

Estimation below the identifiability limit with application to cryo-EM

Tamir Bendory

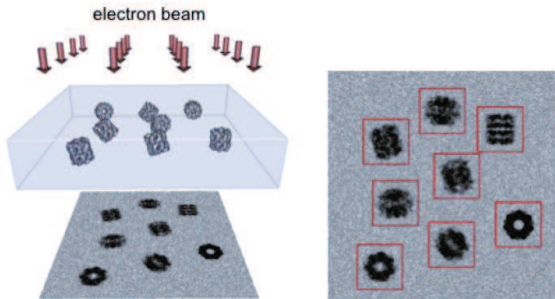
January 23, 2018

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The Program in Applied and Computational Mathematics
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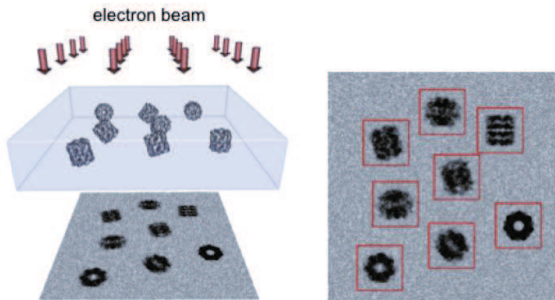
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- 1 Motivation: Cryo-EM
- 2 The multireference alignment problem and its invariants
- 3 Application to cryo-EM: 2D classification
- 4 Future research directions

Single particle reconstruction using cryo-EM



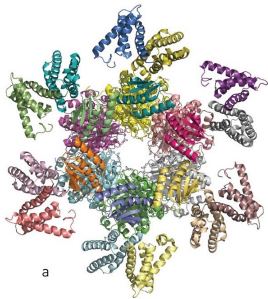
Single particle reconstruction using cryo-EM



Why cryo-EM?

- Mapping the structure of molecules without crystallizing them
- Imaging of heterogeneous samples, with mixtures of molecules or multiple conformations

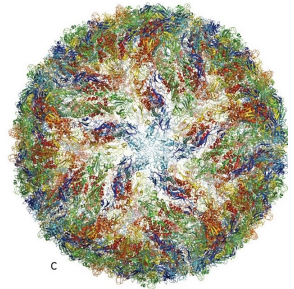
The resolution revolution



A protein complex that governs the circadian rhythm (sleep/wake cycle)

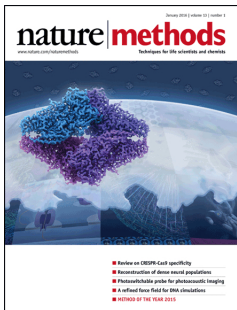


A sensor of the type that reads pressure changes in the ear



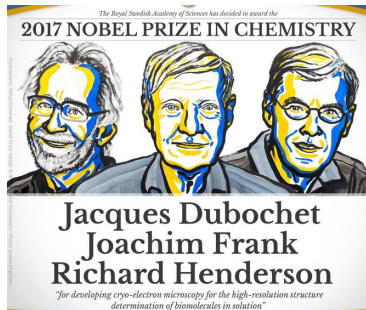
The Zika virus

Exciting times for cryo-EM



Method of the Year 2015

“Single-particle cryo-EM is our choice for Method of the Year 2015 for its newfound ability to solve protein structures at near-atomic resolution.”



Nobel Prize in Chemistry 2017

“for developing cryo-EM for the high-resolution structure determination of biomolecules in solution”

Image formation model and inverse problem

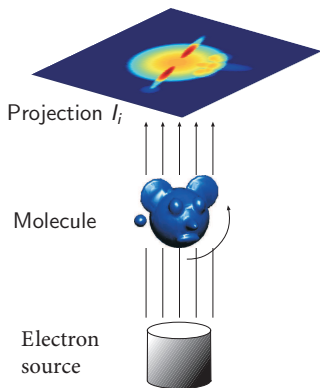


Image formation model

$$I_i = P(R_i \circ X) + \text{noise}, R_i \in SO(3)$$

Image formation model and inverse problem

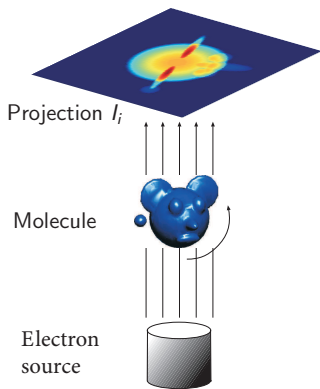


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The cryo-EM problem

Estimate X given I_1, \dots, I_N

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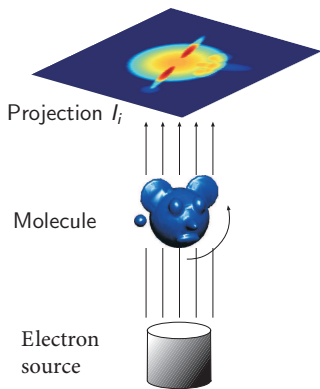


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The heterogeneity problem

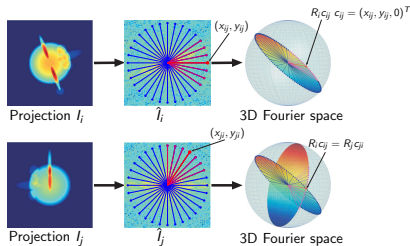
Estimate X_1, \dots, X_K given I_1, \dots, I_N

Fundamental challenges

- 1 Unknown viewing directions

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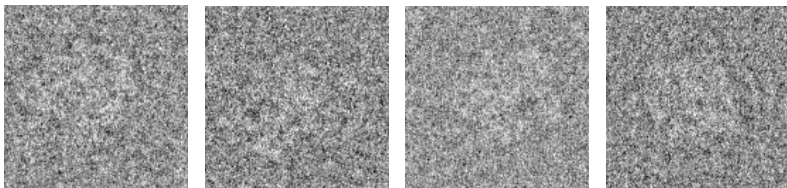


[Singer and Shkolnisky (2011)]

Fundamental challenges

① Unknown viewing directions

② Challenging SNR regime



The images of E. coli 50S ribosomal subunit were provided by Dr. Fred Sigworth, Yale Medical School.

Fundamental challenges

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② Challenging SNR regime

③ Massive datasets

2.2 Å resolution \iff 12.4 TB

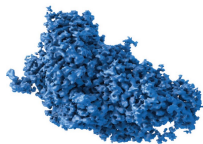


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The multireference alignment problem

Problem: Estimating a signal $x \in \mathbb{R}^L$, up to cyclic-translation, from its noisy circularly-translated copies

$$y_i = R_{r_i}x + \varepsilon_i, \quad i = 1, \dots, N, \quad \varepsilon_i \sim \mathcal{N}(0, \sigma^2 I)$$

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$\sigma = 0$



$\sigma = 0.1$



$\sigma = 1.2$



Connection with the cryo-EM problem

The problem is mainly inspired by the cryo-EM problem:

$$y_i = P(R_i \circ X) + \text{noise}, \quad X \in \mathbb{R}^3, \quad R_i \in SO(3)$$

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	Cryo-EM	MRA
Signal	3D continuous signal	1D discrete signal
Latent variables	3D rotations	cyclic translations
Linear operator	tomographic projection	no projection
Interesting regime	low SNR, large N	low SNR, large N

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Multireference alignment via alignment

Model

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Given the translations, we can estimate

$$\frac{1}{N} \sum_{i=1}^N R_{r_i}^{-1} y_i \rightarrow x$$

This is an unbiased estimator with variance σ^2/N .

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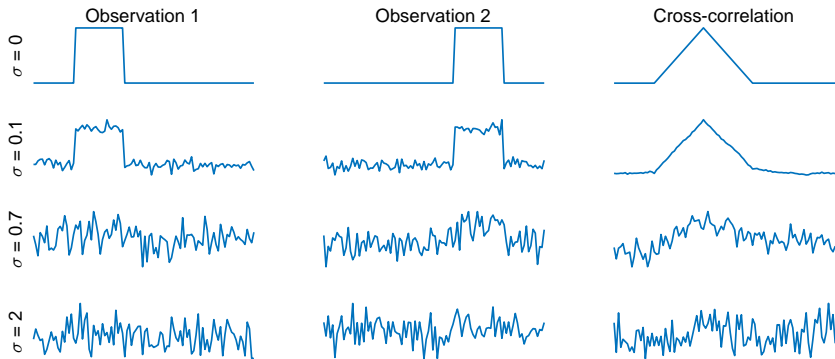
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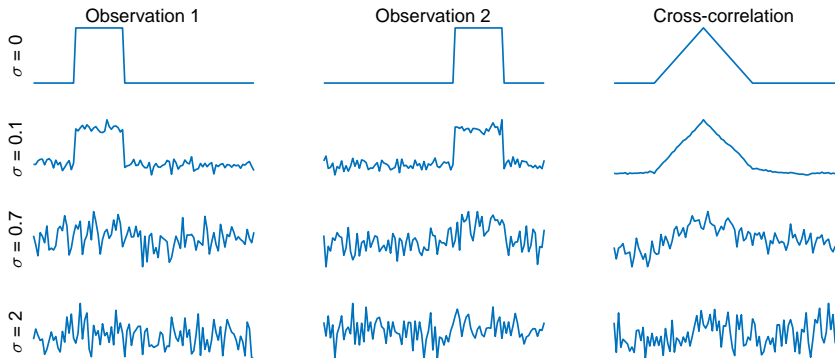
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Can we estimate the translations?

Alignment



Alignment



Alignment is impossible in the low SNR regime!

Information-theoretic limits

Can we reconstruct the signal accurately while estimating most shifts poorly?

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Can we achieve the optimal estimation rate?

Expectation–maximization

An iterative method to find maximum marginal likelihood:

$$x_{k+1} = \frac{1}{N} \sum_{i=1}^N \sum_{\ell=0}^{L-1} \omega_k^{\ell,i} R_{\ell}^{-1} y_i, \quad \omega_k^{\ell,i} \propto \exp \left(-\frac{1}{2\sigma^2} \|R_{\ell} x_k - y_i\|_2^2 \right)$$

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Can we achieve similar performance with only **one pass** over the data?

Cyclic-shift invariant features

We need the first three moments to determine a signal

$$\text{(mean)} \quad \mu_x := \hat{x}[0]/L$$

$$\text{(power spectrum)} \quad P_x[k] := \hat{x}[k]\overline{\hat{x}[k]} = |\hat{x}[k]|^2$$

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Algorithms and analysis are provided in "Bispectrum inversion with application to multireference alignment", Bendory, Boumal, Ma, Zhao and Singer.

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Stable recovery using a non-convex least-squares

$$\min_z |\mu_z - \tilde{\mu}_x|^2 + \lambda_1 \left\| P_z - \tilde{P}_x \right\|_2^2 + \lambda_2 \left\| B_z - \tilde{B}_x \right\|_F^2$$

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Multireference alignment by invariant features

The invariant features can be estimated from the data:

$$M_1 := \frac{1}{N} \sum_{i=1}^N \mu_{y_i} \rightarrow \mu_x, \quad \text{Var}(M_1) \sim \sigma^2/N,$$

$$M_2 := \frac{1}{N} \sum_{i=1}^N P_{y_i} \rightarrow P_x + \sigma^2 L \mathbf{1}, \quad \text{Var}(M_2) \sim \sigma^4/N,$$

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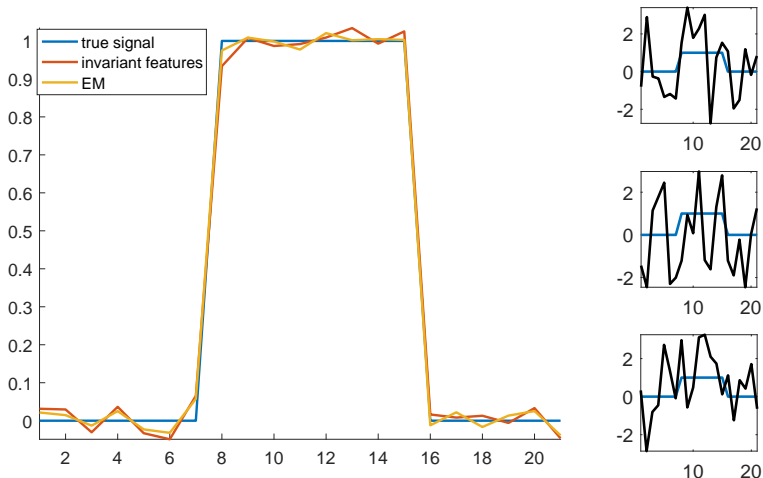
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Achieving the optimal estimation rate σ^6/N

Computational complexity linear in N (requires only one pass over the data)

Numerical example



$N = 10^5$, $\sigma = 1.5$. Running time: invariants features = 2.1 [sec], EM = 67.2 [sec]

Multireference alignment with heterogeneity

Problem: Estimating a set of signals x_1, \dots, x_K from their noisy, unlabeled, circularly-translated copies

$$y_i = R_{r_i} x_{v_i} + \varepsilon_i, \quad i = 1, \dots, N.$$

$v_i = k$ with probability w_i , $w \in \Delta^K$.

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Mixed invariant feature

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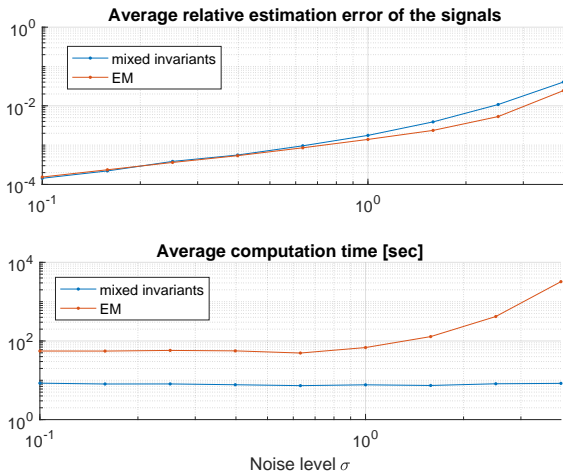
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Signal estimation by non-convex least-squares

Numerical results



$L = 50$, $K = 2$ with i.i.d. normal entries, $N = 2 \times 10^6$, 20 repetitions

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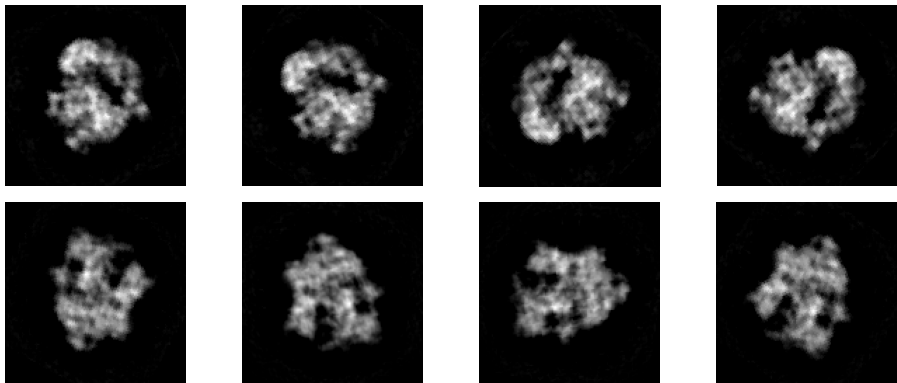
2D classification for cryo-EM

Goal: Given N noisy cryo-EM images, estimate accurately $K \ll N$ representative images

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Simplified example: Set of rotated copies of 2 images

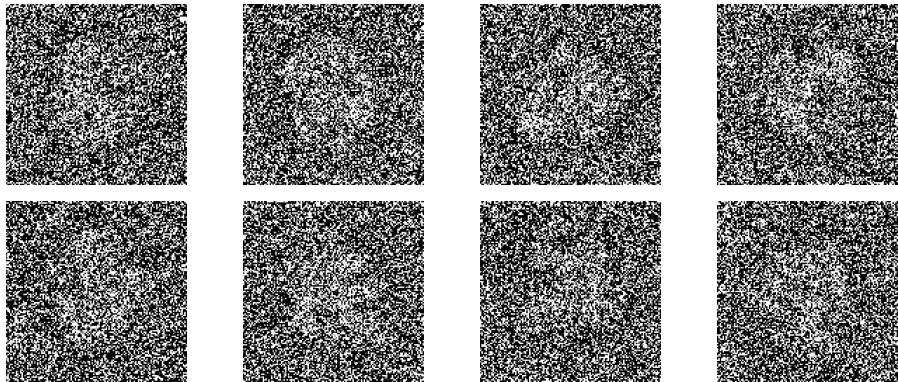


Ongoing project with Chao Ma, Nicolas Boumal and Amit Singer

2D classification for cryo-EM

Simplified example: Estimate 2 images, up to rotation, from their noisy rotated copies

Noisy rotated images: ($\text{SNR} = 1/50$):



Invariants of a steerable basis

Each image can be expanded by a **steerable** basis

$$X(r, \theta) = \sum_{k,q} A_{k,q} u^{k,q}(r, \theta),$$

satisfying

$$X(r, \theta - \alpha) = \sum_{k,q} A_{k,q} e^{-\iota k \alpha} u^{k,q}(r, \theta).$$

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The **rotationally-invariant** features are:

$$\begin{aligned} \text{(mean)} & \quad A_{0,q}, \quad \forall q \\ \text{(power spectrum)} & \quad A_{k,q_1} \overline{A_{k,q_2}}, \quad \forall k, q_1, q_2 \\ \text{(bispectrum)} & \quad A_{k_1,q_1} \overline{A_{k_2,q_2}} A_{k_2-k_1,q_3}, \quad \forall k_1, k_2, q_1, q_2, q_3 \end{aligned}$$

Algorithm for 2D classification

- 1 Expand each image in a steerable basis
- 2 Compute the rotationally-invariant features
- 3 Average over all images to estimate the **mixed** invariant features
- 4 Solve a non-convex least-squares

Numerical example

Estimated images (5000 images per class, $\text{SNR} = 1/50$):

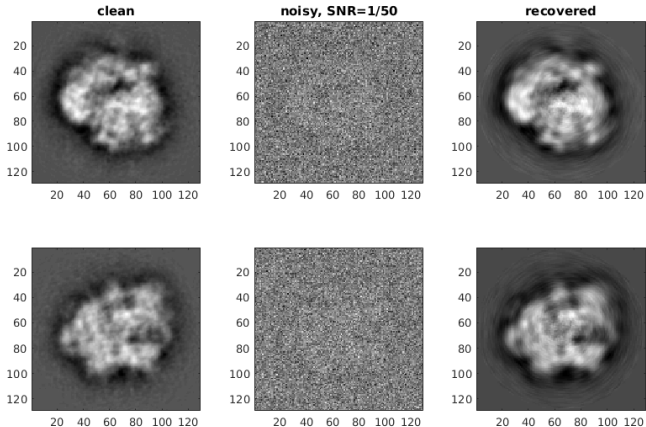


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The invariants of the cryo-EM problem

Theorem (Levin, B, Boumal, Kileel, Singer)

If the viewing directions are drawn from the uniform distribution, then the second moment of the projections and two clean images determine the molecule, up to global rotation.

- Requires one pass over the data (very fast, computational complexity linear in N)

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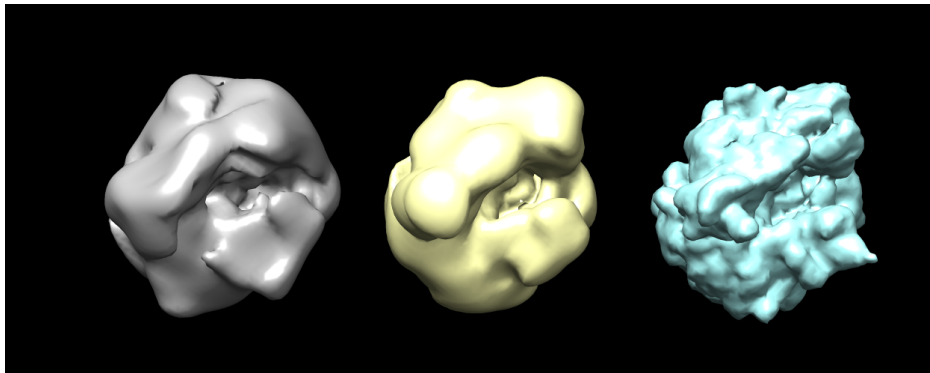
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- Requires one pass over the data (very fast, computational complexity linear in N)
- These conditions are not met in practice and therefore only low-resolution estimation is possible

3D ab-initio modeling

EMDB 5360, 50,000 projections, SNR = 1/10



estimation

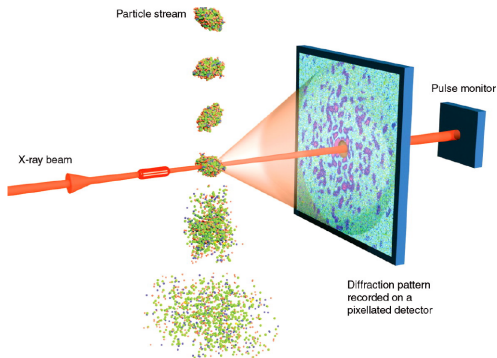
low-resolution
"ground truth"

"ground truth"

Levin, Bendory, Boumal, Kileel and Singer. "3D ab initio modeling in cryo-EM by autocorrelation analysis."

Single Particle Reconstruction using X-ray Free-Electron Laser (XFEL)

XFEL \approx cryo-EM + phase retrieval



[Gaffney and Chapman (2007)]

Main theoretical questions

Non-convex optimization

Why is it so effective for cryo-EM and multireference alignment?

Reasons for optimism: Recent developments in statistical estimation problems, such as phase retrieval, blind deconvolution, matrix completion and phase synchronization

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Information-theoretic limit

What is the sample complexity of the cryo-EM problem?

Partial results:

- Levin, Bendory, Boumal, Kileel and Singer. "3D ab initio modeling in cryo-EM by autocorrelation analysis."
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References

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Thanks for your attention!