

# Dynamic MRI image reconstruction using adaptive regularization methods



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## Ill-posed problems and regularization

- Classical “hand crafted” regularizers

- Data-driven (adaptive / learned) regularizers

- Data-driven regularized MRI via dictionary learning

- Extension: learning low-rank atoms

  - DictioNary with lOw-ranK AToms (DINO-KAT)

## Dynamic MR imaging

- DINO-KAT for dynamic MR

## Summary

## Backup

- DINO-KAT convergence guarantees

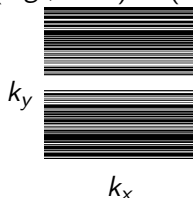
- Online method

$$\mathbf{y} = \mathbf{A}\mathbf{x} + \boldsymbol{\varepsilon}$$

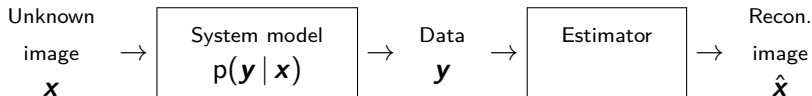
$\mathbf{y}$  : measurements       $\boldsymbol{\varepsilon}$  : noise

$\mathbf{x}$  : unknown image       $\mathbf{A}$  : system matrix (typically wide)

- ▶ compressed sensing (e.g., MRI)      ( $\mathbf{A}$  “random” rows of DFT)



- ▶ deblurring (restoration)      ( $\mathbf{A}$  Toeplitz)
- ▶ in-painting      ( $\mathbf{A}$  subset of rows of  $\mathbf{I}$ )
- ▶ denoising (not ill posed)      ( $\mathbf{A} = \mathbf{I}$ )



If we have a prior  $p(\mathbf{x})$ , then the MAP estimate is:

$$\hat{\mathbf{x}} = \arg \max_{\mathbf{x}} p(\mathbf{x} | \mathbf{y}) = \arg \max_{\mathbf{x}} \log p(\mathbf{y} | \mathbf{x}) + \log p(\mathbf{x}).$$

For gaussian measurement errors and a linear forward model:

$$-\log p(\mathbf{y} | \mathbf{x}) \equiv \frac{1}{2} \|\mathbf{y} - \mathbf{A}\mathbf{x}\|_{\mathbf{W}}^2$$

where  $\|\mathbf{y}\|_{\mathbf{W}}^2 = \mathbf{y}' \mathbf{W} \mathbf{y}$

and  $\mathbf{W}^{-1} = \text{Cov}\{\mathbf{y} | \mathbf{x}\}$  is known  
( $\mathbf{A}$  from physics,  $\mathbf{W}$  from statistics)



- ▶ If all images  $\mathbf{x}$  are “plausible” (have non-zero probability) then

$$p(\mathbf{x}) \propto e^{-\mathbf{R}(\mathbf{x})} \implies -\log p(\mathbf{x}) \equiv \mathbf{R}(\mathbf{x})$$

(from fantasy / imagination / wishful thinking / data)

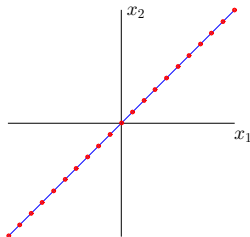
- ▶ MAP  $\equiv$  regularized weighted least-squares (WLS) estimation:

$$\begin{aligned}\hat{\mathbf{x}} &= \arg \max_{\mathbf{x}} \log p(\mathbf{y} | \mathbf{x}) + \log p(\mathbf{x}) \\ &= \arg \min_{\mathbf{x}} \frac{1}{2} \|\mathbf{y} - \mathbf{A}\mathbf{x}\|_{\mathbf{W}}^2 + \mathbf{R}(\mathbf{x})\end{aligned}$$

- ▶ A regularizer  $\mathbf{R}(\mathbf{x})$ , aka log prior, is essential for high-quality solutions to ill-conditioned / ill-posed inverse problems.
- ▶ Why ill-posed? Often high ambitions...

# Subspace model: Alternative to regularization

Assuming  $\mathbf{x}$  lies in a sufficiently low-dimensional subspace could make an inverse problem well conditioned.



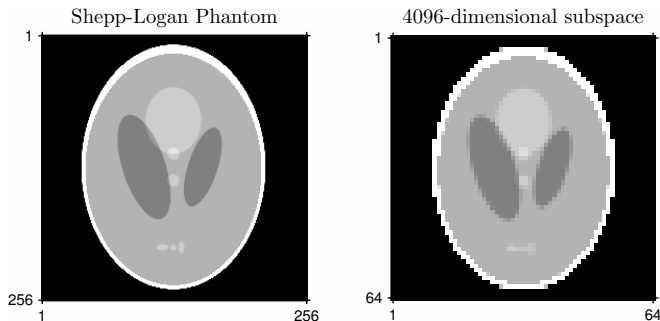
Assume  $\mathbf{x} = \mathbf{D}\mathbf{z}$  where  $\mathbf{D} = \begin{bmatrix} 1 \\ 1 \end{bmatrix}$  and  $\mathbf{z} \in \mathbb{R}^1$

( $\mathbf{z}$  has only one nonzero element so very sparse!?)

Estimate coefficient(s):  $\hat{\mathbf{z}} = \arg \min_{\mathbf{z}} \|\mathbf{y} - \mathbf{A}\mathbf{D}\mathbf{z}\|_2^2$ , then  $\hat{\mathbf{x}} = \mathbf{D}\hat{\mathbf{z}}$ ,  
where usually  $\text{cond}(\mathbf{D}'\mathbf{A}'\mathbf{A}\mathbf{D}) \ll \text{cond}(\mathbf{A}'\mathbf{A})$ .

# Why not use subspace models?

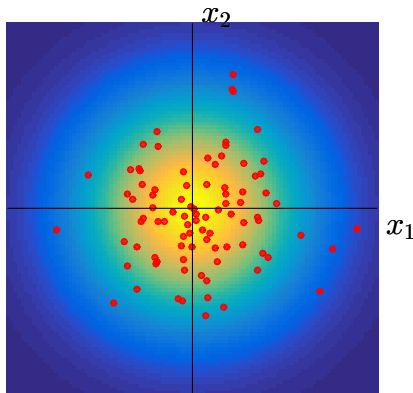
Candès and Romberg (2005) [1] used 22 (noiseless) CT projection views (*i.e.*, 22 pseudo-radial lines in MRI), each with 256 samples.  
 $\implies 22 \cdot 256 = 5632$  measured values,  
vs  $256^2 = 65536$  unknown pixels



Subspace representation (using pixel basis) is undesirably coarse.

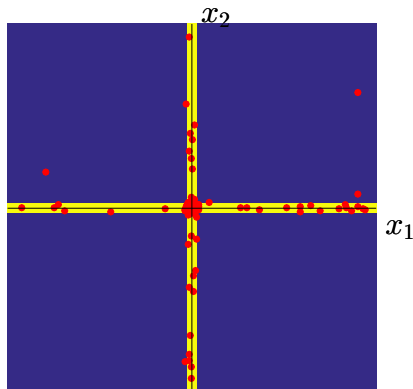
- ▶ Tikhonov regularization (IID gaussian prior)
- ▶ Roughness penalty (Basic MRF prior)
- ▶ Sparsity in ambient space
- ▶ Edge-preserving regularization
- ▶ Total-variation (TV) regularization
- ▶ Black-box denoiser like NLM

$$R(\mathbf{x}) = \beta \|\mathbf{x}\|_2^2$$



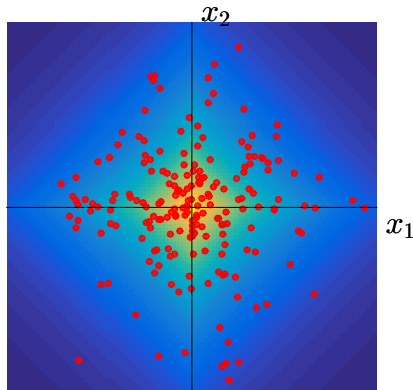
- ▶ Colors show equivalent (normalized) prior  $p(\mathbf{x}) / p(\mathbf{0}) = e^{-R(\mathbf{x})}$
- ▶ Equivalent to IID gaussian prior on  $\mathbf{x}$
- ▶ Makes any ill-conditioned / ill-posed problem well conditioned
- ▶ Ignores correlations between pixels

$$R(\mathbf{x}) = \beta \|\mathbf{x}\|_0 = \beta \sum_j \mathbb{I}_{\{x_j \neq 0\}}$$

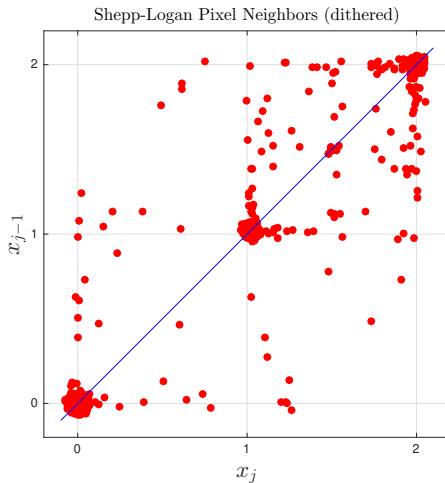
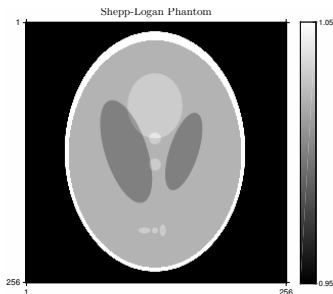


- ▶ Approximate Bayesian interpretation
- ▶ Non-convex
- ▶ IID  $\implies$  also ignores correlations

$$R(\mathbf{x}) = \beta \|\mathbf{x}\|_1 = \beta \sum_j |x_j|$$



- ▶ Equivalent to IID Laplacian prior on  $\mathbf{x}$
- ▶ Also ignores correlations



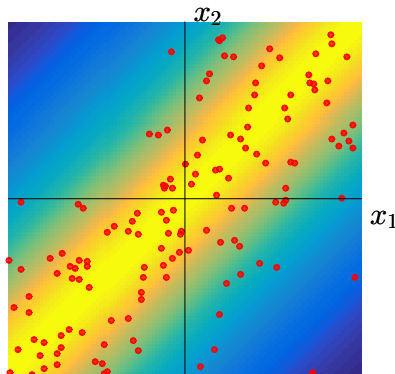
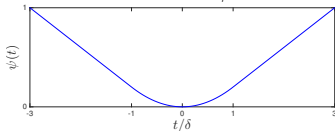
Caution: Shepp-Logan phantom [2] was designed for testing non-Bayesian methods, not for designing signal models. Q: What causes the spread??



Neighboring pixels tend to have similar values except near edges:

$$R(\mathbf{x}) = \beta \sum_j \psi(x_j - x_{j-1})$$

Potential function  $\psi$ :



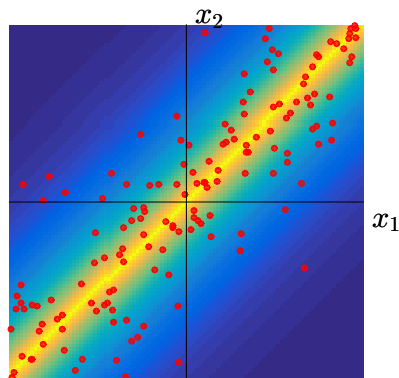
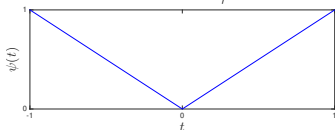
- Equivalent to improper prior (agnostic to DC value)
- Accounts for spatial correlations, but only very locally
- Used clinically now for low-dose X-ray CT image reconstruction

# Total-variation (TV) regularization

Neighboring pixels tend to have similar values except near edges (“gradient sparsity”):

$$\begin{aligned} R(\mathbf{x}) &= \beta \text{TV}(\mathbf{x}) = \beta \|\mathbf{C}\mathbf{x}\|_1 \\ &= \beta \sum_j |x_j - x_{j-1}| \end{aligned}$$

Potential function  $\psi$ :



- ▶ Equivalent to improper prior (agnostic to DC value)
- ▶ Accounts for correlations, but only very locally
- ▶ Well-suited to piece-wise constant Shepp-Logan phantom!
- ▶ Used in many academic publications...

Noisy image  $\rightarrow$  Denoiser  $\rightarrow$  Denoised image

- ▶ Example: Non-local means (NLM)
- ▶ Corresponding regularizer [3–5]:

$$\mathbf{R}(\mathbf{x}) = \beta \frac{1}{2} \|\mathbf{x} - \text{NLM}(\mathbf{x})\|_2^2$$

- ▶ Encourages self-consistency with denoised version of image
- ▶ No evident Bayesian interpretation
- ▶ Variable splitting can facilitate minimization [6].

- ▶ Transforms: wavelets, curvelets, ...
- ▶ Markov random field models
- ▶ Graphical models
- ▶ ...

## Ill-posed problems and regularization

Classical “hand crafted” regularizers

**Data-driven (adaptive / learned) regularizers**

Data-driven regularized MRI via dictionary learning

Extension: learning low-rank atoms

DictioNary with lOw-ranK AToms (DINO-KAT)

## Dynamic MR imaging

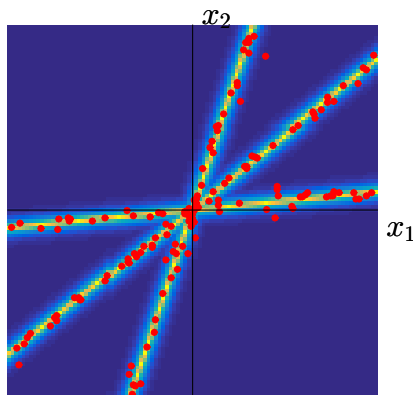
DINO-KAT for dynamic MR

## Summary

## Backup

DINO-KAT convergence guarantees

Online method



- ▶ Dimensionality reduction?
- ▶ *cf.* classification / clustering motivation [7]
- ▶ (Extension to union of “flats” (linear varieties) is possible [8].)

Given (?) collection of  $K$  subspace bases  $\mathbf{D}_1, \dots, \mathbf{D}_K$   
(dictionaries with full column rank  $\implies$  tall).

Assume  $\mathbf{x} \approx \mathbf{D}_k \mathbf{z}_k$  for some  $k$  and some (non-sparse) coefficients  $\mathbf{z}_k$ .

Natural regularizer for this model is:

$$\begin{aligned} \mathbf{R}(\mathbf{x}) &= \underbrace{\min_k}_{\text{"classification"}} \underbrace{\min_{\mathbf{z}_k} \beta \frac{1}{2} \|\mathbf{x} - \mathbf{D}_k \mathbf{z}_k\|_2^2}_{\text{regression}} \\ &= \min_k \beta \frac{1}{2} \left\| \mathbf{x} - \mathbf{D}_k \mathbf{D}_k^+ \mathbf{x} \right\|_2^2. \end{aligned}$$

- ▶  $\mathbf{R}(\mathbf{x}) = 0$  if  $\mathbf{x}$  lies in the span of any of the dictionaries  $\{\mathbf{D}_k\}$ .
- ▶ Otherwise, distance to nearest subspace (discourage, not constrain).
- ▶ Non-convex (highly?) (cf. preceding picture) due to min
- ▶ Apply to image patches to be practical.
- ▶ Equivalent Bayesian interpretation? (Not a mixture model here.)
- ▶ Given? Learned from training **data**.

Assume  $\mathbf{x} \approx \mathbf{D}\mathbf{z}$  where

- ▶  $\mathbf{D}$  is a dictionary (often over-complete  $\implies$  wide)
- ▶  $\mathbf{z}$  is a *sparse* coefficient vector (subset of columns of  $\mathbf{D}$ ).

Corresponding regularizers:

$$\mathbf{R}(\mathbf{x}) = \min_{\mathbf{z}: \|\mathbf{z}\|_p \leq s} \beta \frac{1}{2} \|\mathbf{x} - \mathbf{D}\mathbf{z}\|_2^2, \quad \text{or:}$$

$$\mathbf{R}(\mathbf{x}) = \min_{\mathbf{z}} \left( \beta_1 \frac{1}{2} \|\mathbf{x} - \mathbf{D}\mathbf{z}\|_2^2 + \beta_2 \|\mathbf{z}\|_p \right).$$

- ▶ Convex in  $\mathbf{z}$  (for given  $\mathbf{x}$ ) if  $p \geq 1$  and  $\mathbf{D}$  given.
- ▶  $\mathbf{R}(\mathbf{x})$  typically non-convex in  $\mathbf{x}$ , due to  $\|\cdot\|_p$ .
- ▶ Could be equivalent to a union-of-subspaces regularizer if  $\mathbf{D} = [\mathbf{D}_1 \ \dots \ \mathbf{D}_K]$  and if we constrain coefficient vector  $\mathbf{z}$  in a non-standard way.



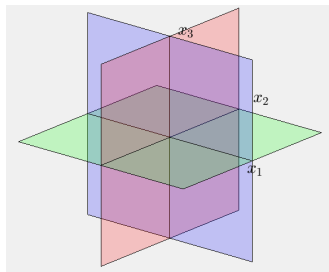
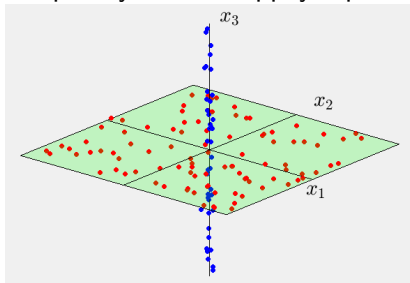
# Union-of-subspaces vs sparse-coding-with-dictionary

Consider union-of-subspaces model with  $\mathbf{D}_1 = \begin{bmatrix} 1 & 0 \\ 0 & 1 \\ 0 & 0 \end{bmatrix}$ ,  $\mathbf{D}_2 = \begin{bmatrix} 0 \\ 0 \\ 1 \end{bmatrix}$ .

So  $\mathbf{D}_1$  spans x-y plane and  $\mathbf{D}_2$  spans z-axis.

A dictionary model with  $\mathbf{D} = [\mathbf{D}_1 \ \mathbf{D}_2] = \begin{bmatrix} 1 & 0 & 0 \\ 0 & 1 & 0 \\ 0 & 0 & 1 \end{bmatrix}$

and sparsity  $s = 2$ , happily represents all three cardinal planes.



Thus dictionary model seems “less constrained” than union-of-subspaces model.

(Still, focus on sparse dictionary representation hereafter.)

# New dictionary learning method (SOUP-DIL)

Joint work with Sai Ravishankar and Raj Nadakuditi [9–11]

- In practice, must **learn  $D$  from data**, say  $X$
- Write sparse representation as Sum of Outer Products (SOUP):

$$X \approx DZ = DC' = \sum_{j=1}^J d_j c_j'$$

where  $Z' = C = [c_1 \dots c_J] \in \mathbb{R}^{N \times J}$  (coefficients for each atom)

- Replace individual atom sparsity constraint  $\|z_n\|_0 \leq s$  of K-SVD with aggregate sparsity regularizer:  $\|Z\|_0 = \|C\|_0$ .
  - ▶ Natural for Dictionary Learning (DIL) from training data.
  - ▶ Unnatural for image compression using sparse coding.

SOUP-DIL  $\ell_0$  formulation:

$$D^* = \arg \min_{D \in \mathbb{R}^{d \times J}} \min_{C \in \mathbb{R}^{N \times J}} \|X - DC'\|_F^2 + \lambda^2 \|C\|_0 \quad \text{s.t.} \quad \begin{aligned} \|d_j\|_2 &= 1 \quad \forall j \\ \|c_j\|_\infty &\leq L \quad \forall j \end{aligned}$$

SOUP-DIL formulation [9–11]:

$$\mathbf{D}^* = \arg \min_{\mathbf{D} \in \mathbb{R}^{d \times J}} \min_{\mathbf{C} \in \mathbb{R}^{N \times J}} \|\mathbf{X} - \mathbf{D}\mathbf{C}'\|_F^2 + \lambda^2 \|\mathbf{C}\|_0 \quad \text{s.t.} \quad \begin{aligned} \|\mathbf{d}_j\|_2 &= 1 \quad \forall j \\ \|\mathbf{c}_j\|_\infty &\leq L \quad \forall j \end{aligned}$$

- ▶ Block coordinate descent (BCD) algorithm
  - Sparse coding step for  $\mathbf{C}$
  - Dictionary update step for  $\mathbf{D}$
- ▶ Very simple update rules (low compute cost)
- ▶ Monotone descent of cost function  $\Psi(\mathbf{D}, \mathbf{C})$
- ▶ Convergence theorem: for any given initialization  $(\mathbf{D}^0, \mathbf{C}^0)$ , all accumulation points of sequence  $(\mathbf{D}, \mathbf{C})$ 
  - are critical points of cost  $\Psi$  and
  - are equivalent (reach same cost function value  $\Psi^*$ ).
  - Furthermore:  $\left\{ \|\mathbf{D}^{(k)} - \mathbf{D}^{(k-1)}\| \right\} \rightarrow 0$ . Same for  $\left\{ \mathbf{C}^{(k)} \right\}$ .

$$\mathbf{D}^* = \arg \min_{\mathbf{D} \in \mathbb{R}^{d \times J}} \min_{\mathbf{C} \in \mathbb{R}^{N \times J}} \|\mathbf{X} - \mathbf{D}\mathbf{C}'\|_F^2 + \lambda^2 \|\mathbf{C}\|_0 \quad \text{s.t.} \quad \begin{aligned} \|\mathbf{d}_j\|_2 &= 1 \quad \forall j \\ \|\mathbf{c}_j\|_\infty &\leq L \quad \forall j \end{aligned}$$

Alternate: update one column  $\mathbf{c}_j$  of  $\mathbf{C}$  then one column  $\mathbf{d}_j$  of  $\mathbf{D}$ .

- Sparse coding step: update  $\mathbf{c}_j$  with residual  $\mathbf{E}_j \triangleq \mathbf{X} - \sum_{k \neq j} \mathbf{d}_k \mathbf{c}'_k$ :

$$\min_{\mathbf{c}_j} \|\mathbf{E}_j - \mathbf{d}_j \mathbf{c}'_j\|_F^2 + \lambda^2 \|\mathbf{c}_j\|_0 \quad \text{s.t.} \quad \|\mathbf{c}_j\|_\infty \leq L.$$

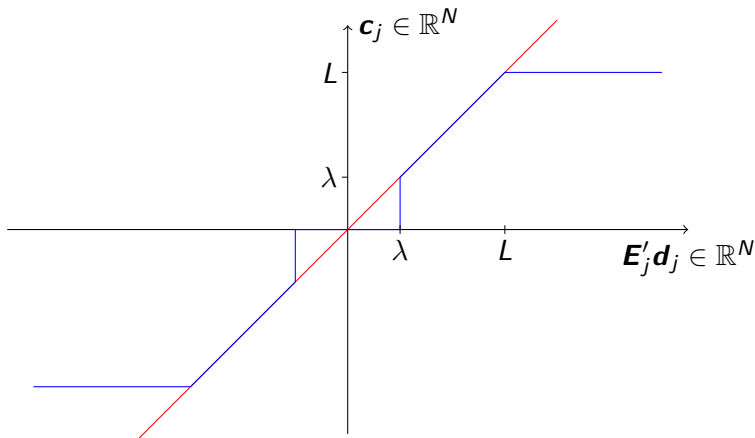
Truncated (via  $L$ ) hard thresholding of  $\mathbf{E}'_j \mathbf{d}_j$  with threshold  $\lambda$ .

- Dictionary atom step: update  $\mathbf{d}_j$

$$\min_{\mathbf{d}_j} \|\mathbf{E}_j - \mathbf{d}_j \mathbf{c}'_j\|_F^2 \quad \text{s.t.} \quad \|\mathbf{d}_j\|_2 = 1.$$

Constrained least-squares solution:  $\mathbf{d}_j = (\mathbf{E}_j \mathbf{c}_j) / \|\mathbf{E}_j \mathbf{c}_j\|_2$ .

# Truncated hard thresholding for SOUP-DIL



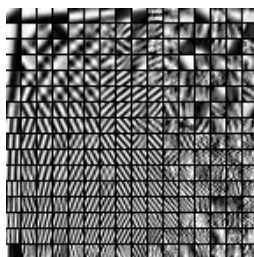
(Acts element-wise.) (In practice take  $L = \infty$ .)

(Algorithm also provides a simple sparse coding method.)

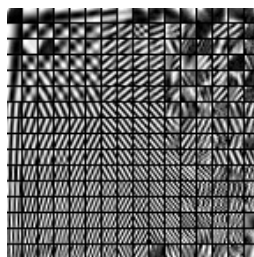
# Example: dictionary learning for Barbara



Barbara



K-SVD  $D$



SOUP-DIL  $D$

Denoising PSNR (dB) from [9]

$\sigma$	Noisy	O-DCT	K-SVD	SOUP-DIL
20	22.13	29.95	<b>30.83</b>	30.79
25	20.17	28.68	29.63	<b>29.64</b>
30	18.59	27.62	28.54	<b>28.63</b>
100	8.11	21.87	21.87	<b>21.97</b>

SOUP-DIL faster than K-SVD

- ▶ Large image  $\mathbf{x} \implies$  extract  $M$  patches  $\mathbf{X} = [\mathbf{P}_1\mathbf{x} \dots \mathbf{P}_M\mathbf{x}]$ .
- ▶ Assume patch  $\mathbf{x}_m = \mathbf{P}_m\mathbf{x} \approx \mathbf{D}\mathbf{z}_m$  has (aggregate) sparse representation in dictionary  $\mathbf{D} \in \mathbb{R}^{d \times J}$  where  $d$  is patch size.
- ▶ Two variations:
  - Use dictionary  $\mathbf{D}$  from training data:

$$\mathbf{R}(\mathbf{x}) = \mathbf{R}(\mathbf{X}) = \min_{\mathbf{C} \in \mathcal{C}} \|\mathbf{X} - \mathbf{D}\mathbf{C}'\|_F^2 + \lambda^2 \|\mathbf{C}\|_0$$

- Learn  $\mathbf{D}$  while reconstructing (blind / adaptive)

$$\mathbf{R}(\mathbf{x}) = \min_{\mathbf{D} \in \mathcal{D}} \min_{\mathbf{C} \in \mathcal{C}} \|\mathbf{X} - \mathbf{D}\mathbf{C}'\|_F^2 + \lambda^2 \|\mathbf{C}\|_0$$

$$\mathcal{D} = \{\mathbf{D} \in \mathbb{R}^{d \times J} : \|\mathbf{d}_j\|_2 = 1 \ \forall j\}, \quad \mathcal{C} = \{\mathbf{C} \in \mathbb{R}^{M \times J} : \|\mathbf{c}_j\|_\infty \leq L \ \forall j\}$$

- ▶  $\mathbf{R}(\mathbf{x}) \approx 0$  if patches can be represented closely with “sufficiently few” non-zero coefficients (depends on  $\lambda$ ).
- ▶ Ignore constraint  $\|\mathbf{c}_j\|_\infty \leq L$  in practice.
- ▶ Bayesian interpretation?

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Dictionary-blind MR image reconstruction:

$$\hat{\mathbf{x}} = \arg \min_{\mathbf{x}} \frac{1}{2} \|\mathbf{y} - \mathbf{A}\mathbf{x}\|_2^2 + \beta \mathbf{R}(\mathbf{x})$$
$$\mathbf{R}(\mathbf{x}) = \min_{\mathbf{D} \in \mathcal{D}} \min_{\mathbf{Z}' \in \mathcal{C}} \sum_{m=1}^M \left( \|\mathbf{P}_m \mathbf{x} - \mathbf{D} \mathbf{z}_m\|_2^2 + \lambda^2 \|\mathbf{z}_m\|_0 \right)$$

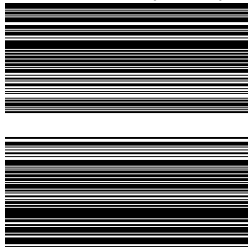
where  $\mathbf{P}_m$  extracts  $m$ th of  $M$  image patches.

*In words: of the many images...*

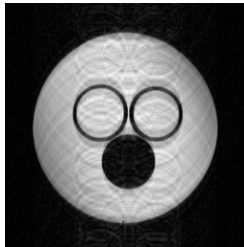
Alternating (nested) minimization:

- ▶ Fixing  $\mathbf{x}$  and  $\mathbf{D}$ , update each  $\mathbf{z}_j$  via hard-thresholding
- ▶ Fixing  $\mathbf{x}$  and  $\mathbf{Z}$ , update  $\mathbf{D}$  using SOUP-DIL
- ▶ Fixing  $\mathbf{Z}$  and  $\mathbf{D}$ , updating  $\mathbf{x}$  is a quadratic problem.
  - Efficient FFT solution for single-coil Cartesian MRI.
  - Use CG for non-Cartesian and/or parallel MRI.
- ▶ Non-convex, but monotone decreasing and some convergence theory [9].

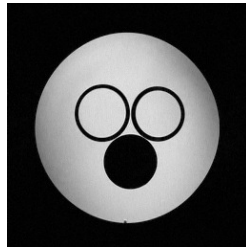
Sampling ( $2.5\times$ )



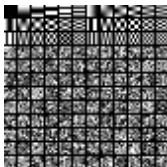
Zero-Filled



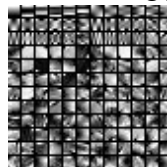
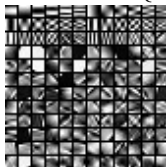
SOUP-DILLO-MRI



Initial  $D$



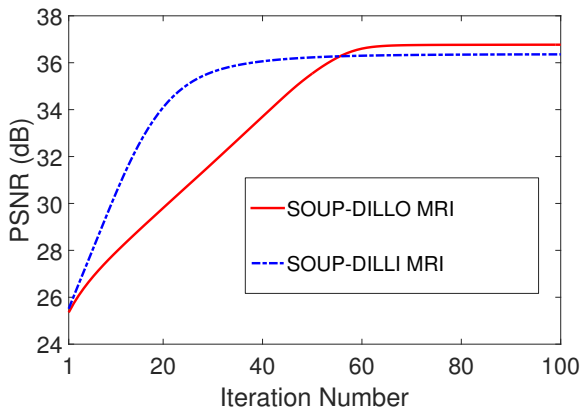
Learned real  $\{D\}$     Learned imag  $\{D\}$



$6 \times 6$  patches

$$D \in \mathbb{C}^{6^2 \times 144}$$

[9]



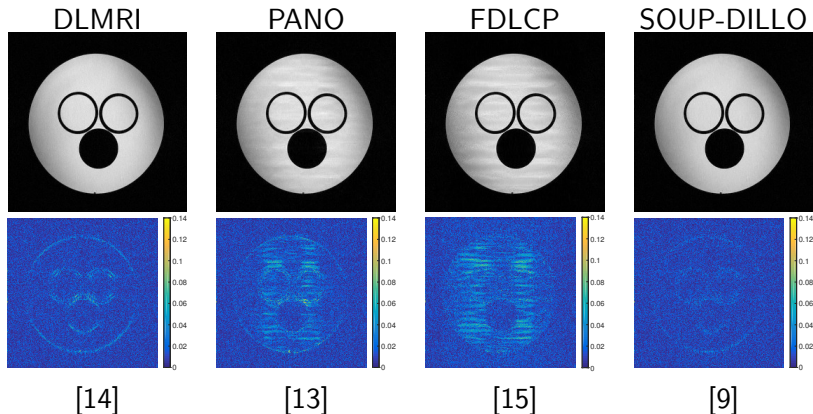
(SNR compared to fully sampled image.)

Using  $\|\mathbf{z}_m\|_0$  leads to higher SNR than  $\|\mathbf{z}_m\|_1$ .

Adaptive case is non-convex anyway...



Im.	Samp.	Acc.	0-fill	Sparse MRI	PANO	DLMRI	SOUP-DILLI	SOUP-DILLO
a	Cart.	7x	27.9	28.6	<b>31.1</b>	<b>31.1</b>	30.8	<b>31.1</b>
b	Cart.	2.5x	27.7	31.6	41.3	40.2	38.5	<b>42.3</b>
c	Cart.	2.5x	24.9	29.9	34.8	36.7	36.6	<b>37.3</b>
c	Cart.	4x	25.9	28.8	<b>32.3</b>	32.1	32.2	<b>32.3</b>
d	Cart.	2.5x	29.5	32.1	36.9	38.1	36.7	<b>38.4</b>
e	Cart.	2.5x	28.1	31.7	40.0	38.0	37.9	<b>41.5</b>
f	2D rand.	5x	26.3	27.4	30.4	30.5	30.3	<b>30.6</b>
g	Cart.	2.5x	32.8	39.1	41.6	41.7	42.2	<b>43.2</b>
Ref.				[12]	[13]	[14]	[9]	[9]



Summary: 2D static MR reconstruction from under-sampled data with adaptive dictionary learning and convergent algorithm, faster than K-SVD approach of DLMRI.

## Ill-posed problems and regularization

Classical “hand crafted” regularizers

Data-driven (adaptive / learned) regularizers

Data-driven regularized MRI via dictionary learning

Extension: learning low-rank atoms

DictioNary with lOw-ranK AToms (DINO-KAT)

## Dynamic MR imaging

DINO-KAT for dynamic MR

## Summary

## Backup

DINO-KAT convergence guarantees

Online method

Recall SOUP-DIL  $\ell_0$  formulation for dictionary learning from data  $\mathbf{X}$ :

$$\mathbf{D}^* = \arg \min_{\mathbf{D} \in \mathbb{R}^{d \times J}} \min_{\mathbf{C} \in \mathbb{R}^{N \times J}} \|\mathbf{X} - \mathbf{D}\mathbf{C}'\|_F^2 + \lambda^2 \|\mathbf{C}\|_0 \quad \text{s.t.} \quad \begin{aligned} \|\mathbf{d}_j\|_2 &= 1 \quad \forall j \\ \|\mathbf{c}_j\|_\infty &\leq L \quad \forall j. \end{aligned}$$

Recent extension [10]

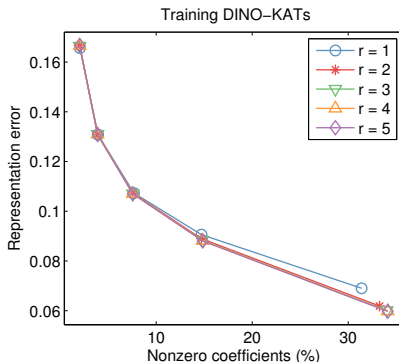
Dictionary with lOw-rank AToms (DINO-KAT) model:

$$\mathbf{D}^* = \arg \min_{\mathbf{D} \in \mathbb{R}^{d \times J}} \min_{\mathbf{C} \in \mathbb{R}^{N \times J}} \|\mathbf{X} - \mathbf{D}\mathbf{C}'\|_F^2 + \lambda^2 \|\mathbf{C}\|_0 \quad \text{s.t.} \quad \begin{aligned} \|\mathbf{d}_j\|_2 &= 1 \quad \forall j \\ \|\mathbf{c}_j\|_\infty &\leq L \quad \forall j \\ \text{rank}\{\text{reshape}(\mathbf{d}_j)\} &\leq r, \end{aligned}$$

where  $\text{reshape}(\mathbf{d}_j)$  reshapes dictionary atom  $\mathbf{d}_j$  into a 2D array.

# DINO-KAT: why low-rank atoms?

- ▶ Low-rank atoms are less prone to over-fitting.
- ▶ Model structure (e.g., temporal correlation) of dynamic data.
- ▶ Learned dictionary atoms on patch data often have only a few dominant singular values.



Representation error  $\|\mathbf{X} - \mathbf{D}\mathbf{C}'\|_F / \|\mathbf{X}\|_F$  versus sparsity  $\lambda$  for several atom ranks  $r$  for  $8 \times 8 \times 5$  space-time patches from (fully sampled) cardiac perfusion images.



Dictionary with low-rank AToms (DINO-KAT) model:

$$D^* = \arg \min_{D \in \mathbb{R}^{d \times J}} \min_{C \in \mathbb{R}^{N \times J}} \|X - DC'\|_F^2 + \lambda^2 \|C\|_0 \quad \text{s.t.} \quad \begin{aligned} &\|d_j\|_2 = 1 \quad \forall j \\ &\|c_j\|_\infty \leq L \quad \forall j \\ &\text{rank}\{\text{reshape}(d_j)\} \leq r, \end{aligned}$$

Block coordinate descent (BCD) algorithm (monotone descent)  
with simple update rules (low compute cost)

- Sparse coding step for  $C$  uses same truncated hard thresholding
- Dictionary atom update step for  $d_j$ :

$$\arg \min_{d_j} \|E_j - d_j c_j'\|_F^2 \quad \text{s.t.} \quad \|d_j\|_2 = 1, \text{rank}\{\text{reshape}(d_j)\} \leq r$$

Simple solution:  $\text{reshape}(d_j) = \frac{U_r \Sigma_r V_r'}{\|\Sigma_r\|_F}$   
 $U_r \Sigma_r V_r'$  is the rank- $r$  truncated SVD of  $\text{reshape}(E_j c_j)$ .

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“dynamic” = changing over time = motion [16–19]

► Nuisance motions:

- Breathing
- Cardiac
- Peristalsis
- Tremors
- Kids ...

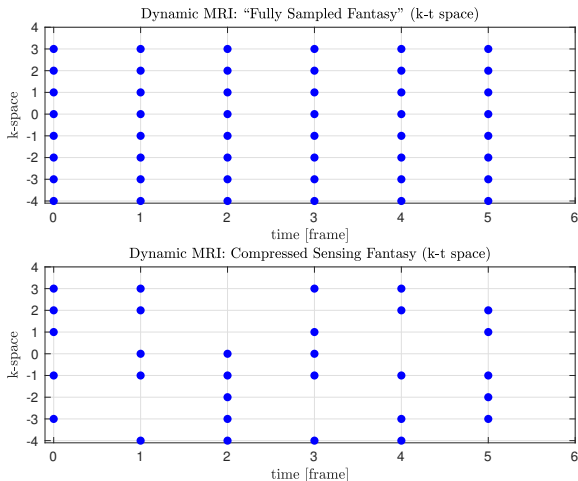
⇒ Faster scans (shorter time) can help reduce motion blur

► Motions of interest (true “dynamic” scans):

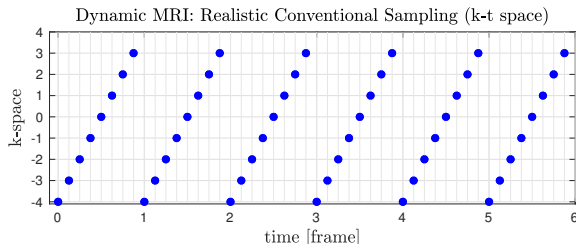
- Vocalization (for speech studies)
- Cardiac (for function)
- Joint articulation (musculoskeletal scans)
- Contrast agent (blood flow / perfusion)
- Diffusion

⇒ Trade-offs between temporal resolution and spatial resolution

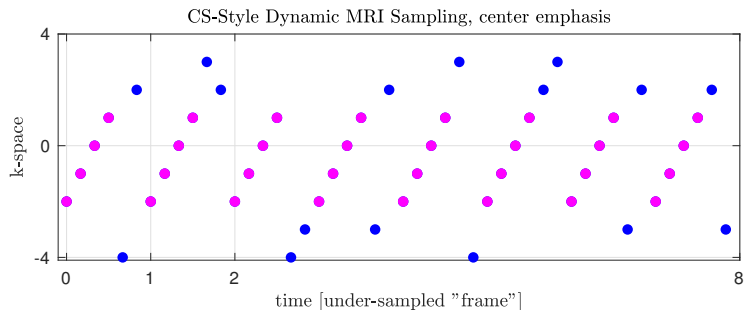
# Dynamic MRI sampling: Fantasy edition



- ▶ Scan “twice as fast” !?
- ▶ Matrix completion problem!?  $\Rightarrow$  ... robust PCA (L+S) ...  
[20, 21]



- ▶ All 3D dynamic MRI data is *inherently under-sampled*
- ▶ No real “fully sampled” data exists, now or ever
- ▶ Unlikely to satisfy any “matrix completion” sufficient conditions ( $N$  measurements but  $N^2$  unknowns per frame)
- ▶ Retrospective “under sampling” of “fully sampled” dynamic data is dubious
- ▶ Opportunity: powerful **signal models** needed for reconstruction from such data
- ▶ Challenge: validation of signal models given such highly incomplete data (low-rank / locally low rank / tensors / wavelets / non-local patches / ...)



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- Extension: learning low-rank atoms

  - DictioNary with lOw-ranK AToms (DINO-KAT)

## Dynamic MR imaging

- DINO-KAT for dynamic MR

## Summary

## Backup

- DINO-KAT convergence guarantees

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DINO-KAT as an adaptive (data-driven) regularizer:

$$\hat{\mathbf{x}} = \arg \min_{\mathbf{x}} \frac{1}{2} \|\mathbf{y} - \mathbf{A}\mathbf{x}\|_2^2 + \beta \mathbf{R}(\mathbf{x})$$

$$\mathbf{R}(\mathbf{x}) = \min_{\mathbf{D} \in \mathbb{C}^{d \times J}} \min_{\mathbf{Z} \in \mathbb{C}^{J \times M}} \sum_{m=1}^M \|\mathbf{P}_m \mathbf{x} - \mathbf{D} \mathbf{z}_m\|_2^2 + \lambda^2 \|\mathbf{z}_m\|_0$$

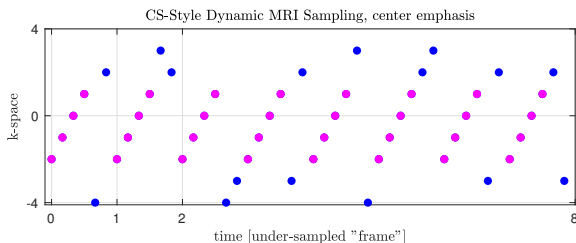
$$\text{s.t. } \|\mathbf{d}_j\|_2 = 1 \ \forall j, \ \|\mathbf{z}_m\|_\infty \leq L \ \forall m, \ \text{rank}\{\text{reshape}(\mathbf{d}_j)\} \leq r$$

Block coordinate descent (BCD) algorithm (monotone descent)

- Update coefficients  $\mathbf{Z}$ : sparse coding via hard thresholding
- Dictionary atom update of  $\mathbf{d}_j$ : uses residual, SVD
- Image update uses FFT (single coil Cartesian) or CG

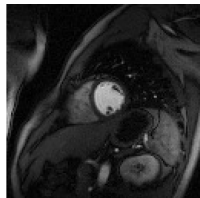
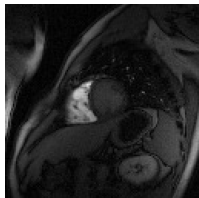


- ▶ Latent signal vector  $\mathbf{x} \in \mathbb{C}^{n_y n_x n_t}$  modeled as  $n_t$  frames, each of dimension  $N = n_x \times n_y$  or  $N = n_x \times n_y \times n_z$ .
- ▶  $k$ -space data  $\mathbf{y} \in \mathbb{C}^{n_{\text{sample}} n_c}$  acquired using  $n_c$  coils.
- ▶ Sensing matrix  $\mathbf{A}$  includes:
  - coil sensitivity maps,
  - 2D or 3D spatial Fourier transform,
  - $k$ -space sampling pattern.
- ▶  $\mathbf{y}$  is undersampled, so regularization is required to estimate dynamic image sequence  $\mathbf{x}$ .



- ▶ **Low-rank and sparse (k-t SLR) [22]:**
  - Model:  $\mathbf{x}$  reshaped into an  $N \times n_t$  space-time matrix, is both low-rank and (transform) sparse
- ▶ **Low-rank plus sparse ( $\mathbf{L} + \mathbf{S}$ ) [20, 21]**
  - model:  $\mathbf{x} = \mathbf{x}_L + \mathbf{x}_S$ ,
  - $\mathbf{x}_L$  reshaped into a  $N \times n_t$  space-time matrix is low-rank,
  - $\mathbf{x}_S$  is (transform) sparse.
- ▶ **DINO-KAT for dynamic MRI [10, 23]:**
  - Extract  $p \times p \times q$  patches of  $\mathbf{x}_S$ .
  - Model patches as sparse w.r.t. an adaptive (learned) dictionary  $\mathbf{D}$ .
  - Model dictionary atoms  $\{\mathbf{d}_j\}$  as low-rank when reshaped into  $p^2 \times q$  space-time matrices.
  - Blind compressed sensing model [14].

Cardiac perfusion data (ref. frames 7, 13)



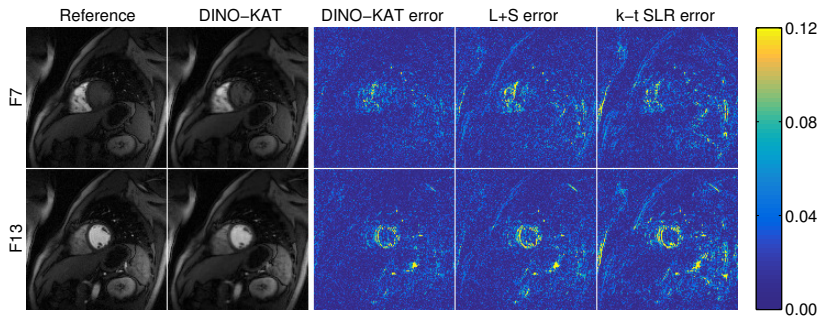
$128^2 \times 40$  fr.  
 $3.2^2 \times 8$  mm<sup>3</sup>  
12 coil  
 $\Delta T = 307$  ms  
Otazo et al. [21] (L+S)

PINCAT data (reference frames 16, 25)

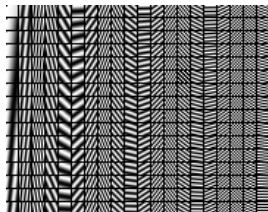


$128^2 \times 50$  fr.  
 $1.5$  mm<sup>2</sup>  
1 coil?  
 $9 \times$  acc.  
Lingala et al. [22] (k-t SLR)

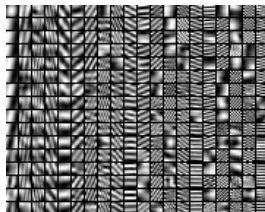
8 $\times$  acceleration



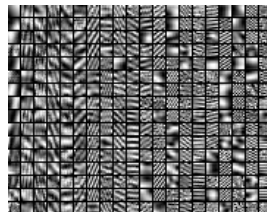
Initial atoms  
(DCT matrix)



Real-part  
of learned atoms

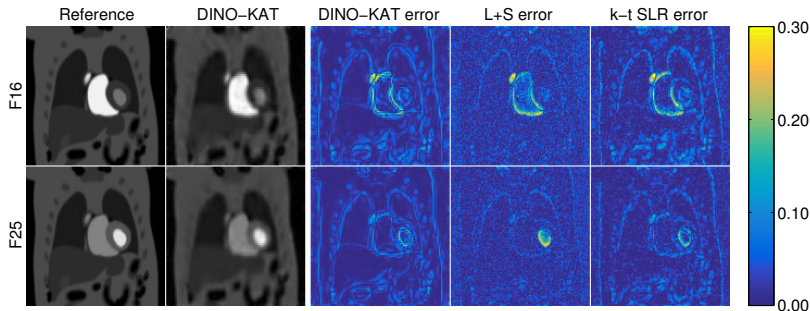


Imaginary-part  
of learned atoms



- ▶ First temporal slices of  $8 \times 8 \times 5$  atoms
- ▶ Learned atoms adapt to structure of data

9× acceleration



- ▶ Two representative frames of each reconstruction
- ▶ DINO-KAT method shows less error than the L+S and k-t SLR (L&S) methods

8× acceleration

9× acceleration



# Quantitative results for cardiac perfusion data

Acceleration	4x	8x	12x	16x	20x	24x
NRMSE (L+S) %	10.93	14.00	15.80	18.87	21.33	23.36
NRMSE (Fixed D) %	11.29	13.76	15.33	18.31	20.77	22.82
NRMSE ( $r = 5$ ) %	10.85	13.08	14.37	17.01	19.19	21.35
NRMSE ( $r = 1$ ) %	<b>10.57</b>	<b>12.90</b>	<b>14.20</b>	<b>16.77</b>	<b>18.74</b>	<b>20.91</b>
Gain over L + S (dB)	0.29	0.71	0.92	1.03	1.13	0.96
Gain over $r = 5$ (dB)	0.23	0.12	0.10	0.13	0.21	0.18

- ▶ Data-driven / adaptive regularization
  - Beneficial for under-sampled MRI reconstruction
  - Dictionary atom structure (e.g., low rank) further helpful
  - SOUP provides reasonably computationally efficient methods (vs KSVD)
  - Convergence theory (unlike KSVD)
- ▶ Future work:
  - Synthesis (e.g., dictionary) vs analysis (e.g., transform learning) formulations
  - Online methods for reduced memory, better adaptation [24–27]
  - Other machine-learning methods (deep...) ?
  - T-MI special issue on Machine-Learning for Image Reconstruction

## Ill-posed problems and regularization

- Classical “hand crafted” regularizers

- Data-driven (adaptive / learned) regularizers

- Data-driven regularized MRI via dictionary learning

- Extension: learning low-rank atoms

  - DictioNary with lOw-ranK AToms (DINO-KAT)

## Dynamic MR imaging

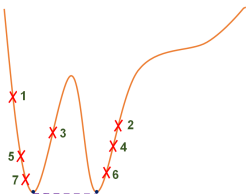
- DINO-KAT for dynamic MR

## Summary

## Backup

- DINO-KAT convergence guarantees

- Online method



## Theorem (Convergence guarantees)

[10, 23, 27]

Let  $\{D^t, C^t\}$  be the iterate sequence generated by the BCD algorithm for DINO-KAT. Then

- ▶ The cost function  $\Psi(D^t, C^t)$  is monotone decreasing and converges to a finite value, say  $\Psi^*$
- ▶ The iterate sequence  $\{D^t, C^t\}$  is bounded and its accumulation points have common cost value  $\Psi^*$

- For frame  $t = 1, 2, 3, \dots$ , solve

$$\begin{aligned}
 \text{(P3)} \quad \{ \hat{W}_t, \hat{x}_t \} = & \arg \min_{W, x_t} \frac{1}{t} \sum_{j=1}^t \{ \overbrace{\|Wy_j - x_j\|_2^2}^{\text{Sparsification Error}} + \overbrace{\lambda_j v(W)}^{\text{Regularizer}} \} \\
 \text{s.t.} \quad & \|x_t\|_0 \leq s, \quad x_j = \hat{x}_j, \quad 1 \leq j \leq t-1.
 \end{aligned}$$

- Minimize the average cost computed over the observed signals.
- $\hat{W}_t^{-1} \hat{x}_t$  is an (e.g., denoised) estimate of  $y_t$ .
- For non-stationary data, use forgetting factor  $\rho \in [0, 1]$ , to diminish the influence of old data.

$$\frac{1}{\sum_{j=1}^t \rho^{t-j}} \sum_{j=1}^t \rho^{t-j} \left\{ \|Wy_j - x_j\|_2^2 + \lambda_j v(W) \right\}$$

- **Sparse Coding:** solve for  $x_t$  in (P3) with fixed  $W = \hat{W}_{t-1}$ .

$$\min_{x_t} \|Wy_t - x_t\|_2^2 \quad \text{s.t.} \quad \|x_t\|_0 \leq s$$

- **Cheap Solution:**  $\hat{x}_t = H_s(Wy_t)$ .

- **Transform Update:** solve for  $W$  with  $x_t = \hat{x}_t$ .

$$\min_W \frac{1}{t} \sum_{j=1}^t \left\{ \|W y_j - x_j\|_2^2 + \lambda_j \left( \|W\|_F^2 - \log |\det W| \right) \right\}$$

$$\hat{W}_t = 0.5 R_t \left( \Sigma_t + (\Sigma_t^2 + 2\beta_t I)^{\frac{1}{2}} \right) Q_t^T L_t^{-1}$$

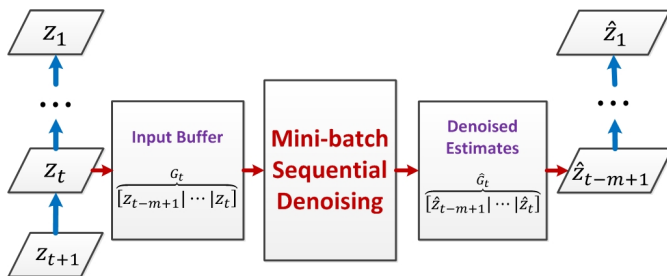
- $t^{-1} \sum_{j=1}^t (y_j y_j^T + \lambda_0 \|y_j\|_2^2 I) = L_t L_t^T$ . **Perform rank-1 update.**
- $\beta_t = \lambda_0 t^{-1} \sum_{j=1}^t \|y_j\|_2^2$ .  $Q_t \Sigma_t R_t^T$  is full SVD of  $L_t^{-1} \Theta_t = t^{-1} \sum_{j=1}^t L_t^{-1} y_j x_j^T$ .
  - $L_t^{-1} \Theta_t \approx (1 - t^{-1}) L_{t-1}^{-1} \Theta_{t-1} + t^{-1} L_t^{-1} y_t x_t^T$   
 $\implies$  **efficient rank-1 SVD update.**

- ▶ **Assumption:**  $y_t$  are i.i.d. random samples from the sphere  $S^{n-1} = \{y \in \mathbb{R}^n : \|y\|_2 = 1\}$ , for an absolutely continuous probability measure  $p$ .
- ▶ We consider minimizing the **expected [28] learning cost:**

$$g(W) = \mathbb{E}_y \left[ \|Wy - H_s(Wy)\|_2^2 + \lambda_0 \|y\|_2^2 v(W) \right].$$

- ▶ **Main Result [25]:**  $\hat{W}_t$  in OTL converges to the set of stationary points of  $g(W)$  almost surely.  
 $\hat{W}_{t+1} - \hat{W}_t \sim O(1/t)$ .



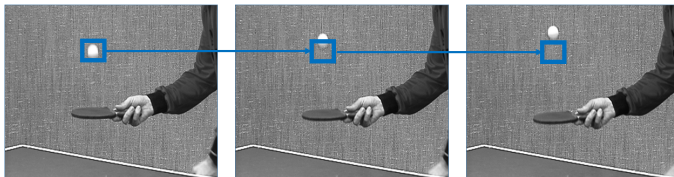


- ▶  $z_t$ : noisy frame,  $\hat{z}_t$ : denoised version.
- ▶  $G_t$ : 3D array with  $m$  frames formed using a sliding window scheme.
- ▶ Overlapping 3D patches in the  $G_t$ 's are denoised sequentially with OTL.
- ▶ Denoised patches averaged at 3D locations to yield frame estimates.

# 3D Patches in Proposed VIDOSAT Method

Direct  
Extraction

VIDOSAT



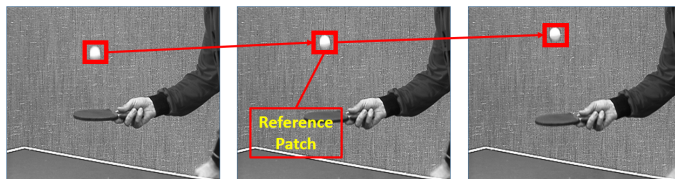
$z_{t-1}$

$z_t$

$z_{t+1}$

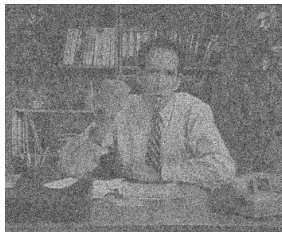
Block  
Matching

VIDOSAT-BM

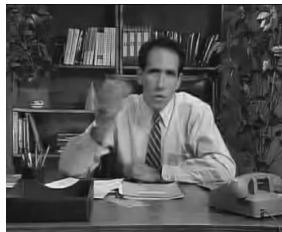


- Transform is learned online from sequentially extracted 3D patches.

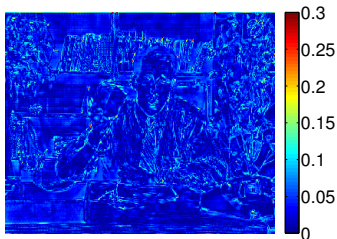
# Video Denoising Example: Salesman



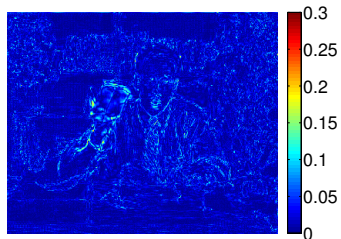
Noisy frame



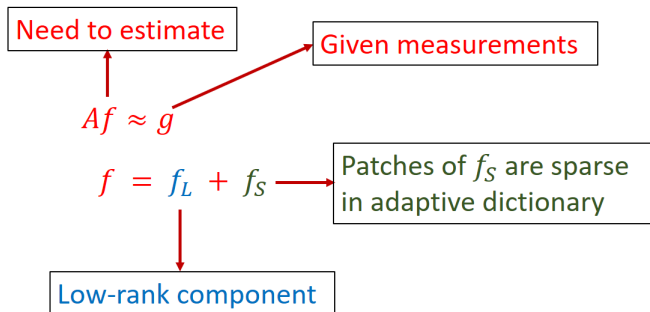
VIDOSAT (PSNR = 30.97 dB)



VBM4D [29] Error (PSNR = 27.20 dB)



VIDOSAT Error



- ▶ Low-rank + **Adaptive** Sparse Signal (**LASSI**) model for dynamic data.
- ▶ **Goal:** learn dictionary and reconstruct  $(f_L, f_S)$  from limited  $g$ .
- ▶ Efficient algorithm for LASSI estimation proposed recently [23]

open

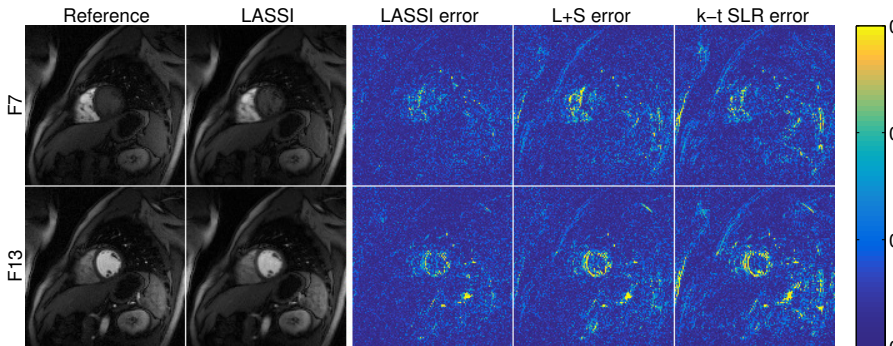
`~/tex/paper/done/17/ravishankar-17-lra/gifs/tmi_otazo_8x_lassi_vs_lps.gif`

open

`~/tex/conf/misc/ivmsp,16/ravishankar-lassi/talk/tmi_otazo_8x_lassi_vs_lps.avi`

Click for LASSI and L+S Results

# Cardiac Perfusion Results: 8x Acceleration



- Error maps for LASSI show smaller distortions than for the k-t SLR [22] (L & S) method and the L+S [21] method.

- [1] E. Candès and J. K. Romberg. "Signal recovery from random projections." In: *Proc. SPIE 5674 Computational Imaging III*. 2005, 76–86.
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