Full-field Phase Imaging in practice and the impact of improved coherence

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Outline

- Introduction to full-field imaging in the Fresnel regime I. Overview of routines |.
- Imaging at various scales of coherence I. Lab systems II. Synchrotron III. XFEL ||.
- Challenges
- Perspectives IV.





Tomographic microscopy at synchrotron



Hierarchical imaging of biological samples



Phase contrast in free space propagation



Free space phase imaging

Linear scenarios for near-field phase retrieval:



Contrast Transfer Function (Weak scattering and absorption object)



Phase (contrast) tomography

Finding the phase of the object (real part of the refractive index)

$$I_{D}(\mathbf{x}) = \left| \text{FRT}_{k}[\text{T}_{A,\varphi}(\mathbf{x})] \right|^{2}$$
$$\phi(\mathbf{x}) = \arg\min_{\phi} \left\| \left| \text{FRT}_{k}[\text{T}_{A,\phi}(\mathbf{x})] \right|^{2} - I_{D}(\mathbf{x}) \right\|^{2}$$



Motivation to utilize phase in full-field imaging

Contrast enhancement

- Improve CNR == less noisy images == reliably image interpretation / segmentation
- reduce dose enables high resolution in-situ and in vivo imaging



• enhance acquisition speed to avoid motion blur due to sample movement





Handling coherence in full-field imaging

How do we practically handle interference of coherent beams in full-field imaging?

- Retrieve the phase and attenuation deploying CTF or TIE based algorithms -> used mainly in nanoimaging with projection microscope
- Filter the projections with the Paganin filter to obtain the projected density -> very successful at beamlines because it is robust and simple
- Use the edge enhanced image -> often suitable for visual analysis
- We simply remove fringes to get clear attenuation image (e.g. decoherer on beamlines / Bronikov aided correction for lab sources)



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Starting with low coherence Lab microtomograph

4D IMAGING LAB@LTH

SoftiMAX

Soft X-rays (in construction -> 2020)



The nanofocus beamline (users in 2017)

2016

DanMAX

Imaging & diffraction (in construction -> 2020)

maxiv.lu.se

2017 2018 2019 2020



Little coherence: lab sources

Edge enhancement vs. Paganin filtering for Versa 520

2019



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Medíum coherence

Standard ímagíng beamlíne



Mokso et al. J. Phys. D, 2013

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Image quality vs. radiation damage

2.9 µm pixel-size optics

G. Lovric et al., J. Appl. Cryst. 46 (4) 2013

901 projections



361 projections

M. Kitchen, CT dose reduction fac	ctors in the thousands using X-ra	ay phase contrast, Sci. Rep. 7 (2017)
	0.24 s	0.17 s
	CNR	
	2.3	1.5
	Entrance dose	Entrance dose
	12 Gy	9 Gy

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Lung alveoli microstructure in vivo









In vivo tomographic microscopy Lovric et al. Sci. Rep. 2017



Híghly coherent beams

Coherent ímagíng beamlíne

Nanoimaging with focused beam



LiNi type particles in new cathode material



- Hierarchical morphology of secondary particles in Li-rich cathode materials.
- Develop at longer charging time 50cycle >10cycle







Y. Yang, Y. Liu, et al., Adv. Energy Mater. (2019)



Quantification of Heterogeneous Degradation in Li-ion Batteries



Development in phase retrieval algorithms

Working towards a more robust phasing tool

Direct model in phase retreival, CTF

$$\tilde{I}_D(\mathbf{f}) = \delta(\mathbf{f}) - 2 \cos\left(\pi \lambda D |\mathbf{f}|^2\right) \tilde{B}(\mathbf{f}) + 2 \sin\left(\pi \lambda D |\mathbf{f}|^2\right) \tilde{\phi}(\mathbf{f})$$

Solution by minimization of the Euklidian distance



Needs regularisation:

$$J(\mathbf{x}) = \|\mathbf{y} - \mathbf{H}\mathbf{x}\|^2 + \lambda_{\mathrm{R}}\Delta(\mathbf{x}, \mathbf{x}_0)$$

Robust determination of the regularisation parameter under Bayesian framework

Unsupervised solution for in-line holography phase retrieval using Bayesian inference

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Fig. 6. Phase map of a red blood cell obtained through a) standard regularization and b) Bayesian inference.



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Phase retrieval with compressed sensing

Phase problem:

$$\bar{\phi} = \min_{\phi} [|\mathbf{H}\phi - I|^2 + \eta R(\phi)], \text{ where } I = \mathbf{H}\phi;$$

Contrast Transfer Function

Weak scattering <u>absorption</u> object

$$\hat{I}_z(\boldsymbol{f}_x) = 2\hat{\psi}(\boldsymbol{f}_x)\sin(\pi\lambda z|\boldsymbol{f}_x|^2) - 2\hat{B}(\boldsymbol{f}_x)\cos(\pi\lambda z|\boldsymbol{f}_x|^2)$$

 $\boldsymbol{H} = \mathcal{F}^{-1}(2\sin(\alpha|f|^2))\mathcal{F}$



b) Analytical reconstruction

(c) ADMM reconstruction



ADMM lagrangian.

$$\mathcal{L}(\phi, u, \alpha) = \frac{1}{2} \left| |H\phi - I| \right|_{2}^{2} + \lambda \sum_{i} \left| |u_{i}| \right|_{2} - \alpha^{T} (u - \nabla \phi) + \frac{\beta}{2} \left| |u - \nabla \phi| \right|_{2}^{2}$$
ADMM iteratively minimizes L

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by solving smaller problems

Optics Letters

Neglecting attenuation

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Contrast-transfer-function phase retrieval based on compressed sensing

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Sensitivity, resolution and photon flux



Total number of scattered photons

$$N_S = \Phi \sigma_S = \Phi \iint_{-\infty} |1 - t(x,y)|^2 dxdy = \Phi \pi \sigma^2 |\phi_{max}|^2$$

Coherent flux
Total scattering cross section
 $Gaussian feature width$
 $SNR_{eff}^{PM} \equiv 2\sqrt{N_S \omega} |A| \simeq 2\sqrt{N_S} \frac{4}{\sqrt{\pi}} \beta \frac{2\sigma}{FOV_{PM}}$
 $SNR_{eff}^{CDI} \equiv \sqrt{N_S} \frac{2\sqrt{\pi}\sigma}{FOV_{CDI}} \frac{2\pi}{\pi q_{max}^2} \int_0^{q_{max}} e^{-\frac{\sigma^2 q^2}{2}} q dq$
 $\approx \sqrt{N_S} \frac{2\sigma}{FOV_{CDI}}$.
Villanueva et al. Opt. Exp. 2016

Sensitivity, resolution and photon flux





Villanueva et al. Opt. Exp. 2016



Sensitivity, resolution and photon flux



$$| D[Gy] = \phi\left(\frac{\mu}{\varrho}\right) \cdot h\nu \cdot t_{\text{scan}} \qquad \begin{array}{c} \phi \dots \text{flux [photons/s]} \\ h\nu \dots \text{energy per photon} \\ t_{\text{scan}} \dots \text{total scan time} \end{array}$$



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Total entrance dose

Full-field imaging demo from MAX IV

NanoMAX experiment



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(Fully) coherent beams

European XFEL

Highest coherence

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optica

Letter

Megahertz x-ray microscopy at x-ray free-electron laser and synchrotron sources

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E = 9.3 keV,

MHz radiography

the pulse train was filled with 128 x-ray pulses with a repetition rate of 1.128 MHz. The effective pixel size of the imaging system was 3.2 μ m







Image reconstruction and analysis remains the bottleneck

Turning data into science



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MAXIN

Processing data streams for quality check and reduction



Data reduction in tomo: minimize number of scans saved per experiment. E.g. foam dynamics studies contain lot of static scans. One could correlate consecutive 3D volumes and if they match one is deleted.



Dynamic tomographic reconstruction Time domain decomposition

$$\mathcal{R}_{\alpha}f(s,z,\theta) = \iiint f(x,y,z,t)\delta(x\cos\theta + y\sin\theta - s)\delta(\theta - \alpha t)dx\,dy\,dt$$

Decomposition

$$f(x, y, z, t) \approx \sum_{j=0}^{M-1} f_j(x, y, z) \varphi_j(t),$$

where $\{f_j(x, y, z)\}_{i=0}^{M-1}$ are decomposition coefficients. Example: Fourier basis $\varphi_i(t) = e^{2\pi i t\xi_i}$. By using the linearity property of the projection operator:



2016 x 2016 x 1800 130 rotations

$$\mathcal{R}^*_{\alpha}g(x,y,z,t) = \sum_{j=0}^{M-1} \varphi_j(t) \mathcal{R}^*\left(g\hat{\varphi}_j\right)(x,y,z),$$

where $\mathcal{R}, \mathcal{R}^*$ are standard projection and back-projection operators in the static case.

Dynamic tomography: ceramic particles, ¹⁰⁰

Tomography at 2BM, APS 2560 x 2560 x 700, 12 x 900 projections

FBP reconstruction





Iterative, basis size M=24

2000

1500

x 1000

650um





y 2000 2500 1500 2000

500

1500

1000

500

2000

1500

1000 y

500



... and here is how to use the new tool

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Foam data <u>https://tomobank.readthedocs.io</u>

In this study, we investigate the rheology of liquid foams by fast synchrotron X-ray tomographic microscopy [B30]. Foams are complex cellular systems which require artifact free tomographic reconstruction for a reliable quantification of their time-dependent properties such as deformation fields of bubbles. In our example we acquire X-ray projections of the liquid foam flowing through a constriction and being rotated around the tomographic axis. The experiment was performed at the TOMCAT beamline of the Swiss Light Source using the fast acquisition setup [B28].

To load the data sets and perform reconstruction use the 📩 tomopy_rectv.py python script.

Reconstruction by Gridrec

python tomopy_rectv.py dk_MCFG_1_p_s1_.h5 --type subset --nsino 0.75 --binning 2 --frame 95

Reconstruction by the method with suppressing motion artifacts [Nikitin et. al, 2018] requires module *rectv* that can be installed from https://github.com/math-vrn/rectv_gpu. In this case, the algorithm run with option *-tv True*

```
python tomopy_rectv.py dk_MCFG_1_p_s1_.h5 --type subset --nsino 0.75 --binning 2 --tv True --frame
```

tomo_ID	00080
Image preview	
Downloads	tomo_00080
Instrument	SLS TOMCAT
Sample name	dk_MCFG_1_p_s1
Energy	16 keV
Sample-to-detector Distance	250 mm
Scan Range	180 degree



Space for the missing tools

Off-line

- Interactive visualization and interactive scripting / annotation
- Feature quantification and visualization
- Vector field visualisation





weeks

Streaming

- Detect changes in structure in real time (relate 2D to 3D)
- Interactive stream viewing (and decision taking)

seconds

38

hour<u>s</u>

Time relative to experiment start

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Complexity of the

middle

easy

analysis

difficult

Challenges

small field of view – large sample

High spatial resolution in large samples



Local tomography: a multiresolution approach

Iterative reconstruction from combined high-low resolution sinogram pairs



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Local tomography: a multiresolution approach



Perspectives

PP repair and

New science?

Fast tomographic microscopy: 3D



Real life system dynamics

The rise of the Aluminum foam: From nucleation to film rupture Acquisition speed: 208 tomo / s TOMCAT beamline, SLS

0.00 s



1 mm

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ARTICLE

https://doi.org/10.1038/s41467-019-11521-1 OPEN

Using X-ray tomoscopy to explore the dynamics of foaming metal

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Liquid Aluminum

ım

0 s



Rajmund.Mok 27.595000 s

Exploiting new geometries: multiprojection



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Perspectives: shorter exposure times – less dose

Considerations for radiation dose optimization on the new sources:

- Fringe visibility is proportional to the transversal coherence length at the sample position (= source size and distance). The contrast (CNR or SNR) comes through detecting the oscillation(s) of the Fresnel pattern.
- Phase retrieval algorithms perform better for samples with weak (no) attenuation => it makes sense to increase X-ray energy until the attenuation can be neglected (<5%)</p>
- Looking into most efficient photon detection schemes on the optics / detector side (e.g. simultaneous 2 distance acquisition)
- Make use of modern tomographic reconstruction algorithms

Deep sub-micron in vivo imaging of photoreceptor movement in drosophila 100 fps @ ID168^{mber}

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Evidence for high resolution stereovision in compound eyes



For reliable scientific interpretation In vivo measurement must be complemented by nanotomography with highest possible accuracy (cryo-nanotomo)





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Bee photoreceptors Image: Alexandra Pacureanu, ID16A

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Perspectives: new science?

3D histology (in vivo)

• Image 3D living tissue at the cellular level in 3D

Nanoscale characterization of materials with realistic dimensions

• Pores, defects

Exploiting new phasing methods: convolutional neural networks



Phase retrieval: Bayesian approach

Projection microscopy, ID16

Probability density functions are assigned to this quantities: $p(\mathbf{x})$ as the prior distribution, $p(\mathbf{y}|\mathbf{x})$ as the likelihood and $p(\mathbf{x}|\mathbf{y})$ as the the posterior distribution. Applying the Bayes rule for probabilities [24], the posterior distribution is found to be proportional to the multiple of the previous two:

$$p(\mathbf{x}|\mathbf{y}) = \frac{p(\mathbf{y}|\mathbf{x}) \ p(\mathbf{x})}{p(\mathbf{y})} \propto p(\mathbf{y}|\mathbf{x}) \ p(\mathbf{x}), \tag{30}$$

where the denominator $p(\mathbf{y}) = \int p(\mathbf{y}|\mathbf{x}) p(\mathbf{x}) dx$ is a normalizing constant called evidence, that will be ignored in the following formalism.

Let's consider a white noise distribution for ϵ , $p(\epsilon) = \mathcal{N}(\epsilon|0, v_{\epsilon}\mathbf{I})$, i.e. a Gaussian distribution with zero mean and where v_{ϵ} denotes the variance of the noise. In turn the likelihood follows a Gaussian distribution of mean equal to the model simulation $\mathbf{H}\mathbf{x}$ and variance equal to that of the noise:

$$p(\mathbf{y}|\mathbf{x}) = \mathcal{N}(\mathbf{y}|\mathbf{H}\mathbf{x}, v_{\epsilon}\mathbf{I}) \propto v_{\epsilon}^{-\frac{M}{2}} \exp\{-\frac{1}{2v_{\epsilon}}\|\mathbf{y} - \mathbf{H}\mathbf{x}\|^{2}\}.$$
 (31)

In the case of *maximum a posteriori* (MAP) estimation of the posterior distribution $p(\mathbf{x}, \boldsymbol{\theta} | \mathbf{y})$ the Bayesian approach [25] is able to infer on both the unknown quantity \mathbf{x} and the hyper-parameters of the model, $\boldsymbol{\theta}$:

$$\hat{\mathbf{x}} = \arg \max_{\mathbf{x}} p(\mathbf{x}, \boldsymbol{\theta} | \mathbf{y}) = \arg \min_{\mathbf{x}} J(\mathbf{x}), \quad \text{where } J(\mathbf{x}) = -\ln p(\mathbf{x}, \boldsymbol{\theta} | \mathbf{y}),$$
$$\hat{\boldsymbol{\theta}} = \arg \max_{\boldsymbol{\theta}} p(\mathbf{x}, \boldsymbol{\theta} | \mathbf{y}) = \arg \min_{\boldsymbol{\theta}} J(\mathbf{x})$$
(34)

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Summary

With new sources we can explore a new spatio-temporal domain to understand structure and function of materials and bio samples

Full-field tomography with focused beam will profit the most from the new source. Some hope for further dose optimization also with parallel beams

The main bottleneck remains data reconstruction and analysis (phase retrieval, quantification)

To enable new science with new sources in place we need to invest more in bringing the data handling on the same level by (i) bringing existing tools into routine use and (ii) exploiting new avenues

