

# **Phase Retrieval, MAXCUT and Complex Semidefinite Programming**

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# Introduction

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Focus on the **phase retrieval** problem, i.e.

$$\begin{array}{ll} \text{find} & x \\ \text{such that} & |\langle a_i, x \rangle|^2 = b_i^2, \quad i = 1, \dots, n \end{array}$$

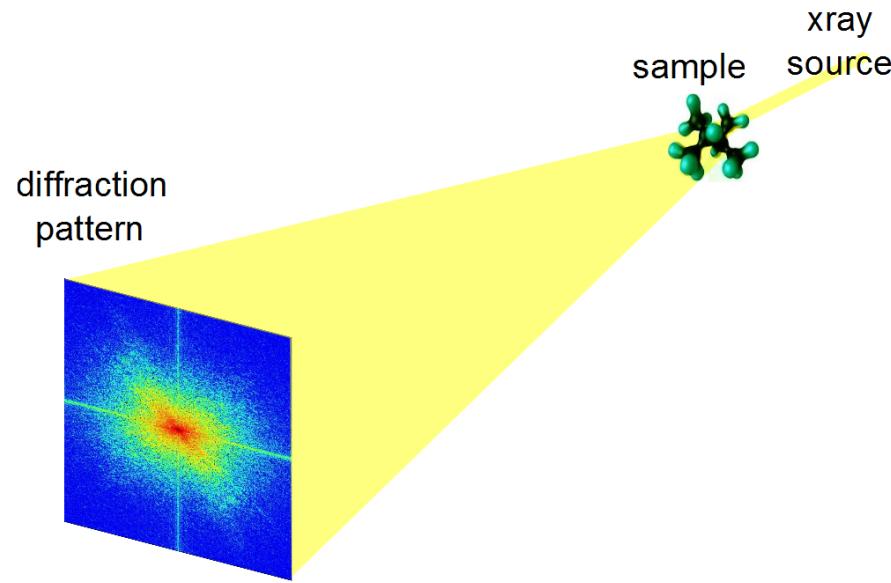
in the variable  $x \in \mathbb{C}^p$ .

- Reconstruct a signal  $x$  from the **amplitude of  $n$  linear measurements**.
- We seek a **tractable** procedure, i.e. a polynomial time algorithm with explicit approximation and complexity bounds.

# Introduction

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Applications in e.g. molecular imaging



(from [Candes et al., 2011b])

- CCD sensors only record the **magnitude** of diffracted rays, and loose the **phase**
- **Fraunhofer diffraction:** phase is required to invert the 2D Fourier transform

# Introduction

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Problem is almost 100 years old, infinite list of references. . .

## Algorithms

- Greedy algorithm [Gerchberg and Saxton, 1972]
- Classical survey of algorithms by [Fienup, 1982].
- NP-complete [Sahinoglou and Cabrera, 1991].
- Matrix completion formulation [Chai, Moscoso, and Papanicolaou, 2011] and [Candes, Strohmer, and Voroninski, 2011a]

## Applications

- X-ray and crystallography imaging [Harrison, 1993], diffraction imaging [Bunk et al., 2007] or microscopy [Miao et al., 2008].
- Audio signal processing [Griffin and Lim, 1984].

# Introduction

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Classical **greedy algorithm** [Gerchberg and Saxton, 1972].

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**Input:** An initial  $y^1 \in \mathbb{C}^n$ , i.e. such that  $|y^1| = b$ .

1: **for**  $k = 1, \dots, N - 1$  **do**

2:   Set

$$w = AA^\dagger y^k$$

3:   Set

$$y_i^{k+1} = b_i \frac{w}{|w|}, \quad i = 1, \dots, n.$$

4: **end for**

**Output:**  $y_N \in \mathbb{C}^n$ .

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Very similar to **alternating projections**:

- Project on  $\mathcal{R}(A)$ .
- Adjust the magnitude to match  $b$
- Repeat . . .

# Introduction

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- [Chai et al., 2011] and [Candes et al., 2011a] use a **lifting** procedure from [Shor, 1987, Lovász and Schrijver, 1991] to write

$$|\langle a_i, x \rangle|^2 = b_i^2 \iff \mathbf{Tr}(a_i a_i^* x x^*) = b_i^2$$

and formulate phase recovery as a **matrix completion** problem

$$\begin{aligned} & \text{Minimize} && \mathbf{Rank}(X) \\ & \text{such that} && \mathbf{Tr}(a_i a_i^* X) = b_i^2, \quad i = 1, \dots, n \\ & && X \succeq 0 \end{aligned}$$

in the matrix  $X \in \mathbf{H}_p$ .

- [Recht et al., 2007, Candes and Recht, 2008, Candes and Tao, 2010] show that under certain conditions on  $A$  and  $x_0$ , it suffices to solve

$$\begin{aligned} & \text{Minimize} && \mathbf{Tr}(X) \\ & \text{such that} && \mathbf{Tr}(a_i a_i^* X) = b_i^2, \quad i = 1, \dots, n \\ & && X \succeq 0 \end{aligned}$$

which is a (convex) **semidefinite program** in  $X \in \mathbf{H}_p$ .

# Outline

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- Introduction
- **MAXCUT formulation**
- Tightness
- Algorithms & Structure
- Numerical Results

# MAXCUT formulation

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We can **decouple** the phase and magnitude reconstruction problems.

- In the noiseless case, write  $Ax = \text{diag}(b)u$  where  $u \in \mathbb{C}^n$  is a **phase vector** with  $|u_i| = 1$ .
- The phase recovery problem can be written

$$\min_{\substack{u \in \mathbb{C}^n, |u_i|=1, \\ x \in \mathbb{C}^p}} \|Ax - \text{diag}(b)u\|_2^2,$$

- The inner minimization problem in  $x$  is a standard least squares, with solution  $x = A^\dagger \text{diag}(b)u$ , so phase recovery becomes

$$\begin{aligned} & \text{minimize} && u^* Mu \\ & \text{subject to} && |u_i| = 1, \quad i = 1, \dots, n, \end{aligned}$$

in  $u \in \mathbb{C}^n$ , where the Hermitian matrix  $M = \text{diag}(b)(\mathbf{I} - AA^\dagger)\text{diag}(b)$  is positive semidefinite.

# MAXCUT formulation

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**MAXCUT.** Classical algorithm in combinatorial optimization.

- Given an undirected graph with weights  $w_{ij}$  on its edges  $(i, j)$ , *MaxCut* seeks to partition the vertices in two sets  $S$  and  $\bar{S}$  to **maximize the weight of the cut**

$$\max_{S \subset [1, n]} \sum_{\{i \in S, j \in \bar{S}\}} w_{ij}$$

- This can be written as a quadratic program

$$\begin{aligned} & \text{maximize} && x^T L x \\ & \text{subject to} && x_i^2 = 1, \quad i = 1, \dots, n \end{aligned}$$

where  $L$  is the graph Laplacian,  $L = \text{diag}(We) - W$ .

- Other interpretations as computing the ground state of spin glass models [Mezard and Montanari, 2009], computing mixed matrix norms [Nemirovski, 2005], approximating the CUT-norm [Alon and Naor, 2004], etc...

# MAXCUT formulation

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**MAXCUT.** We know a lot about how to find an approximate solution

$$\begin{aligned} & \text{maximize} && x^T L x \\ & \text{subject to} && x_i^2 = 1, \quad i = 1, \dots, n \end{aligned}$$

- [Goemans and Williamson, 1995] produce a polynomial algorithm with an approximation ratio of 0.878..., using a semidefinite relaxation

$$\begin{aligned} & \text{maximize} && \text{Tr}(XL) \\ & \text{subject to} && \text{diag}(X) = 1, \quad X \succeq 0 \end{aligned}$$

combined with a randomization argument.

- Approximating the solution with an approximation ratio better than 16/17 is NP-Hard, etc.

# MAXCUT formulation

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The **phase recovery** problem was written (in phase) as

$$\begin{aligned} & \text{minimize} && u^* M u \\ & \text{subject to} && |u_i| = 1, \quad i = 1, \dots, n, \end{aligned}$$

- We can write a relaxation for phase recovery similar to the MAXCUT SDP, and recycle all the efficient algorithms designed for MAXCUT to solve it.
- Nesterov [1998] produces approximation bounds for generic nonconvex quadratic programs. [Goemans and Williamson, 2001, Zhang and Huang, 2006] extend these results to complex valued problems and show a  $\pi/4$  approximation ratio for

$$\begin{aligned} & \text{maximize} && u^* M u \\ & \text{subject to} && |u_i| = 1, \quad i = 1, \dots, n, \end{aligned}$$

when  $M \succeq 0$ .

- Tightness results on very similar maximum-likelihood channel detection problems [Luo et al., 2003, Kisialiou and Luo, 2010, So, 2010].

# Outline

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- Introduction
- MAXCUT formulation
- **Tightness**
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# Outline

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**Tightness.** [Waldspurger, d'Aspremont, and Mallat, 2012] Write a semidefinite relaxation for phase recovery, similar to the MAXCUT SDP

$$\begin{aligned} & \text{Minimize} && \mathbf{Tr}(MU) \\ & \text{such that} && \mathbf{diag}(U) = 1, \quad X \succeq 0 \end{aligned}$$

call it **PhaseCut**. When do we perfectly recover the signal  $x$ ?

- [Candes et al., 2011a] show exact recovery w.h.p. for the **PhaseLift** relaxation

$$\begin{aligned} & \text{Minimize} && \mathbf{Tr}(X) \\ & \text{such that} && \mathbf{Tr}(a_i a_i^* X) = b_i^2, \quad i = 1, \dots, n \\ & && X \succeq 0 \end{aligned}$$

when  $n = O(p \log p)$  observations  $a_i$  are picked uniformly on the unit sphere.

- [Waldspurger et al., 2012] show

**PhaseCut is tight whenever PhaseLift is.**

- Empirically, slightly more robust to noise.

# Outline

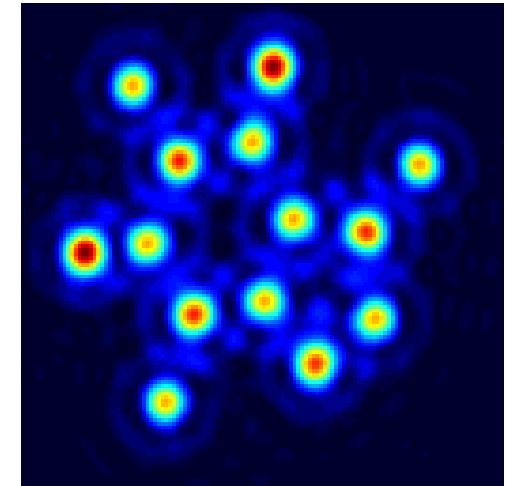
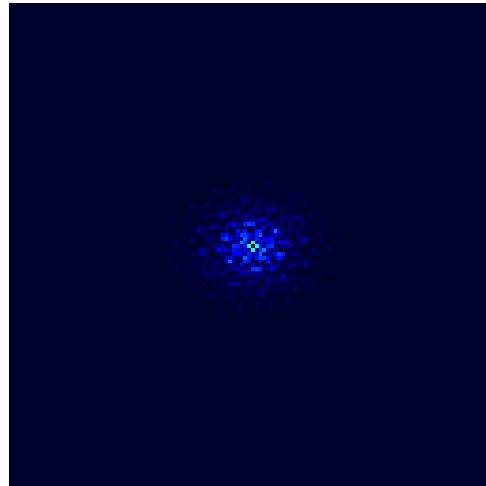
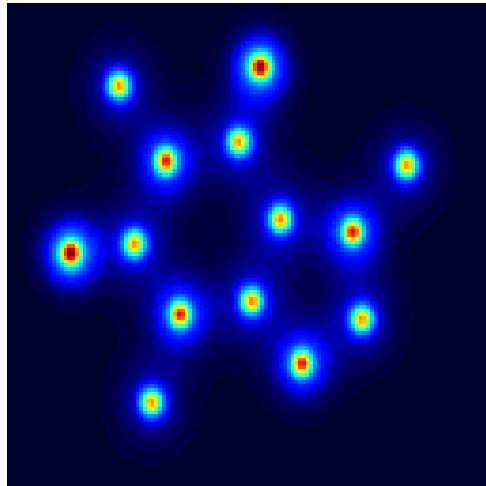
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# Sparsity: known support in 2D

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- Molecular imaging: the samples are approximately sparse with **known support**.
- Most of the coefficient in  $b$  are close to zero.



Electronic density for the caffeine molecule (left), its 2D FFT transform (diffraction pattern, center), the density reconstructed using 3% of the coefficients at the core of the FFT (right).

# Positivity

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- We observe the magnitude of the Fourier transform of a discrete nonnegative signal  $x \in \mathbb{R}^p$  so that

$$|\mathcal{F}x| = b$$

- We seek to reconstruct **positive** signals  $x \geq 0$ .
- This introduces additional **convex** restrictions on the phase vector  $u$ .

A function  $f : \mathbb{R}^s \mapsto \mathbb{C}$  is *positive semidefinite* if and only if the matrix  $B$  with  $B_{ij} = f(x_i - x_j)$  is Hermitian positive semidefinite for any sequence  $x_i \in \mathbb{R}^s$ .

## Theorem

**Bochner.** A function  $f : \mathbb{R}^s \mapsto \mathbb{C}$  is positive semidefinite if and only if it is the Fourier transform of a (finite) nonnegative Borel measure.

# Positivity

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- Reconstruct a phase vector  $u \in \mathbb{C}^n$  such that  $|u| = 1$  and

$$\mathcal{F}x = \text{diag}(b)u.$$

In 1D (for simplicity), if we define the Toeplitz matrix

$$B_{ij}(y) = y_{|i-j|+1}, \quad i, j = 1, \dots, p,$$

so that

$$B(y) = \begin{pmatrix} y_1 & y_2^* & & \cdots & & y_n^* \\ y_2 & y_1 & y_2^* & & \cdots & \\ & y_2 & y_1 & y_2^* & & \vdots \\ \vdots & & \ddots & \ddots & \ddots & \\ & \cdots & & y_2 & y_1 & y_2^* \\ y_n & & \cdots & & y_2 & y_1 \end{pmatrix}$$

- When  $\mathcal{F}x = \text{diag}(b)u$ , Bochner's theorem means  $B(\text{diag}(b)u) \succeq 0$  iff  $x \geq 0$ .
- The constraint  $B(\text{diag}(b)u) \succeq 0$  is a **linear matrix inequality** in  $u$ , hence is convex.

# Algorithms

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**PhaseCut** is a complex semidefinite program, written

$$\begin{aligned} & \text{Minimize} && \mathbf{Tr}(MU) \\ & \text{such that} && \mathbf{diag}(U) = 1, \quad X \succeq 0 \end{aligned}$$

where  $U \in \mathbb{H}_n$  with  $n = Jp$ , where  $p$  is the size of the signal.

- The complexity of solving this SDP using the algorithm in Helmburg et al. [1996] is

$$O\left(J^{3.5} p^{3.5} \log \frac{1}{\epsilon}\right) \quad \text{and} \quad O\left(K J^2 p^{4.5} \log \frac{1}{\epsilon}\right)$$

for *PhaseCut* and *PhaseLift* respectively.

- Solving a generic linear system is  $O(p^3)$ , solving a LP is  $O(p^{3.5})$ . . .
- Using first-order solvers such as TFOCS [Becker et al., 2012], based on [Nesterov, 1983], the dependence on the dimension can be further reduced, to become

$$O\left(\frac{J^3 p^3}{\epsilon}\right) \quad \text{and} \quad O\left(\frac{K J p^3}{\epsilon}\right)$$

for solving *PhaseCut* and *PhaseLift* respectively, serious impact on precision.

# Algorithms

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## Block Coordinate Method. [Wen et al., 2009]

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**Input:** An initial  $X^0 = \mathbf{I}_n$  and  $\nu > 0$  (typically small). An integer  $N > 1$ .

1: **for**  $k = 1, \dots, N$  **do**

2:   Pick  $i \in [1, n]$ .

3:   Compute

$$x = X_{i^c, i^c}^k M_{i^c, i} \quad \text{and} \quad \gamma = x^* M_{i^c, i}$$

4:   If  $\gamma > 0$ , set

$$X_{i^c, i}^{k+1} = X_{i, i^c}^{k+1*} = -\sqrt{\frac{1-\nu}{\gamma}} x$$

else

$$X_{i^c, i}^{k+1} = X_{i, i^c}^{k+1*} = 0.$$

5: **end for**

**Output:** A matrix  $X \succeq 0$  with  $\text{diag}(X) = 1$ .

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Writing  $i^c$  the index set  $\{1, \dots, i-1, i+1, \dots, n\}$ .

## Complexity.

- Each iteration only requires matrix vector products  $O(n^2)$ .
- Cost per iteration very similar to the greedy algorithm by [Gerchberg and Saxton, 1972].
- In signal applications, the matrix vector product can be computed efficiently using the **FFT**, and the cost per iteration is reduced to  $O(n \log n)$ .

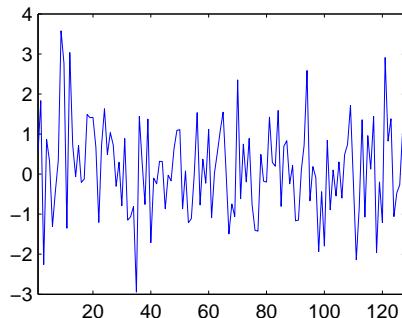
# Outline

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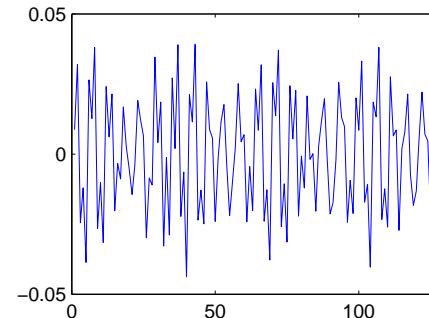
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# Numerical Experiments: 1D

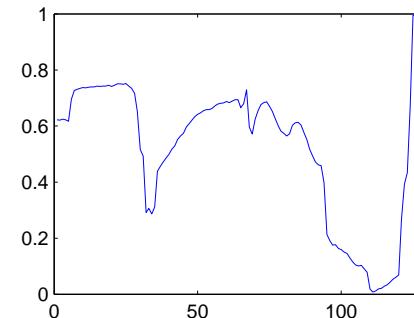
- Three random signal classes: (a) Gaussian white noise. (b) Sum of 6 sinusoids of random frequency & random amplitudes. (c) Random scan-line of an image.



(a)



(b)



(c)

- The linear sampling operator  $A$  is an **oversampled Fourier transform**, multiple filterings with **random filters**, or a **wavelet transform**.
- We measure the error both in signal and in modulus

$$\epsilon(x, \tilde{x}) = \min_{c \in \mathbb{C}, |c|=1} \frac{\|x - c\tilde{x}\|}{\|x\|} \quad \text{and} \quad \epsilon(|Ax|, |A\tilde{x}|) = \frac{\||Ax| - |A\tilde{x}|\|}{\|Ax\|}.$$

# Numerical Experiments: 1D

	Fourier	Random Filters	Wavelets
Gerchberg-Saxton	5%	49%	0%
<i>PhaseLift</i> with reweighting	3%	100%	62%
<i>PhaseCut</i>	4%	100%	100%

Percentage of perfect reconstruction from  $|Ax|$ , over 300 test signals, for the three different operators  $A$  (columns) and the three algorithms (rows).

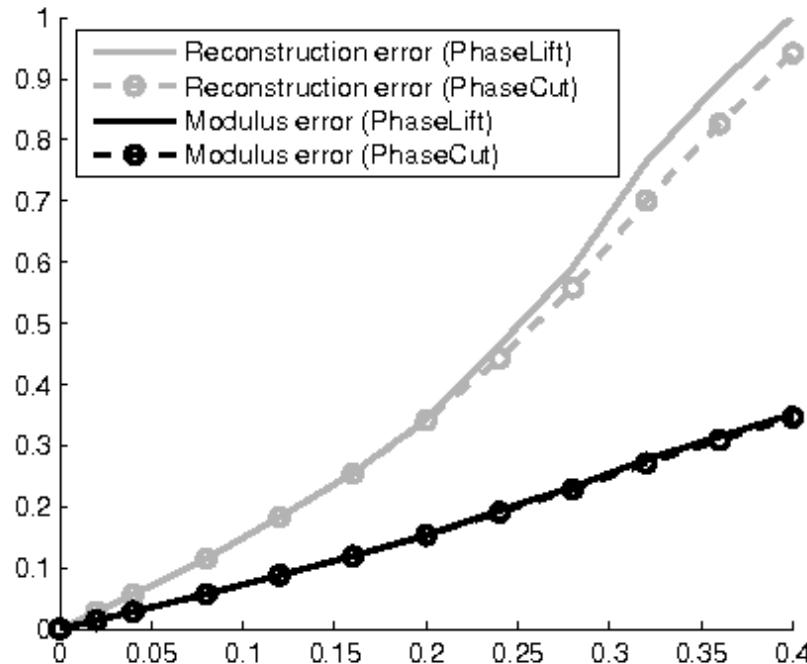
	Fourier	Random Filters	Wavelets
Gerchberg-Saxton	0.9	1.2	1.3
<i>PhaseLift</i> with reweighting	0.8	exact	0.5
<i>PhaseCut</i>	0.8	exact	exact

Average relative signal reconstruction error  $\epsilon(\tilde{x}, x)$  over all test signals that are not perfectly reconstructed, for each operator  $A$  and each algorithm.

	Fourier	Random Filters	Wavelets
Gerchberg-Saxton	$9 \cdot 10^{-4}$	0.2	0.3
<i>PhaseLift</i> with reweighting	$5 \cdot 10^{-4}$	exact	$8 \cdot 10^{-2}$
<i>PhaseCut</i>	$6 \cdot 10^{-4}$	exact	exact

Average relative error  $\epsilon(|A\tilde{x}|, |Ax|)$  of coefficient amplitudes, over all test signals that are not perfectly reconstructed, for each operator  $A$  and each algorithm.

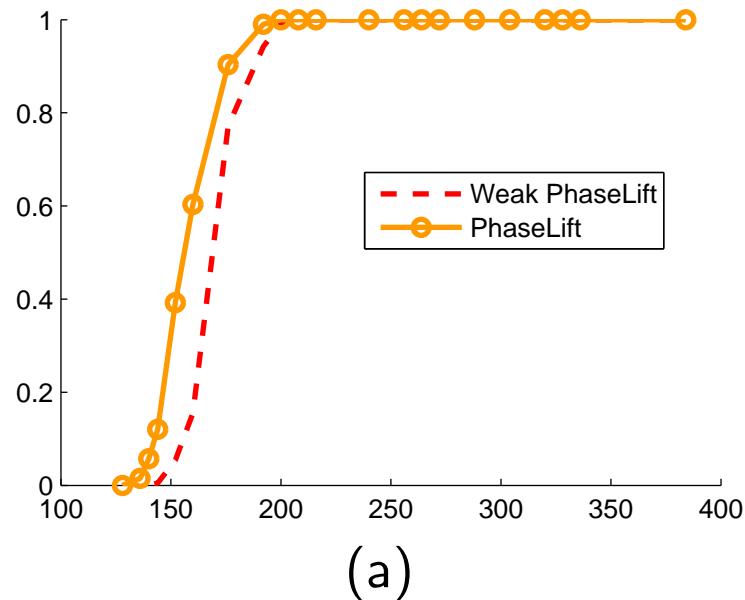
# Numerical Experiments: 1D



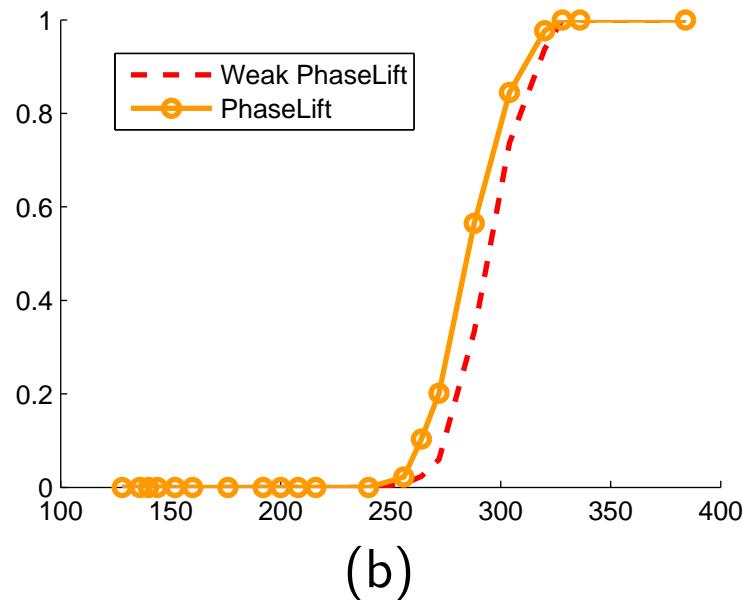
Mean performances of *PhaseLift* and *PhaseCut*, followed by some greedy iterations, for 4 gaussian random illumination filters. The  $x$ -axis represents the relative noise level,  $\|b_{\text{noise}}\|_2/\|Ax\|_2$  and the  $y$ -axis the relative error on the result (signal and modulus).

# Numerical Experiments: 1D

[Demanet and Hand, 2012] show that the solution to the relaxation is unique (trace minimization is unnecessary).



(a)



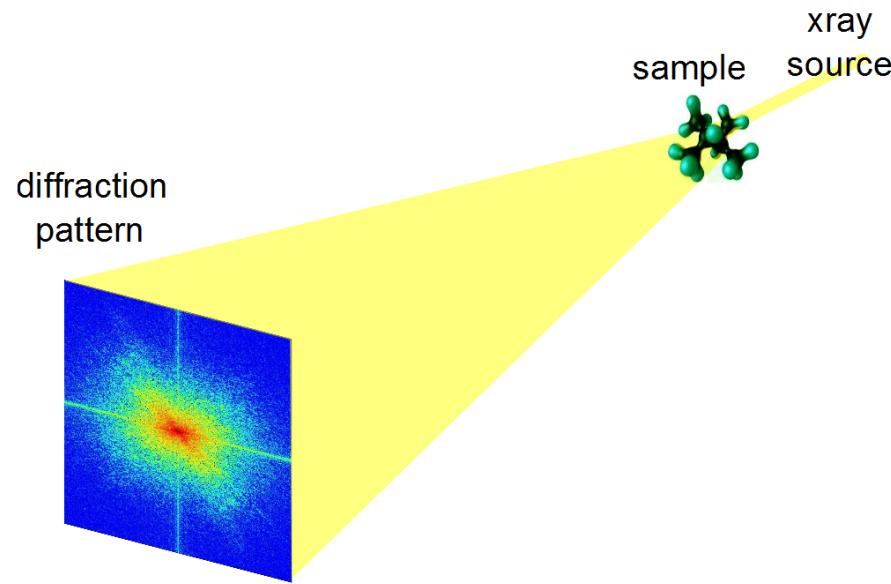
(b)

*PhaseLift* performance, for 64-sized signals, as a function of the number of measurements.

- (a) Proportion of reconstructed signals, postprocessing using after GS iterations.
- (b) Proportion of rank 1 (tight) solutions in the relaxation.

# Numerical Experiments: 2D

Applications in e.g. molecular imaging

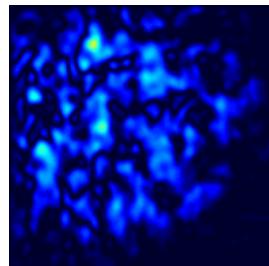


(from [Candes et al., 2011b])

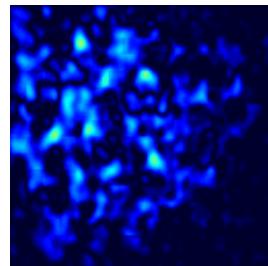
- CCD sensors only record the **magnitude** of diffracted rays, and loose the **phase**
- **Fraunhofer diffraction:** phase is required to invert the 2D Fourier transform
- Simulate diffraction using molecules from PDB and Poisson noise.

# Numerical Experiments: 2D

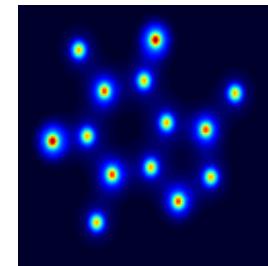
1 ill.,  $\alpha = 0$



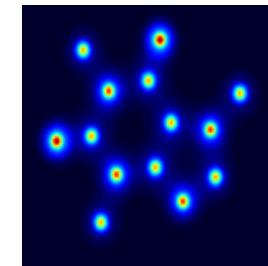
2 ill.,  $\alpha = 0$



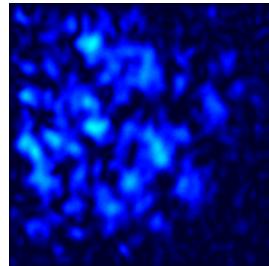
3 ill.,  $\alpha = 0$



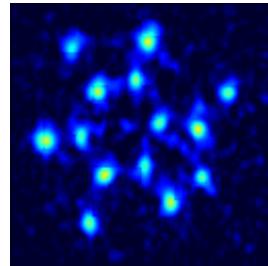
4 ill.,  $\alpha = 0$



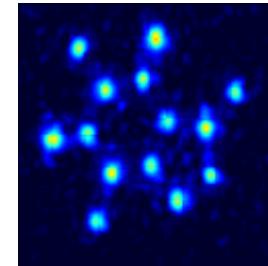
1 ill.,  $\alpha = 10^{-3}$



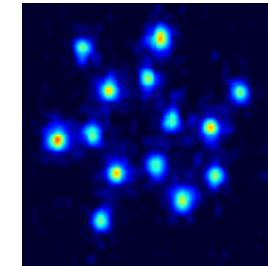
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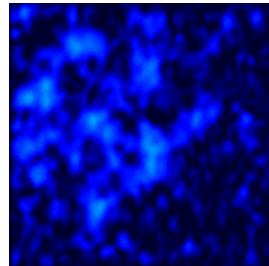
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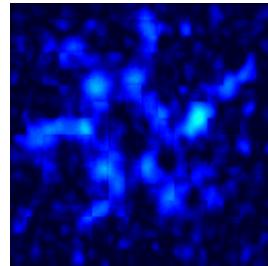
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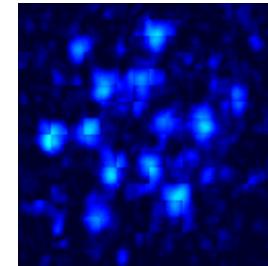
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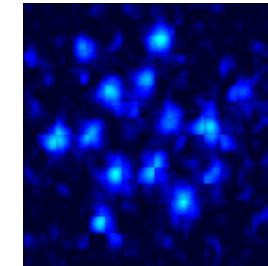
2 ill.,  $\alpha = 10^{-2}$



3 ill.,  $\alpha = 10^{-2}$



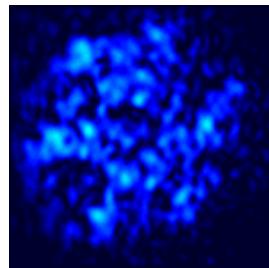
4 ill.,  $\alpha = 10^{-2}$



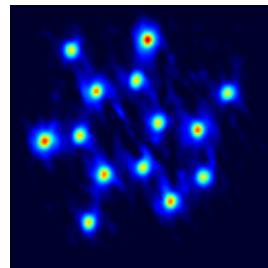
Solution of the greedy algorithm on caffeine molecule, for various values of the number of filters and noise level  $\alpha$ .

# Numerical Experiments: 2D

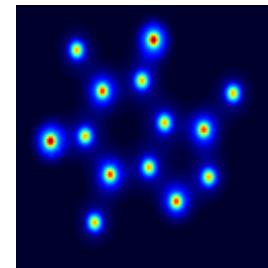
1 ill.,  $\alpha = 0$



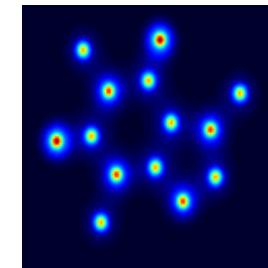
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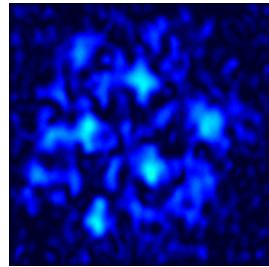
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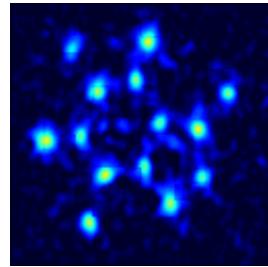
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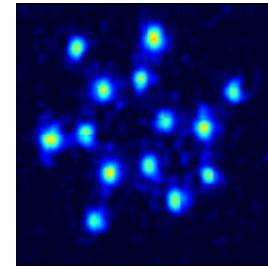
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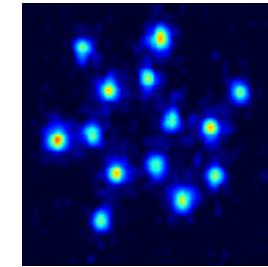
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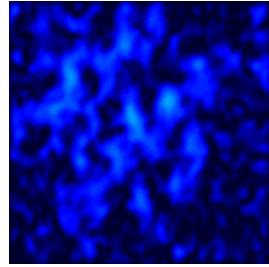
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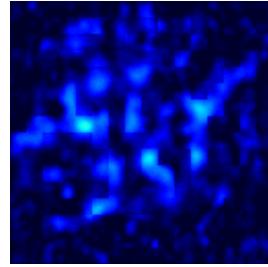
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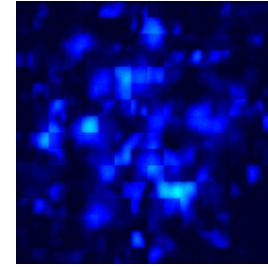
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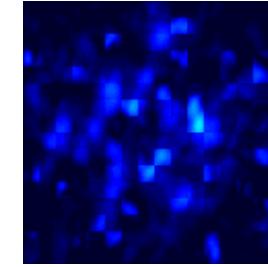
2 ill.,  $\alpha = 10^{-2}$



3 ill.,  $\alpha = 10^{-2}$



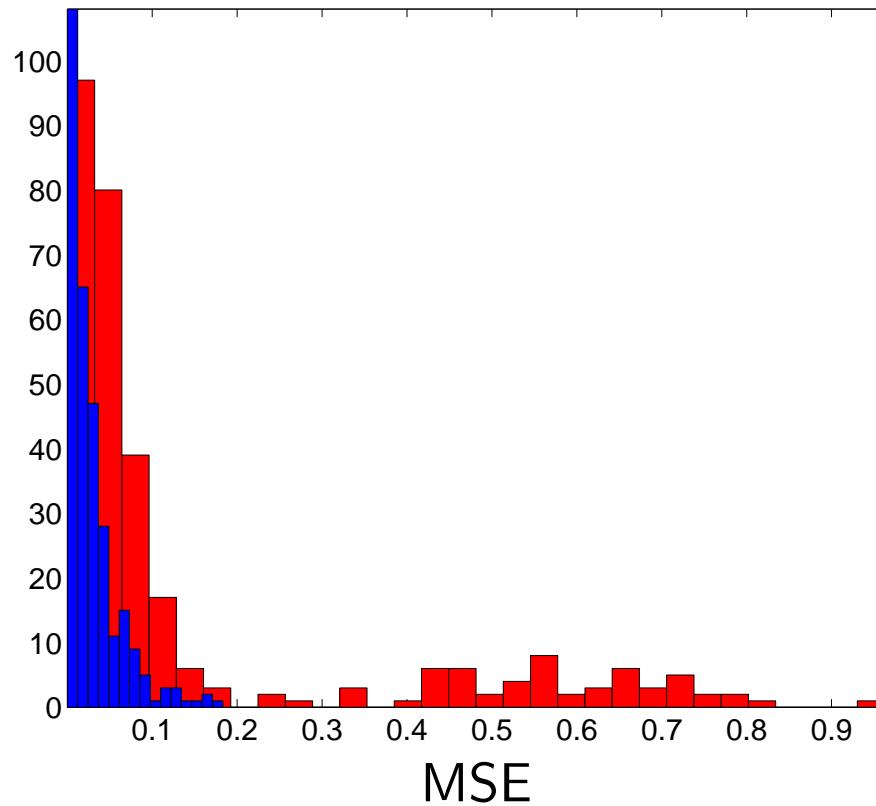
4 ill.,  $\alpha = 10^{-2}$



Solution of the semidefinite relaxation algorithm followed by greedy refinements, for various values of the number of filters and noise level  $\alpha$ .

# Numerical Experiments: 2D

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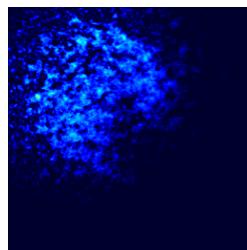


MSE between reconstructed image and true image for 2 illuminations without noise, using SDP then Fienup (**blue**), and Fienup only (**red**).

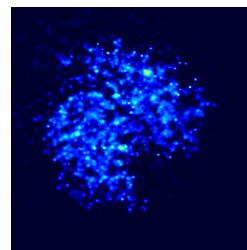
# Numerical Experiments: 2D

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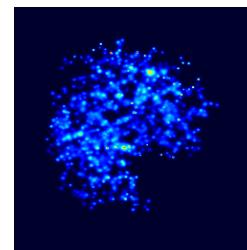
1 ill.,  $\alpha = 0$



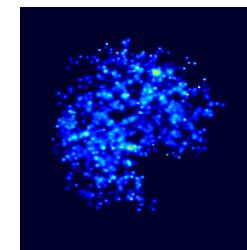
2 ill.,  $\alpha = 0$



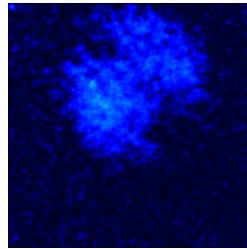
3 ill.,  $\alpha = 0$



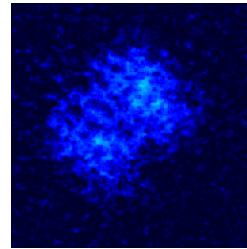
4 ill.,  $\alpha = 0$



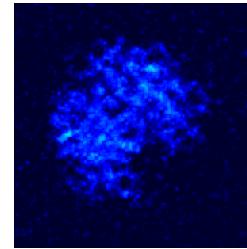
1 ill.,  $\alpha = 10^{-3}$



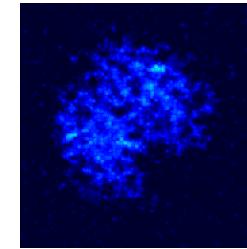
2 ill.,  $\alpha = 10^{-3}$



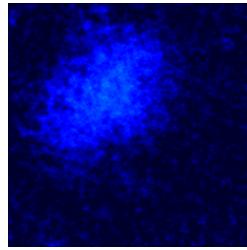
3 ill.,  $\alpha = 10^{-3}$



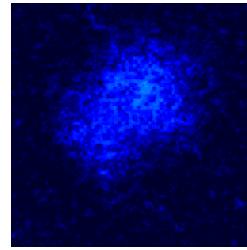
4 ill.,  $\alpha = 10^{-3}$



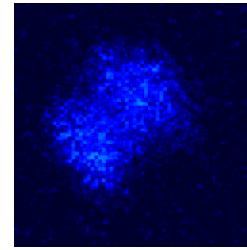
1 ill.,  $\alpha = 10^{-2}$



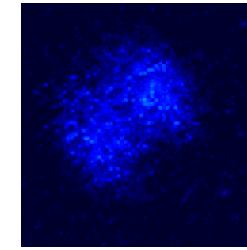
2 ill.,  $\alpha = 10^{-2}$



3 ill.,  $\alpha = 10^{-2}$



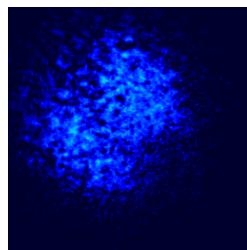
4 ill.,  $\alpha = 10^{-2}$



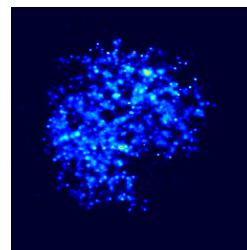
Solution of the greedy algorithm on 2LYZ (Lysozyme), for various values of the number of filters and noise level  $\alpha$ .

# Numerical Experiments: 2D

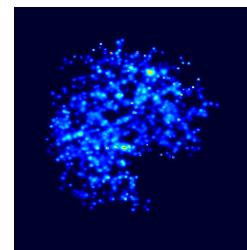
1 ill.,  $\alpha = 0$



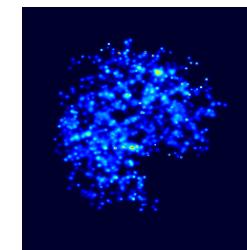
2 ill.,  $\alpha = 0$



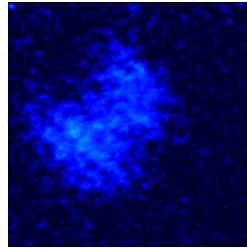
3 ill.,  $\alpha = 0$



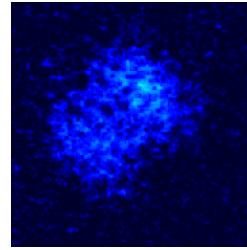
4 ill.,  $\alpha = 0$



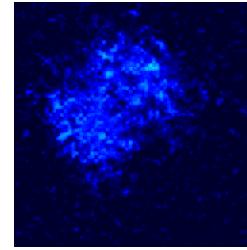
1 ill.,  $\alpha = 10^{-3}$



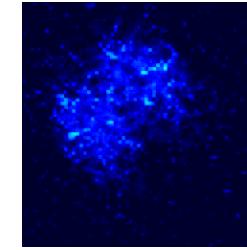
2 ill.,  $\alpha = 10^{-3}$



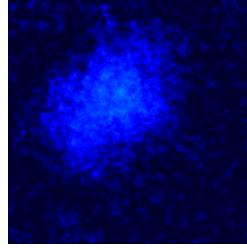
3 ill.,  $\alpha = 10^{-3}$



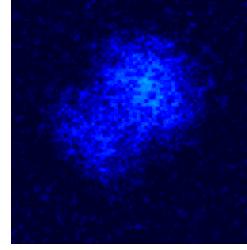
4 ill.,  $\alpha = 10^{-3}$



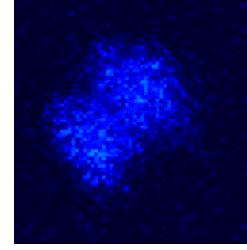
1 ill.,  $\alpha = 10^{-2}$



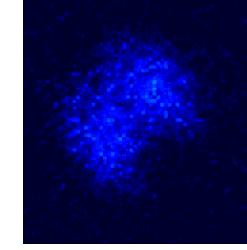
2 ill.,  $\alpha = 10^{-2}$



3 ill.,  $\alpha = 10^{-2}$



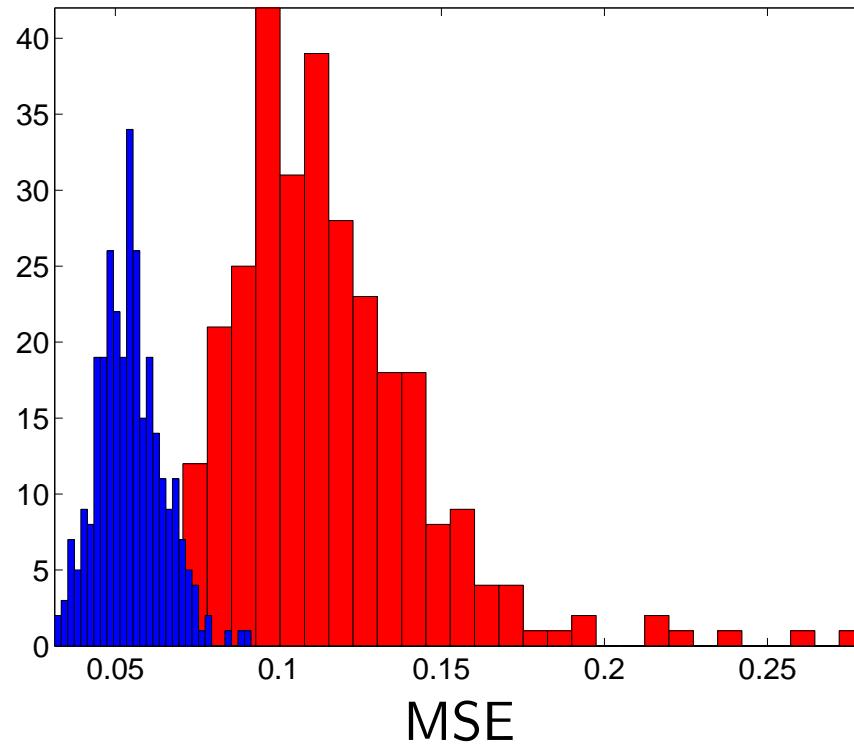
4 ill.,  $\alpha = 10^{-2}$



Solution of the semidefinite relaxation algorithm followed by greedy refinements on 2LYZ (Lysozyme), for various values of the number of filters and noise level  $\alpha$ .

# Numerical Experiments: 2D

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MSE between reconstructed image and true image for 2 illuminations of 2LYZ without noise, using SDP then Fienup (**blue**), and Fienup only (**red**).

# Conclusion

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- Write the phase recovery problem as a MAXCUT like problem.
- Tightness properties equivalent to the matrix completion approach.
- Very fast/scalable algorithms.

Open questions. . . .

- Tightness results in the noisy case, or in the positive case?
- Is the SDP relaxation optimal?

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