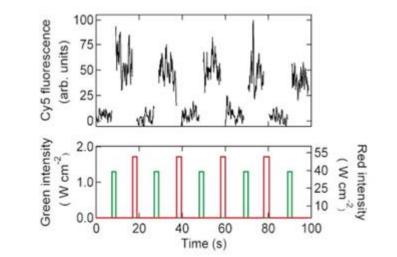


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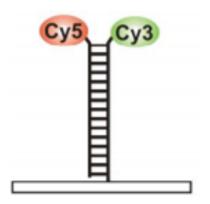


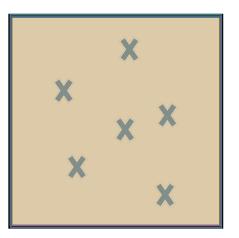
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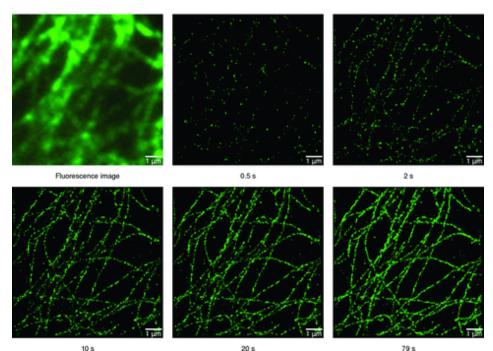


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	1	2	3	4	5	6
1	X	X		Х		
0		Х	Х	Х		
2	Х			Х		Х

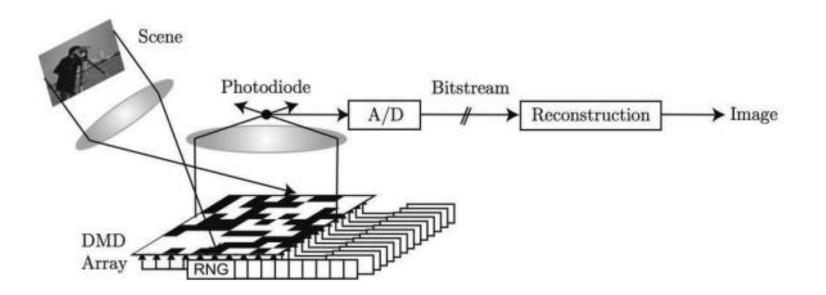
	1	2	3	4	5	6
1	X	0		0		
0		0	0	0		
2	Х			0		Х



Davenport, Duarte, Eldar, Kutynoik, Introduction to Compressed Sensing

Compressed Sensing

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Compressed Sensing

Real Picture (65,536 pixels)





CS Reconstruction (3,300 samples)



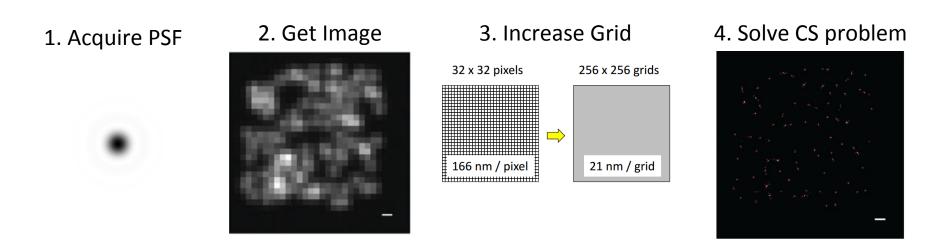
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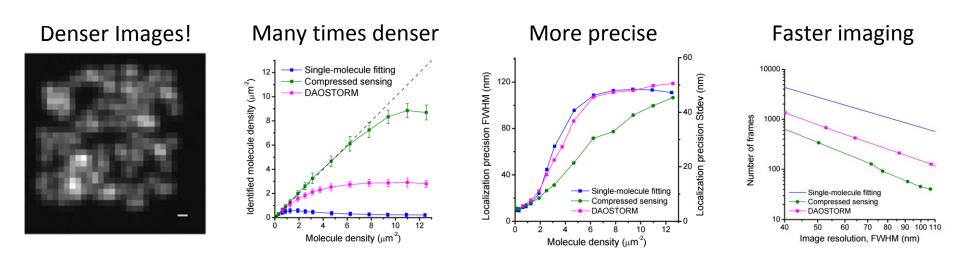
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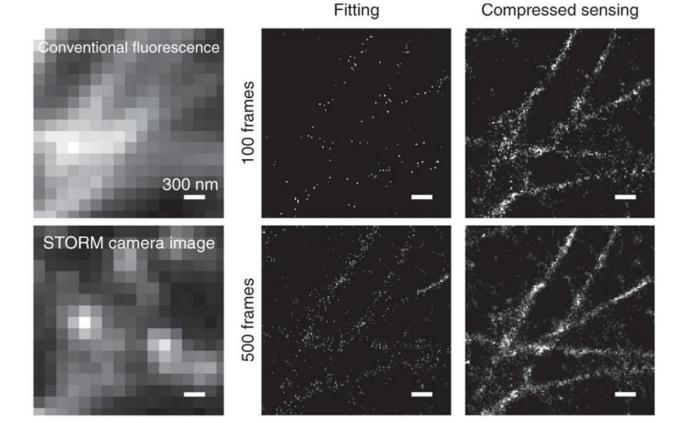


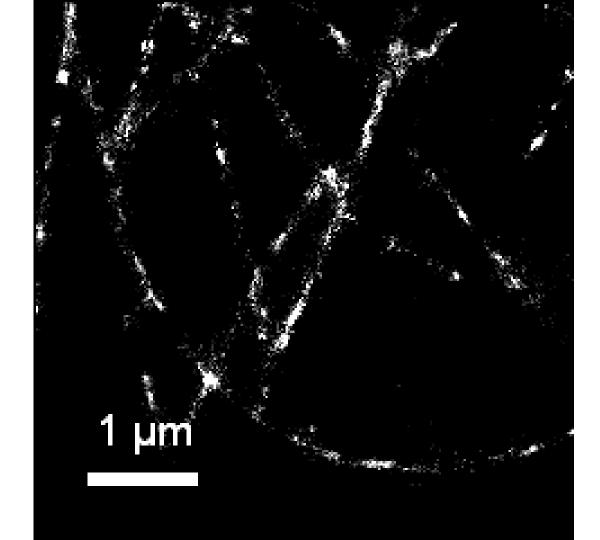
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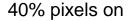


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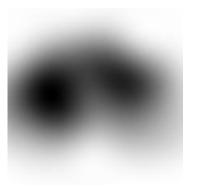


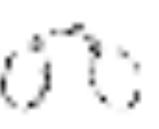


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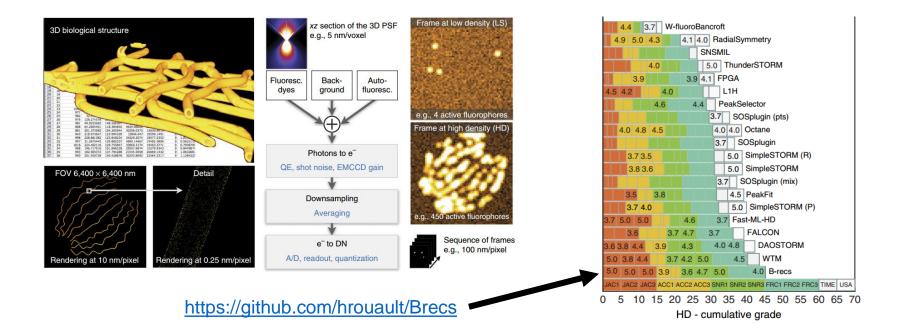






Quantitative Comparison

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Extra Slides

Solve

 $w_0() + \dots + w_{205}() + \dots + w_{819}() + \dots =$ With $\min ||\mathbf{w}||_1$

Gives

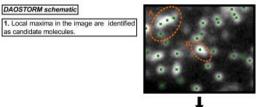
 $\mathbf{w} \in \mathbb{R}^{1024} \Rightarrow \mathbf{w} \in \mathbb{R}^{32x32}$

w =



DAOSTORM

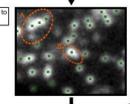
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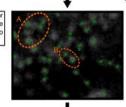
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DAOSTORM schematic

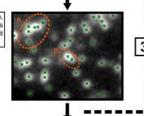
as candidate molecules.



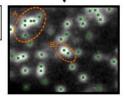
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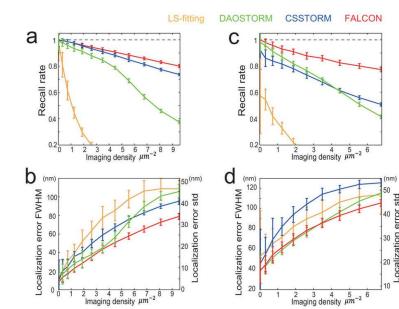


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FALCON

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Algorithm schematic diagram

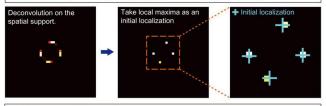
1. Deconvolution with sparsity priors

A deblurred image is generated by using sparsity-promoting priors (weighted I1 norm) on a sub-pixel grid.



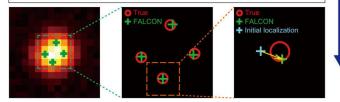
2. Deconvolution with fixed spatial support

Deconvolution by minimizing the least-squares criterion on a fixed spatial support



3. Continuous refinement

Initial localizations are refined by alternatively updating positions and brightnesses. Yellow arrows show directions of refinement.



Supplementary Figure: Switching kinetics

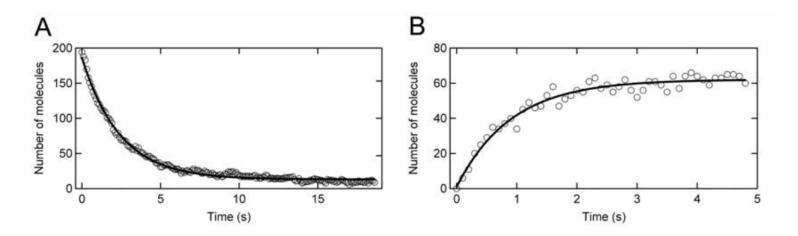


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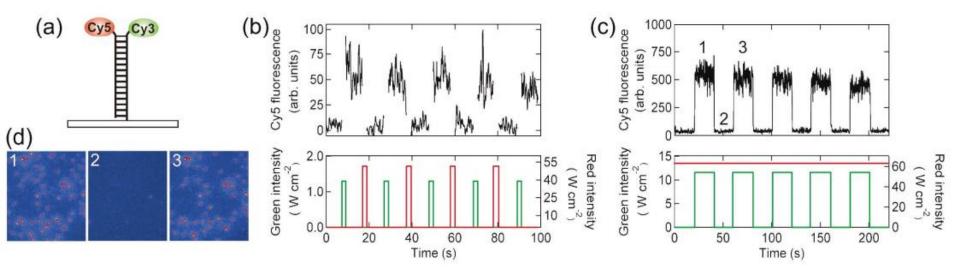
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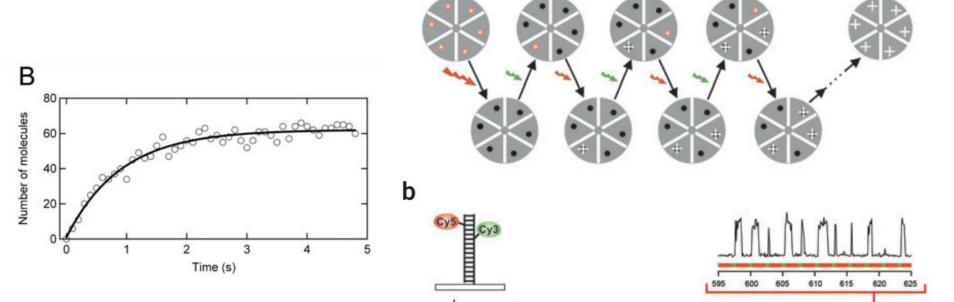
$$||f - f^{\sharp}||_{\ell_2} \le C_p \cdot R \cdot (K/\log N)^{-r}, \quad r = 1/p - 1/2.$$

There is a sense in which this result is optimal; it is generally impossible to obtain a higher accuracy from any set of K measurements whatsoever. The methodology extends to various other random measurement ensembles; for example, we show that similar results hold if one observes few randomly sampled Fourier coefficients of f. In fact, the results are quite general and require only two hypotheses on the measurement ensemble which are detailed.

Bates, Blosser, Zhuang. "Short-range spectroscopic ruler based on a single-molecule optical switch." *Physical review letters* 94.10 (2005)



Rust, Bates, Zhuang. "Stochastic optical reconstruction microscopy (STORM) provides sub-diffraction-limit image resolution." *Nature methods* 3.10 (2006) **a**



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