

New advances in image processing : Some Inverse Problems in Biomedical Imaging

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Outline of the talk

Basic principles of Medical imaging

Inverse problems

- Basics

- Summary

Model-based approaches

- Image restoration

Data-driven approaches / End-to-end : Deep Learning

- SR for Quantitative acoustic microscopy

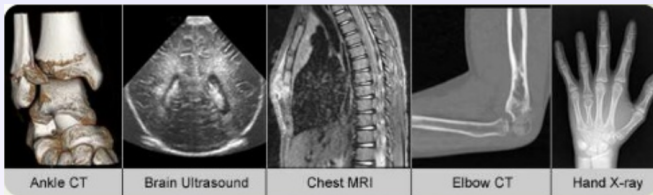
Data-driven approaches / Model-based unfolding Deep Learning

- Flow estimation and clutter rejection

- Denoising

Medical imaging modalities

- ▶ Nuclear medicine (SPECT, PET)
- ▶ Radiology techniques (X-ray radiography, CT, MRI, Ultrasound)



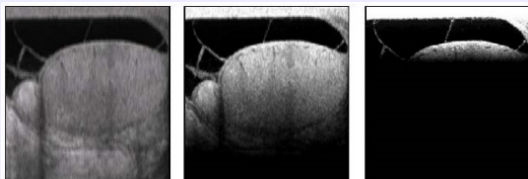
- ▶ Scanners



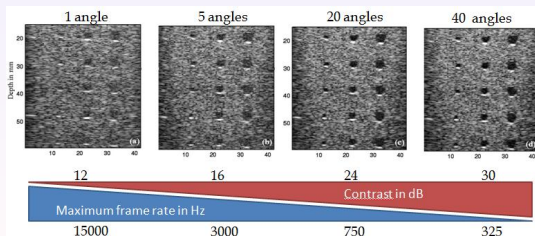
Spatial resolution in medical imaging

▶ Ultrasound imaging

- ▶ Increase the frequency - trade-off with penetration depth

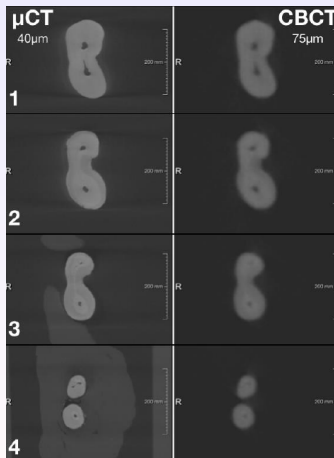


- ▶ Increase the number of fires - trade-off with frame rate



Spatial resolution in medical imaging

- ▶ Computed tomography
 - ▶ Increase the radiation dose - trade-off with patient danger and acquisition time



Summary

Spatial resolution in medical imaging

- ▶ Instrumentation definitely helps but at the cost of...
 - ▶ Field of view
 - ▶ Irradiation dose
 - ▶ Frame rate
 - ▶ Cost
- ▶ **Can we (partially) compensate for the loss of spatial resolution with post-processing methods ?**
 - ▶ Image restoration seen as an inverse problem
- ▶ **This also applies to biological images**

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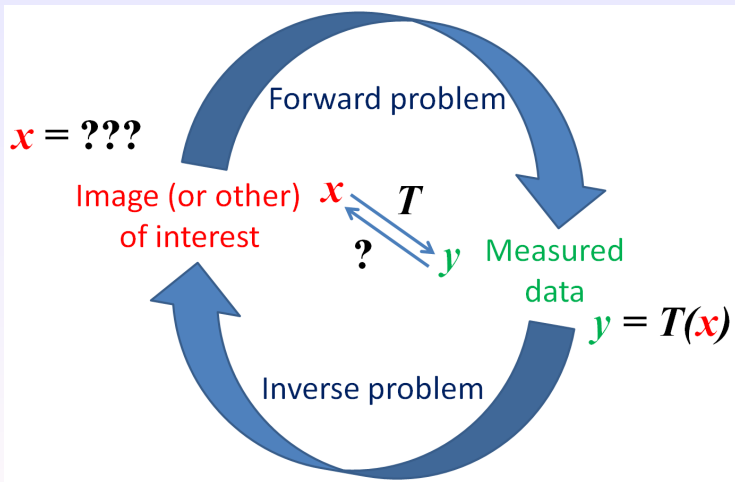
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Schematic view



Ill-posedness

$$\mathbf{y} = T(\mathbf{x}) + \mathbf{n}$$

- ▶ $\mathbf{y} \in \mathbb{C}^M$ is the observed data (image)
- ▶ $\mathbf{x} \in \mathbb{C}^N$ is the image of interest (not observed)
- ▶ $\mathbf{n} \in \mathbb{C}^M$ is the noise

T is the observation (forward) operator

- ▶ known : estimate \mathbf{x} from \mathbf{y}
- ▶ unknown : estimate \mathbf{x} and T from \mathbf{y}
 - ▶ Prior information on T (linear, parametric,...)

Inverse problems in computational medical imaging are usually ill-posed

- ▶ T is not invertible
- ▶ An infinity of solutions may exist
- ▶ A small perturbation on the data may cause an important variation on the estimate (e.g. Fourier measurements)

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General path

From the forward model to its inversion

- ▶ Establish the forward model T linking the unknown (image) to the data
 - ▶ Balance between fidelity to physics and computational tractability
- ▶ Define proper prior information about \mathbf{x} and the noise
 - ▶ Important impact on the solution's pertinence
- ▶ Formalize the inverse problem as a cost function minimization
- ▶ Stochastic simulation or numerical optimization to find the minimizer
 - ▶ Convexity of the cost function
 - ▶ Form of the forward operator T
 - ▶ Continuous and/or discrete variables
- ▶ Is the solution reliable ?

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Image restoration models

$$\mathbf{y} = \mathbf{S}\mathbf{H}\mathbf{x} + \mathbf{n}$$

- ▶ $\mathbf{x} \in \mathbb{R}^N$: image to reconstruct
- ▶ $\mathbf{y} \in \mathbb{R}^M$: observable data
- ▶ $\mathbf{H} \in \mathbb{R}^{N \times N}$: 2D convolution matrix

Deconvolution

- ▶ \mathbf{S} : identity matrix ($M = N$)

Super-resolution

- ▶ \mathbf{S} : subsampling matrix ($M = d^2 N$)

Compressed deconvolution

- ▶ \mathbf{S} : random subsampling matrix ($M \ll N$)

Denoising

- ▶ $\mathbf{S}\mathbf{H} = \mathbf{I}$: identity matrix

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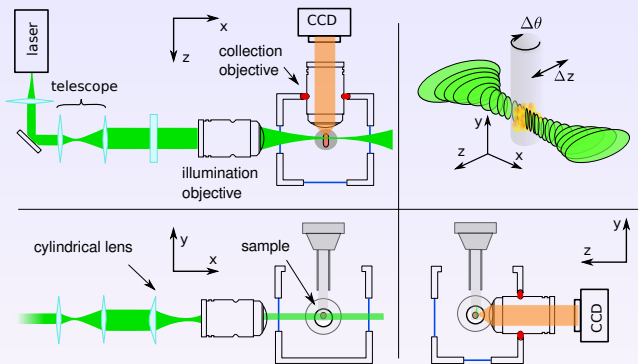
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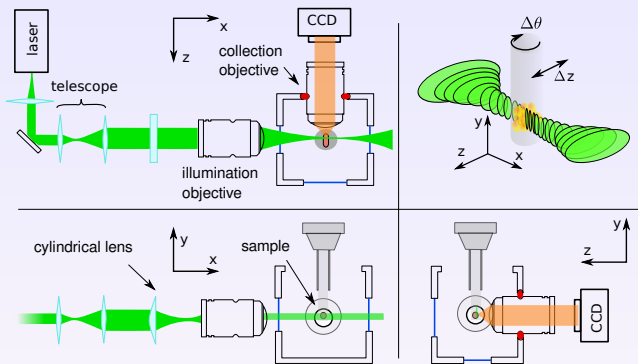
Application Light Sheet Imaging



Restoration of light sheet fluorescence microscopy 3D images :

- ▶ large 3D images (2Mb per “slice”, usually 300-500 slices)
- ▶ optic defects (resolution, blur)
- ▶ acquisition noise (low photon emission)

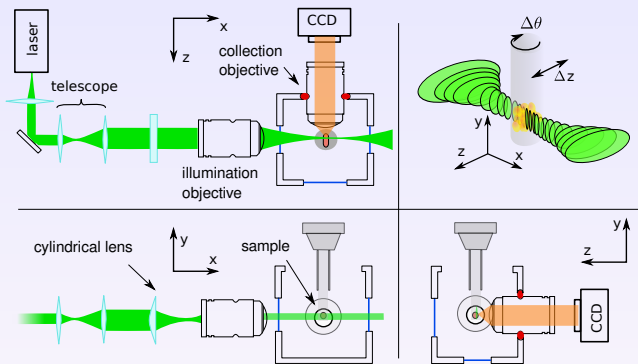
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Forward Model

Assuming :

$$\underbrace{m}_{\text{observed data}} = \underbrace{\mathcal{P}}_{\text{Poisson noise}} \left(\underbrace{H}_{\text{convolution matrix}} \underbrace{x}_{\text{unknown data}} \right)$$

With prior :

$$p(x) \propto \prod_{\text{voxels}} \exp(-\alpha TV(x))$$

Boils down to the unconstrained problem :

$$\arg \min_{x \in \mathbb{R}^n} \langle 1, Hx \rangle - \langle m, \log(Hx) \rangle + \alpha TV(x)$$

Inversion by ADMM

Let $x \in \mathbb{R}^n$, $y \in \mathbb{R}^m$, B a matrix with l rows and n columns, C a matrix with l row and m columns and $b \in \mathbb{R}^l$.

$$\arg \min_{s.t. Bx - Cy = b, x \in X, y \in Y} f_1(x) + f_2(y) \quad (1)$$

$$\begin{aligned} \mathcal{L}(x, y, \lambda) = & \langle \mathbf{1}, \mathbf{w} \rangle - \langle y, \log(\mathbf{w}) \rangle + \alpha \|\mathbf{z}\|_1 + \\ & \langle \lambda, Bx - Cy - b \rangle + \frac{\beta}{2} \|Bx - Cy - b\|_2^2, \end{aligned} \quad (2)$$

- ▶ $x_{k+1} = \arg \min_{x \in \mathbb{R}^n} \mathcal{L}(x, y_k, \lambda_k)$,
- ▶ $y_{k+1} = \arg \min_{y \in \mathbb{R}^m} \mathcal{L}(x_{k+1}, y, \lambda_k)$,
- ▶ $\lambda_{k+1} = \lambda_k + \beta(Bx + Cy - b)$,

until stopping criteria are met.

In practice

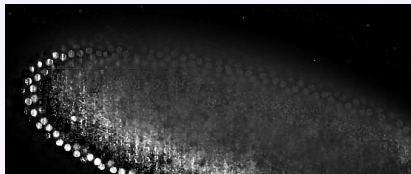
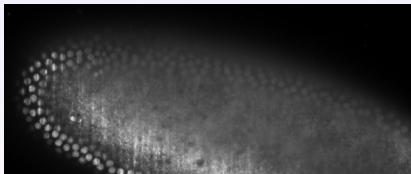
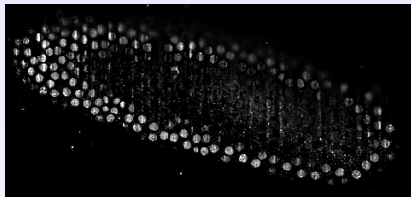
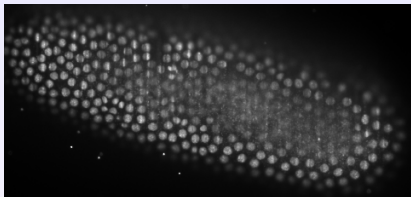
$$x_{k+1} = \mathcal{F}^{-1} \left(\frac{\mathcal{F} \left(H^T \left(w_k - \frac{\lambda_{1,k}}{\beta} \right) + \nabla^T \left(z_k - \frac{(\lambda_{2,k}, \lambda_{3,k}, \lambda_{4,k})}{\beta} \right) \right)}{\mathcal{F} (H^T H + \nabla^T \nabla)} \right). \quad (3)$$

$$w_{k+1} = \frac{q - \frac{1}{\beta}}{2} + \frac{1}{2} \sqrt{\left(\frac{1}{\beta} - q \right)^2 + \frac{4m}{\beta}}, \quad (4)$$

$$z_{k+1} = \text{soft}_{\frac{\alpha}{\beta}} \left(\nabla x_{k+1} + \frac{(\lambda_{2,k}, \lambda_{3,k}, \lambda_{4,k})}{\beta} \right). \quad (5)$$

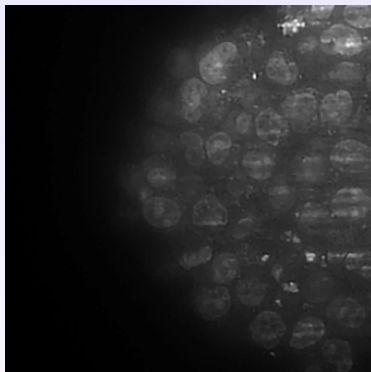
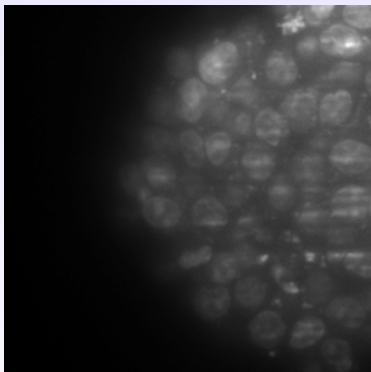
With $q = Hx_{k+1} + \frac{\lambda_{1,k}}{\beta}$ and $\text{soft}_{\gamma}(u) \equiv \frac{u}{\|u\|_1} \times \max(\|u\|_1 - \gamma, 0)$.

Drosophila



$\alpha = 0.002, \beta = 0.002, 50$ iterations.

Spheroids



$\alpha = 0.002, \beta = 0.002, 50$ iterations.

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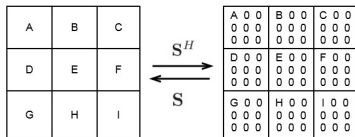
Forward model

$$\mathbf{y} = \mathbf{S}\mathbf{H}\mathbf{x} + \mathbf{n}$$

- ▶ $\mathbf{y} \in \mathbb{R}^{N_l \times 1}$: is the measured image, $N_l = m_l \times n_l$
- ▶ $\mathbf{x} \in \mathbb{R}^{N_h \times 1}$: super-resolved image to be estimated, $N_h = d^2 N_l$
- ▶ $\mathbf{n} \in \mathbb{R}^{N_l \times 1}$: Gaussian noise

Degradation operators

- ▶ $\mathbf{H} \in \mathbb{R}^{N_h \times N_h}$: 2D circulant convolution matrix (PSF of the transducer)
- ▶ $\mathbf{S} \in \mathbb{R}^{N_l \times N_h}$: subsampling operator



Proposed closed-form solution

$$\min_{\mathbf{x}} \frac{1}{2} \|\mathbf{y} - \mathbf{S}\mathbf{H}\mathbf{x}\|_2^2 + \tau \|\mathbf{A}\mathbf{x} - \mathbf{v}\|_2^2$$

- ▶ Lemma

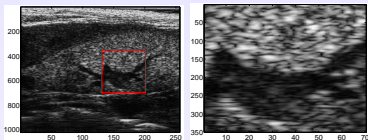
$$\mathbf{F}\mathbf{S}^H\mathbf{S}\mathbf{F}^H = \frac{1}{d^2} (\mathbf{J}_d \cdot \mathbf{I}_{m_l}) \cdot (\mathbf{J}_d \cdot \mathbf{I}_{n_l})$$

where $\mathbf{J}_d \in \mathbb{R}^{d \times d}$ is a matrix of ones, \mathbf{I}_{m_l} and \mathbf{I}_{n_l} are identity matrices and \cdot is the Kronecker product.

- ▶ Proposed solution $\mathcal{O}(N_h \log N_h)$

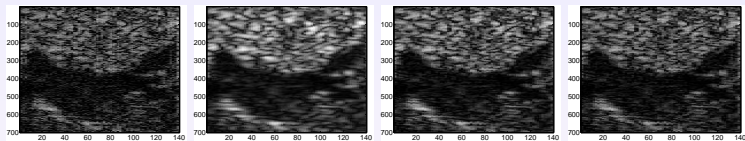
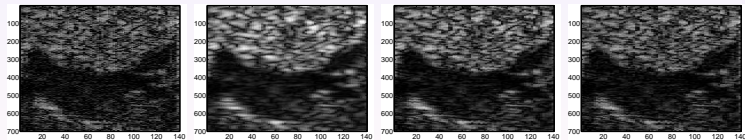
$$\hat{\mathbf{x}} = \frac{1}{2\tau} \mathbf{F}^H \Psi \mathbf{F} \mathbf{r} - \frac{1}{2\tau} \mathbf{F}^H \Psi \underline{\Lambda}^H \left(2\tau d \mathbf{I}_{N_l} + \underline{\Lambda} \Psi \underline{\Lambda}^H \right)^{-1} \underline{\Lambda} \Psi \mathbf{F} \mathbf{r}$$

where $\mathbf{r} = \mathbf{H}^H \mathbf{S}^H \mathbf{y} + 2\tau \mathbf{A}^H \mathbf{v}$, $\Psi = \mathbf{F} (\mathbf{A}^H \mathbf{A})^{-1} \mathbf{F}^H$ and $\underline{\Lambda} \in \mathbb{C}^{N_l \times N_h}$ is block diagonal

Super-resolution result on *in vivo* data

(a) Mouse kidney

(b) Observation

(c) Proposed (ℓ_1)(d) Proposed (ℓ_2)(e) Proposed ($\ell_{1.5}$)(f) Proposed ($\ell_{4/3}$)(g) Classical (ℓ_1)(h) Classical (ℓ_2)(i) Classical ($\ell_{1.5}$)(j) Classical ($\ell_{4/3}$)

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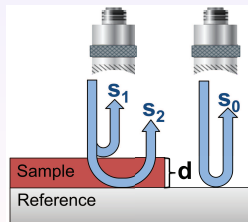
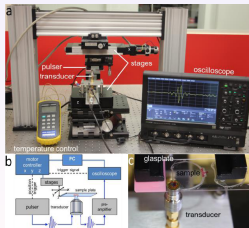
SR for Quantitative acoustic microscopy

Basics of acoustic microscopy

- ▶ Very high frequency : 250 and 500MHz
- ▶ Transmits short ultrasound pulses
- ▶ Receives the RF echo signals reflected from the sample

Sample

- ▶ Thin section of soft tissue ($12\mu m$) affixed to a microscopy slide



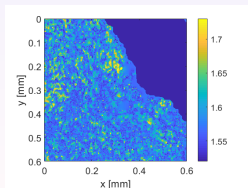
Spatial resolution in acoustic microscopy

Dependent on the central frequency

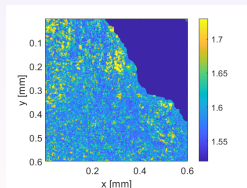
- ▶ Increasing the frequency comes with
 - ▶ Increased costs associated with the transducer and then necessary electronics
 - ▶ Experimental difficulties also arise (e.g., sensitivity to nm scale vibrations and temperature)

Example of impedance images on a section of cancerous human lymph node

- ▶ Thin section of soft tissue ($12\mu\text{m}$) affixed to a microscopy slide



250MHz



500MHz

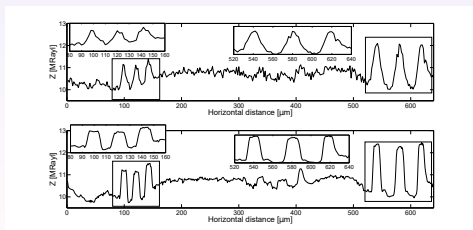
Model-based super-resolution

USAF 1951 resolution phantom

- ▶ Super-resolution factor $d = 2$



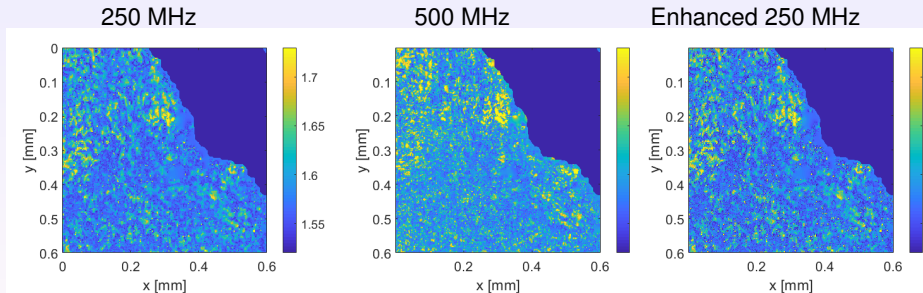
Horizontal profile



Model-based super-resolution

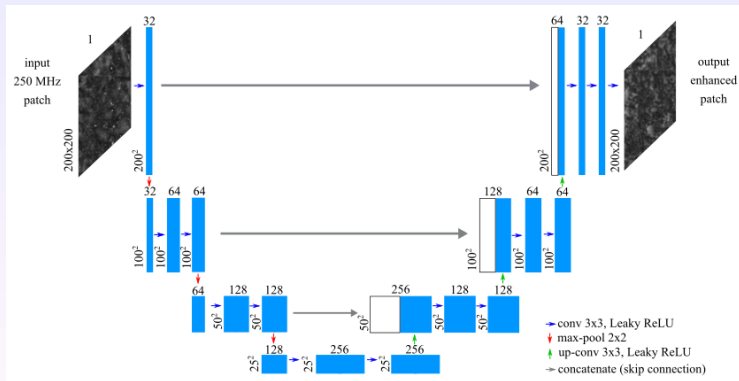
Fails on ex vivo samples

- Convolution with the PSF not sufficient to model the 250 MHz images



Data-driven super-resolution

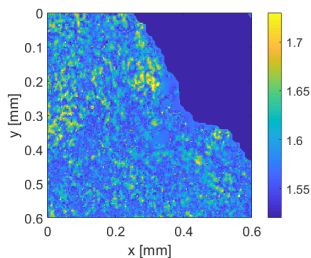
Fully convolution neural network (U-net) trained on 250 and 500 MHz images



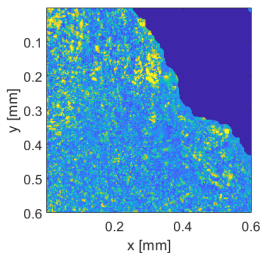
J. Mamou, T. Pellegrini, D. KouamÃ©, A. Basarab, A convolutional neural network for 250-MHz quantitative acoustic-microscopy resolution enhancement, *IEEE EMBC*, 2019.

Results

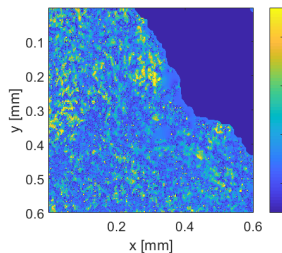
250 MHz



500 MHz



Enhanced 250 MHz



Model-based vs Data driven
approach

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Introduction

Background

- ▶ The contours of the tumor is important during surgery because it is an area from which the tumor recurs.
- ▶ Existing techniques do not allow to precisely visualize this area.



"ElastoGli" project

- ▶ To characterize in vivo the elasticity and perfusion of the cerebral gliomas and this peritumoral zone.



Photographed by Chloe, another intern at the INSERM.

Data Pre-processing

Casorati matrix

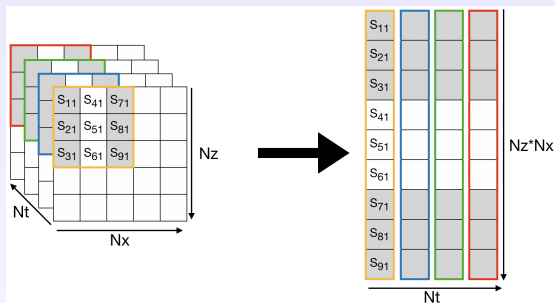


FIGURE: The construction of the Casorati matrix S from a 3D data block

$$data_{(N_z, N_x, N_t)} \implies S_{(N_z \times N_x, N_t)}$$

- ▶ Axis N_z : the propagation direction of the ultrasound ;
- ▶ Axis N_x : the opening of the ultrasound array ;
- ▶ Axis N_t : the acquisition of 2D ultrasound plans.

The inversion algorithm

Model Set-up

RPCA : A method to find the optimal sparse (Blood) and low rank (Tissue) estimation from the sequences converted to Casorati matrix, by solving a simple convex estimation problem.

$$\mathbf{S} = \mathbf{T} + \mathbf{B} + \mathbf{N}$$

- ▶ $\mathbf{S} \in \mathbb{C}^{N_z N_x \times N_t}$ is the Casorati matrix obtained from the experimental 3D Doppler data with depth N_z , probe width N_x and acquisition time N_t ;
- ▶ $\mathbf{T} \in \mathbb{C}^{N_z N_x \times N_t}$, $\mathbf{B} \in \mathbb{C}^{N_z N_x \times N_t}$ and $\mathbf{N} \in \mathbb{C}^{N_z N_x \times N_t}$ the tissue, blood and noise respectively.

Inversion

$$(\hat{\mathbf{B}}, \hat{\mathbf{T}}) = \arg \min_{\mathbf{B}, \mathbf{T}} \|\mathbf{S} - \mathbf{H}_1 \mathbf{B} - \mathbf{H}_2 \mathbf{b} \mathbf{m} \mathbf{T}\|_F^2 + \lambda \|\mathbf{B}\|_1 + \|\mathbf{T}\|_*$$



ISTA algorithm (could also be done by ADMM)

Iterations

- ▶ two main steps

$$\begin{aligned}\mathbf{L}^{k+1} &= \text{SVT}_{\lambda_1/L_f} \left\{ \left(\mathbf{I} - \frac{1}{L_f} \mathbf{H}_1^H \mathbf{H}_1 \right) \mathbf{L}^k - \mathbf{H}_1^H \mathbf{H}_2 \mathbf{S}^k + \mathbf{H}_1^H \mathbf{D} \right\} \\ \mathbf{S}^{k+1} &= \mathcal{T}_{\lambda_2/L_f} \left\{ \left(\mathbf{I} - \frac{1}{L_f} \mathbf{H}_2^H \mathbf{H}_2 \right) \mathbf{S}^k - \mathbf{H}_2^H \mathbf{H}_1 \mathbf{L}^k + \mathbf{H}_2^H \mathbf{D} \right\}\end{aligned}$$

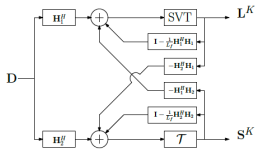
ISTA algorithm (could also be done by ADMM)

Iterations

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$$\mathbf{L}^{k+1} = \text{SVT}_{\lambda_1/L_f} \left\{ \left(\mathbf{I} - \frac{1}{L_f} \mathbf{H}_1^H \mathbf{H}_1 \right) \mathbf{L}^k - \mathbf{H}_1^H \mathbf{H}_2 \mathbf{S}^k + \mathbf{H}_1^H \mathbf{D} \right\}$$

$$\mathbf{S}^{k+1} = \mathcal{T}_{\lambda_2/L_f} \left\{ \left(\mathbf{I} - \frac{1}{L_f} \mathbf{H}_2^H \mathbf{H}_2 \right) \mathbf{S}^k - \mathbf{H}_2^H \mathbf{H}_1 \mathbf{L}^k + \mathbf{H}_2^H \mathbf{D} \right\}$$



(a) Iterative algorithm for sparse and low-rank separation.

Deep Unfolded version

Convolution come in the iterations

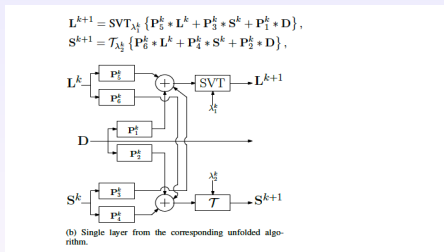
► Thus,

$$\begin{aligned} \mathbf{L}^{k+1} &= \text{SVT}_{\lambda_1^k} \{ \mathbf{P}_5^k * \mathbf{L}^k + \mathbf{P}_3^k * \mathbf{S}^k + \mathbf{P}_1^k * \mathbf{D} \}, \\ \mathbf{S}^{k+1} &= \mathcal{T}_{\lambda_2^k} \{ \mathbf{P}_6^k * \mathbf{L}^k + \mathbf{P}_4^k * \mathbf{S}^k + \mathbf{P}_2^k * \mathbf{D} \}, \end{aligned}$$

Deep Unfolded version

Convolution come in the iterations

► Thus,

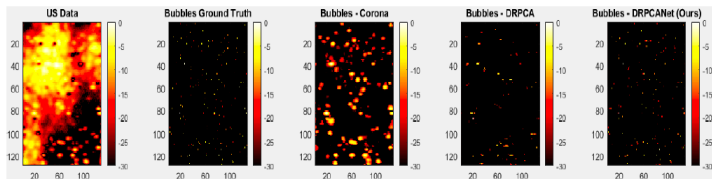


Oren Solomon, Regev Cohen, Yi Zhang, Yi Yang, He Qiong, Jianwen Luo, Ruud J.G. van Sloun, and Yonina C. Eldar, Deep Unfolded Robust PCA with Application to Clutter Suppression in Ultrasound, IEEE Trans. Med. Imaging, 2020

Some results

Classical vs unfolded

► Illustrations,



	Corona	DRPCA	Deconvolved Corona	Unfolded DRPCA (Ours)
PSNR (dB)	24.41	27.41	17.85	36.79
SSIM3	0.13	0.49	0.14	0.95

► Some more illustrations

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Hamiltonian operator

- ▶ Construct an adaptive basis using the solutions of Schrödinger equation.
- ▶ The Schroedinger equation :

$$-\frac{\hbar^2}{2m} \nabla^2 \psi = -V(a)\psi + E\psi$$

- ▶ Can be rewritten as an eigenvalue problem :

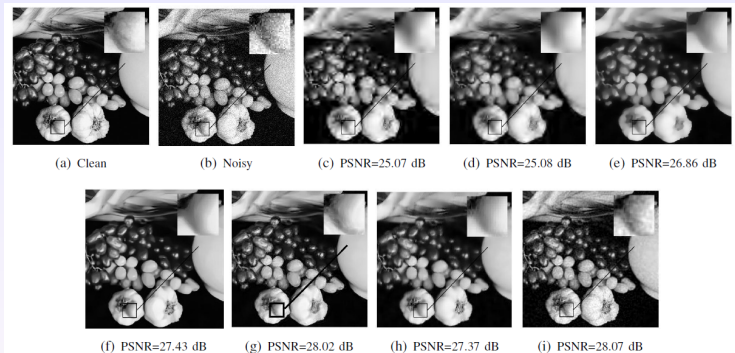
$$\mathbf{H}_{QAB}\psi = E\psi,$$

where $\mathbf{H}_{QAB} = -\frac{\hbar^2}{2m} \nabla^2 + V$ is the Hamiltonian operator.

- ▶ **Main idea** : replace $V(a)$, the potential of the system, by an image pixels' values.
- ▶ The Hamiltonian operator associated to an image

$$\mathbf{H}_{QAB}[i, j] = \begin{cases} \mathbf{x}[j] + 4\frac{\hbar^2}{2m} & \text{for } i = j, \\ -\frac{\hbar^2}{2m} & \text{for } i = j \pm 1, \\ -\frac{\hbar^2}{2m} & \text{for } i = j \pm n, \\ 0 & \text{otherwise,} \end{cases}$$

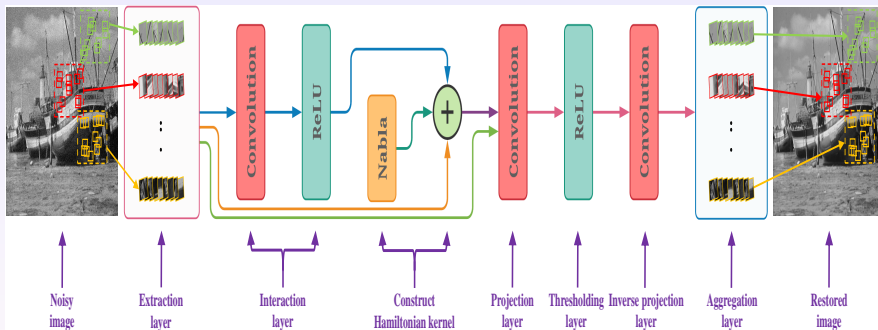
Denoising results



- ▶ (a) Clean Fruits image, (b) Image corrupted with Gaussian noise corresponding to a SNR of 15 dB.
- ▶ Denoising results obtained using : (c) wavelet hard thresholding, (d) wavelet soft thresholding, (e) total variation regularization, (f) graph signal processing, (g) non-local means, (h) dictionary learning and (i) proposed method.

TDeep unfolding

The architecture of the proposed Deep-Quantum-based image denoising



Some results

Classical vs unfolded

New advances in image processing : Some Inverse Problems in Biomedical Imaging

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