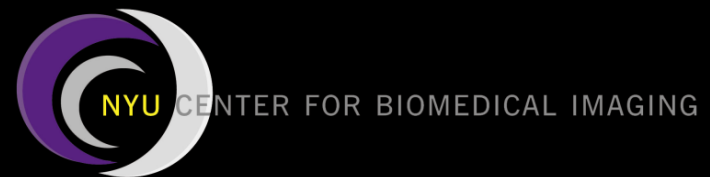


*Imaging with Modulated/Undersampled Data 2014*

*SBF Workshop, Graz, Austria*

# Low-rank plus sparse (L+S) matrix reconstruction for accelerated dynamic MRI

Ricardo Otazo



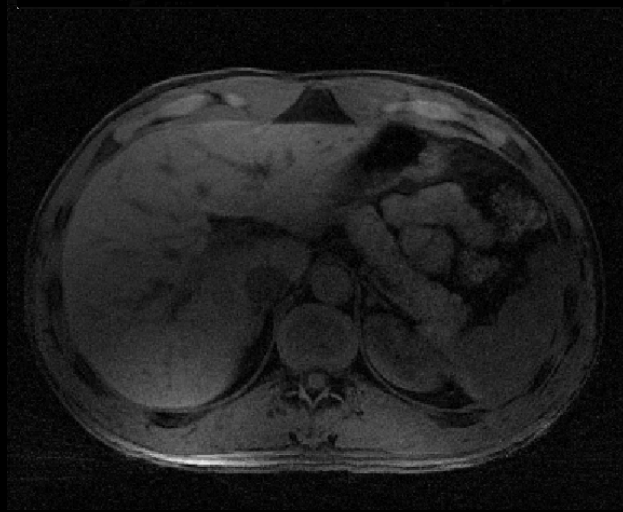
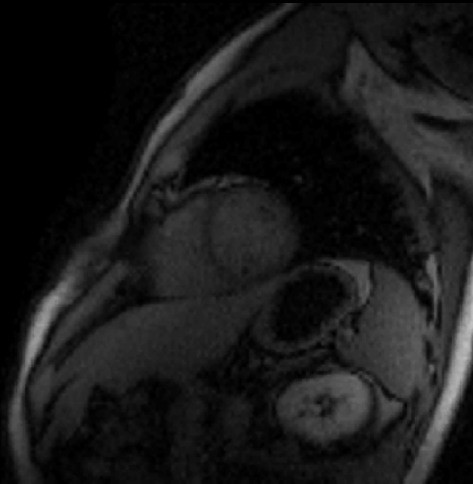
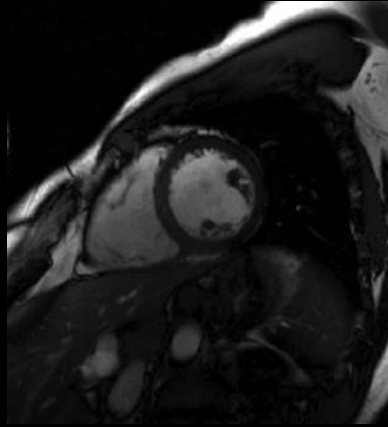
# Need for speed in dynamic MRI

*I feel the need ... the need for speed – Top Gun movie*



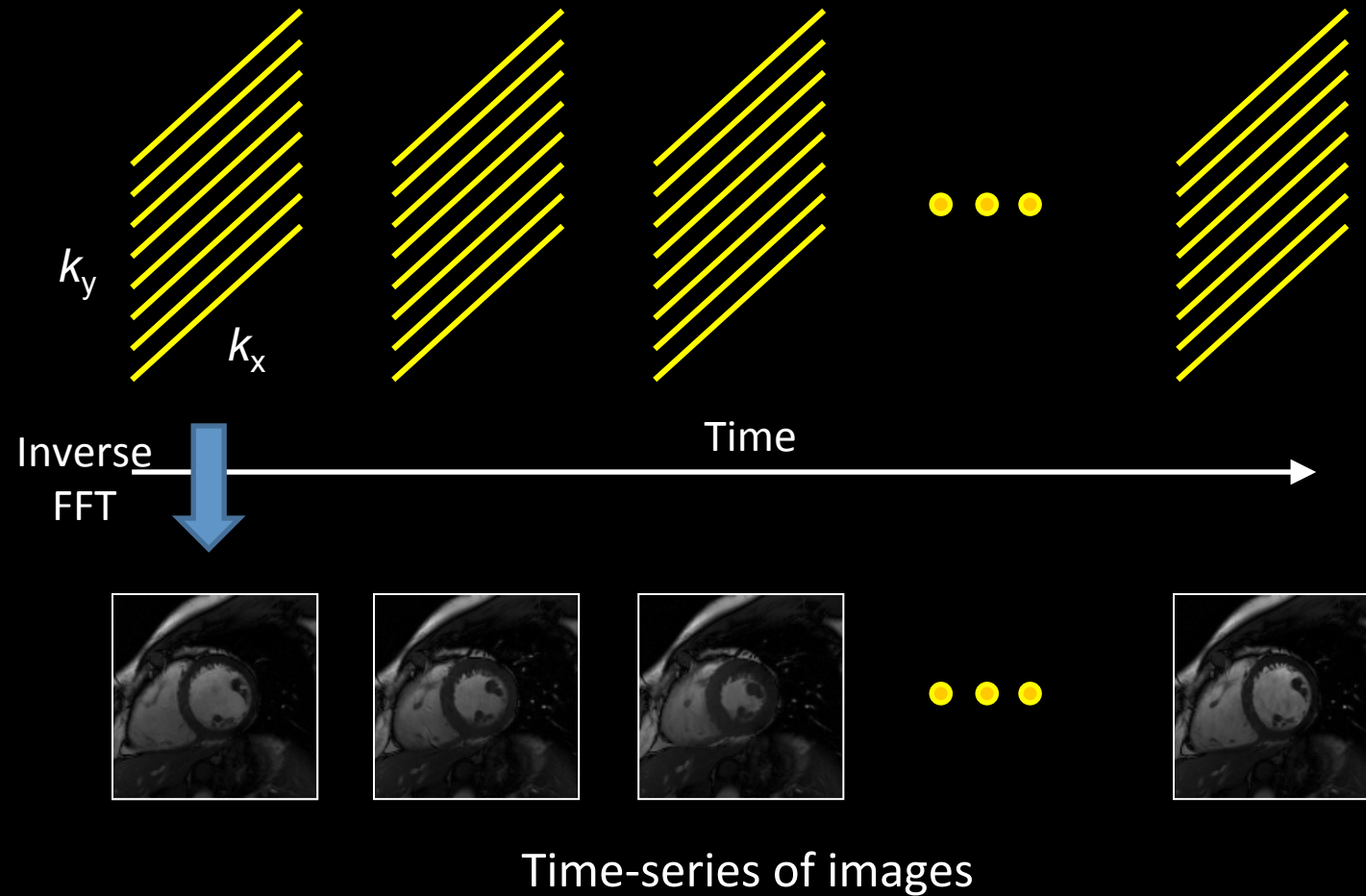
# Dynamic MRI

- Time-series of images



# Dynamic MRI

- k-t data acquisition

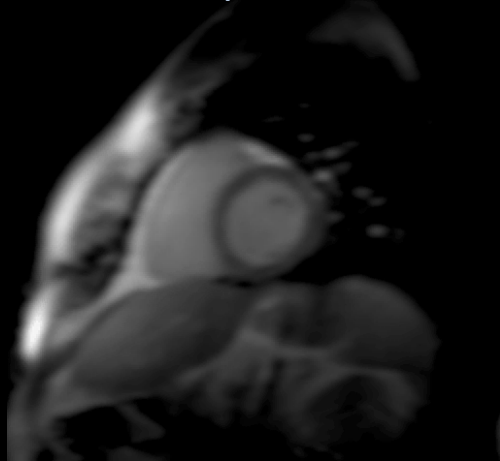


# Dynamic MRI feels the need for speed

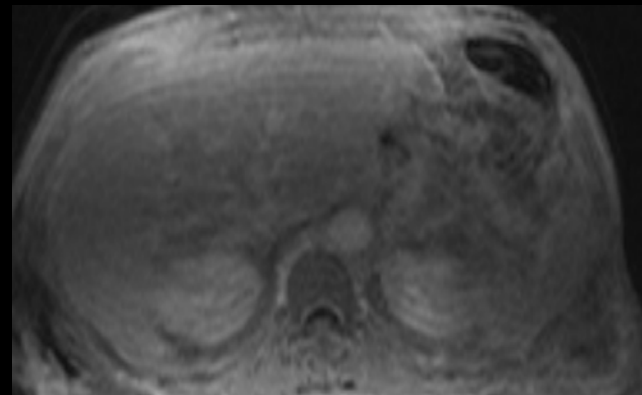
- Current dynamic MRI techniques are slow
  - Switching rate of magnetic field gradients
  - Noise amplification in parallel imaging
- Consequences



*Sacrifice spatial resolution  
for better temporal resolution*

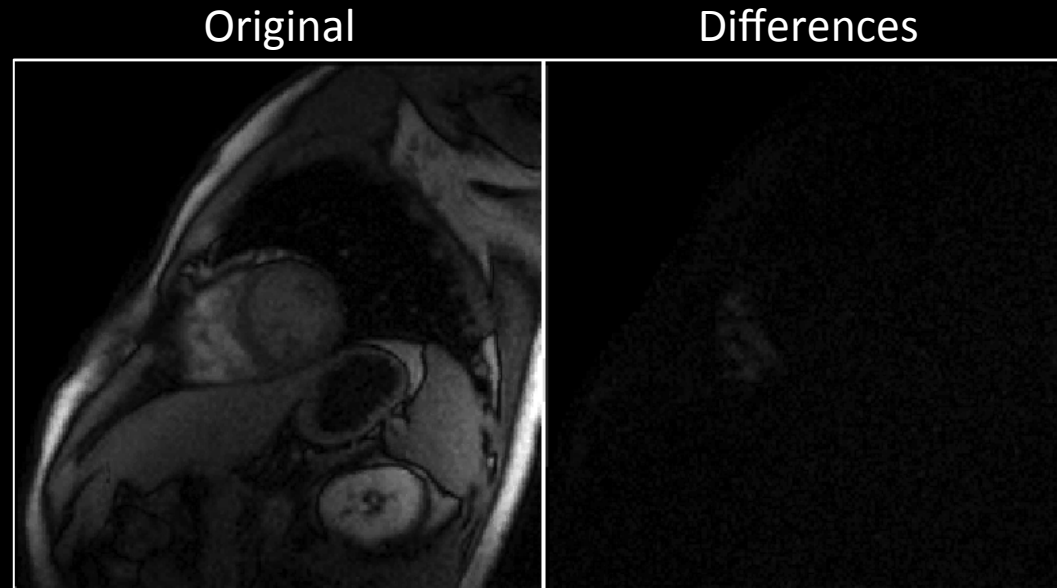


*Motion-related artifacts*



# Dynamic MRI feels the need for speed

- Full k-space sampling for each time point is redundant



- Shouldn't we be just sampling the differences from frame-to-frame?
  - Sampling rate should match the information rate

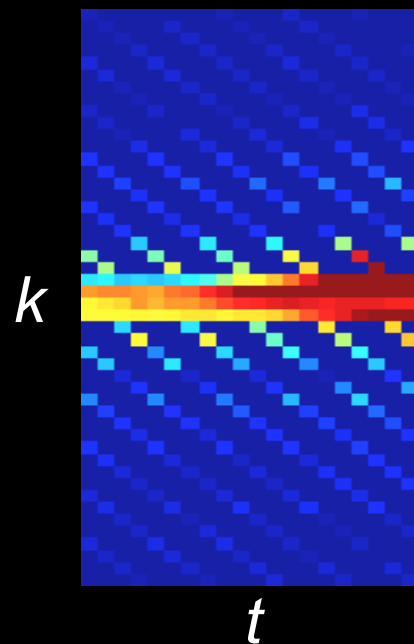
# Accelerated dynamic MRI



# k-t undersampling

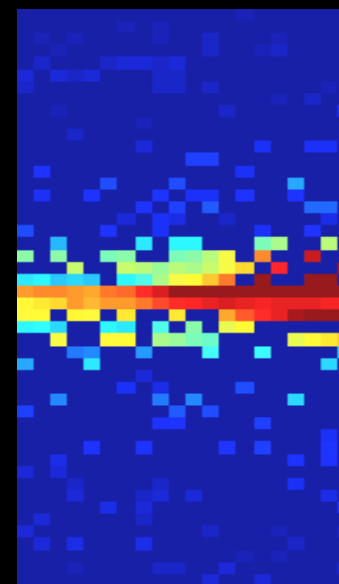
- Different sampling pattern for each time point

Uniform



UNFOLD  
k-t BLAST/SENSE

Random



Compressed sensing  
Matrix completion



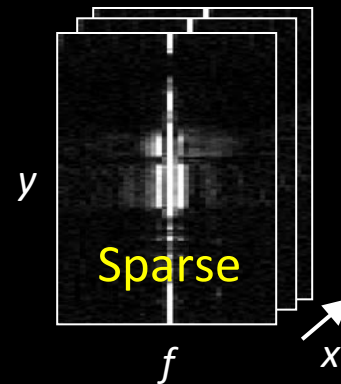
# Reconstruction of undersampled k-t data

- Exploit spatiotemporal correlations
- Dynamic MRI data are **compressible/sparse**
  - *A movie is easier to compress than an image*

Original



Temporal frequencies

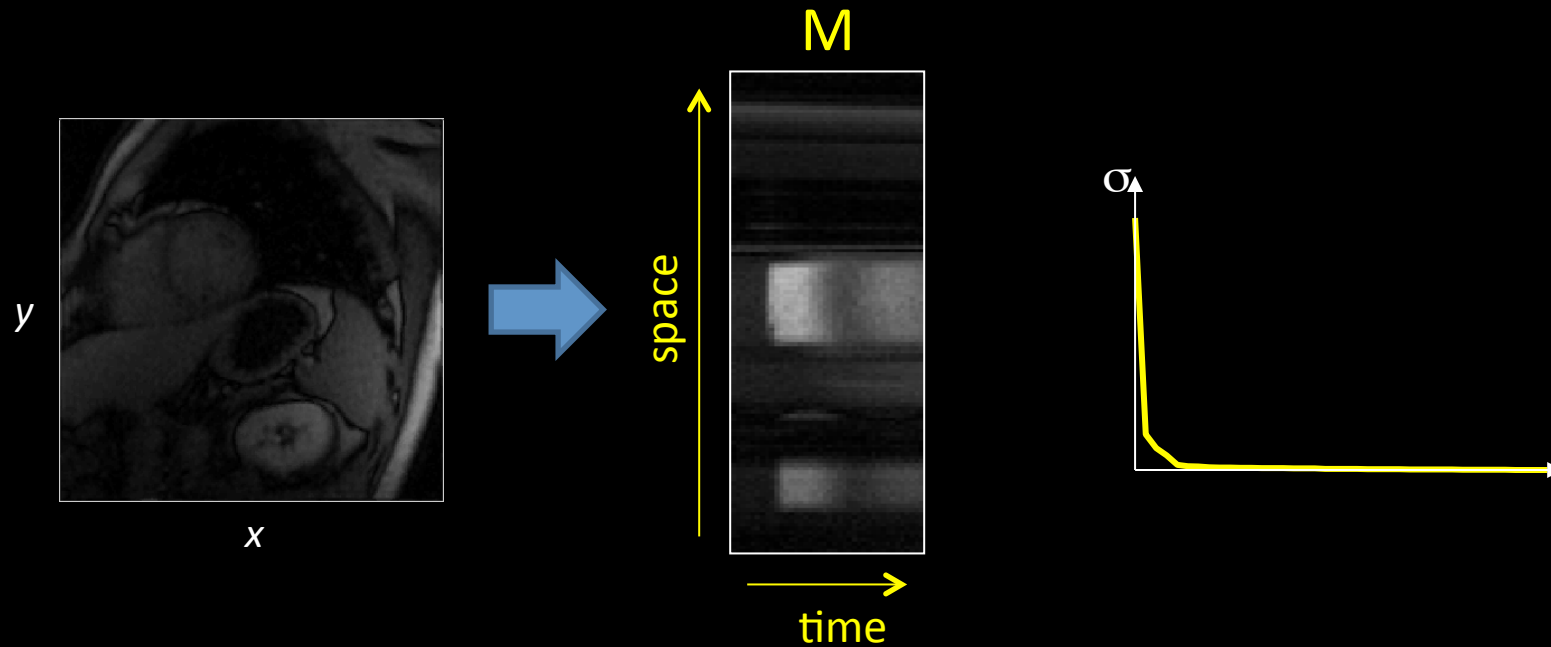


x20 Compressed



# Reconstruction of undersampled k-t data

- Exploit spatiotemporal correlations
- Dynamic MRI data are **low-rank**



# Reconstruction of undersampled k-t data

- Compressed sensing (transform sparsity)

$$\text{minimize } \|Tx\|_1 \text{ subject to } Ex = y$$

Lustig M et al. ISMRM 2006  
Gamper U et al. MRM 2008

Jung H et al. MRM 2009  
Otazo R et al. MRM 2010

- Matrix completion (low-rank)

$$\text{minimize } \|M\|_* \text{ subject to } EM = y$$

Haldar J et al. IEEE ISBI 2010  
Otazo R et al. ISMRM 2012

# Low-rank plus sparse reconstruction

# L+S reconstruction of undersampled k-t data

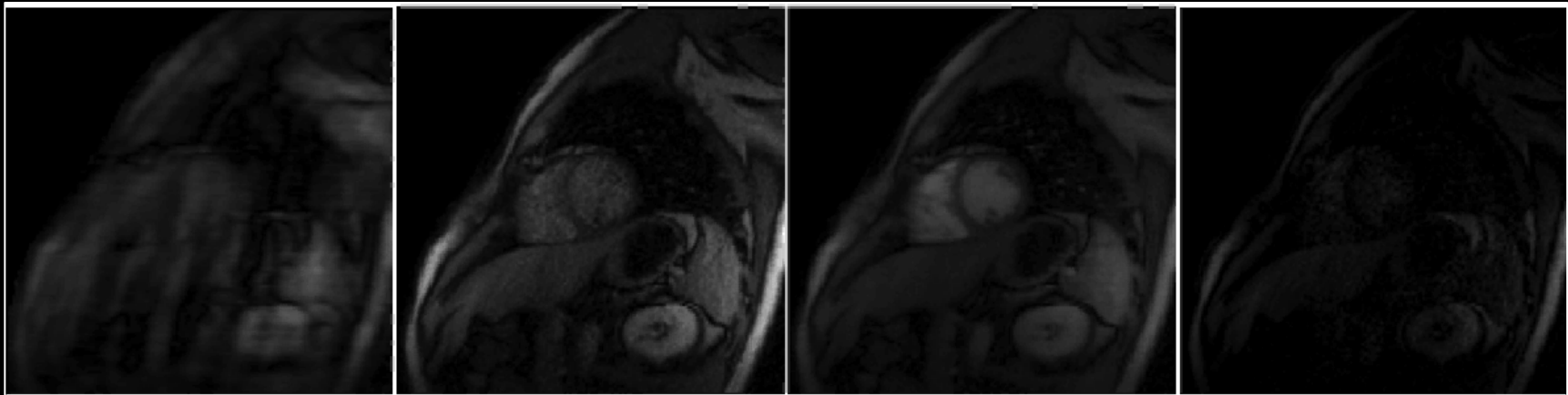
- Combination of matrix completion and compressed sensing
- Based on L+S matrix decomposition (a.k.a. robust PCA)

Conventional DFT

L+S

L

S



8-fold undersampling

# Low-rank plus sparse matrix decomposition

- a.k.a. robust principal component analysis (RPCA)

$$\mathbf{M} = \mathbf{L} + \mathbf{S}$$

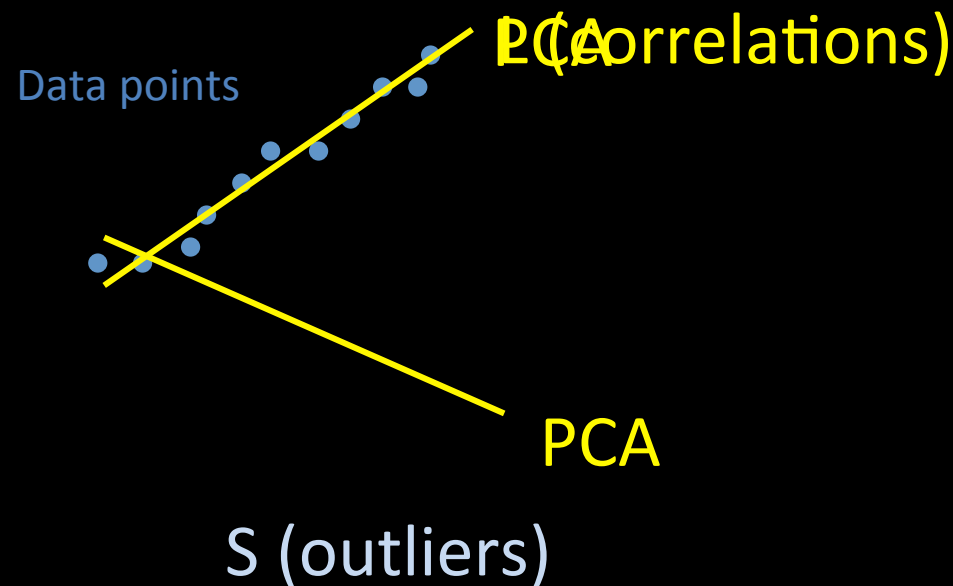
M: data matrix

L: low-rank matrix

S: sparse matrix

# Low-rank plus sparse matrix decomposition

- Motivation: standard PCA is sensitive to outliers



# Low-rank plus sparse matrix decomposition

- Convex optimization problem
- Find  $L$  and  $S$  to

$$\begin{aligned} & \text{minimize } \|L\|_* + \lambda \|S\|_1 \\ & \text{subject to } M = L + S \end{aligned}$$

$\|\cdot\|_*$ : nuclear-norm (sum of singular values)

$\|\cdot\|_1$ :  $l_1$ -norm (sum of absolute values)



# Low-rank plus sparse matrix decomposition

- Requirements

- Singular vectors of  $L$  cannot be sparse

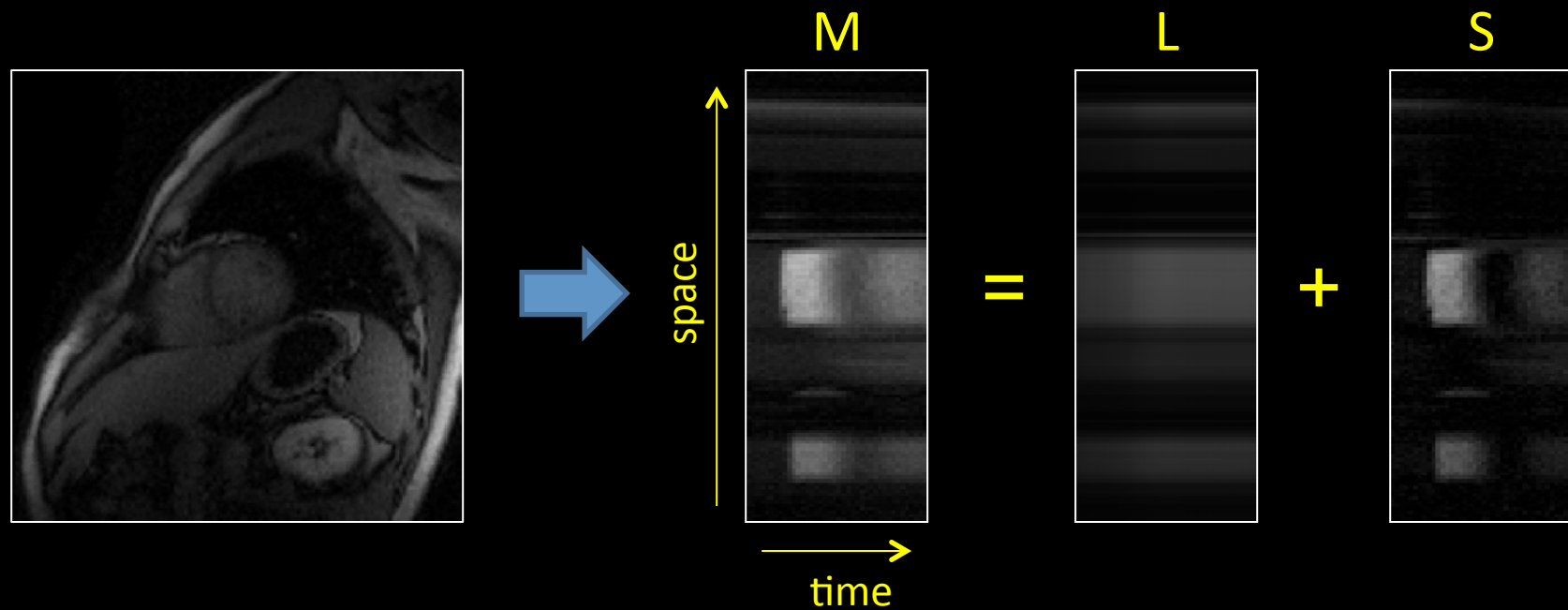
- $S$  cannot have a low-rank representation

} Rank-sparsity  
incoherence

- No need to know the rank of  $L$  or the sparsity of  $S$

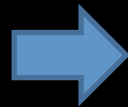
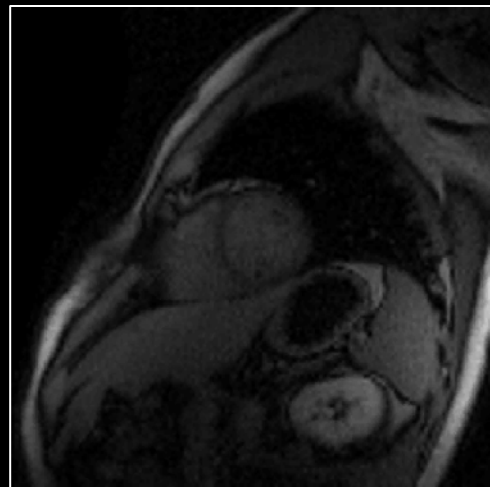
# L+S decomposition of dynamic MRI data

- Cardiac perfusion example



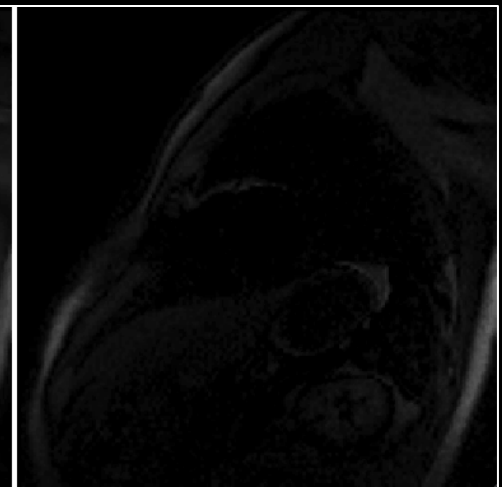
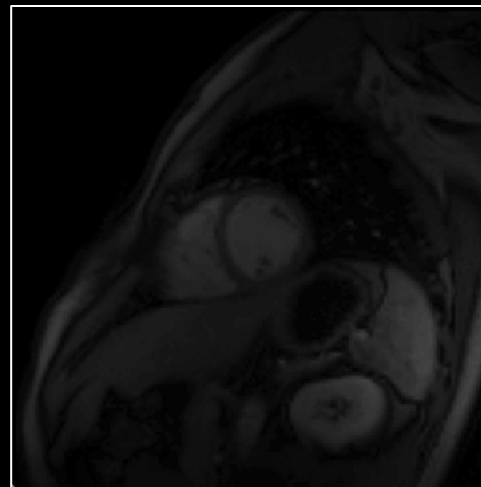
# L+S decomposition of dynamic MRI data

- Cardiac perfusion example



L

S



Background

Innovations

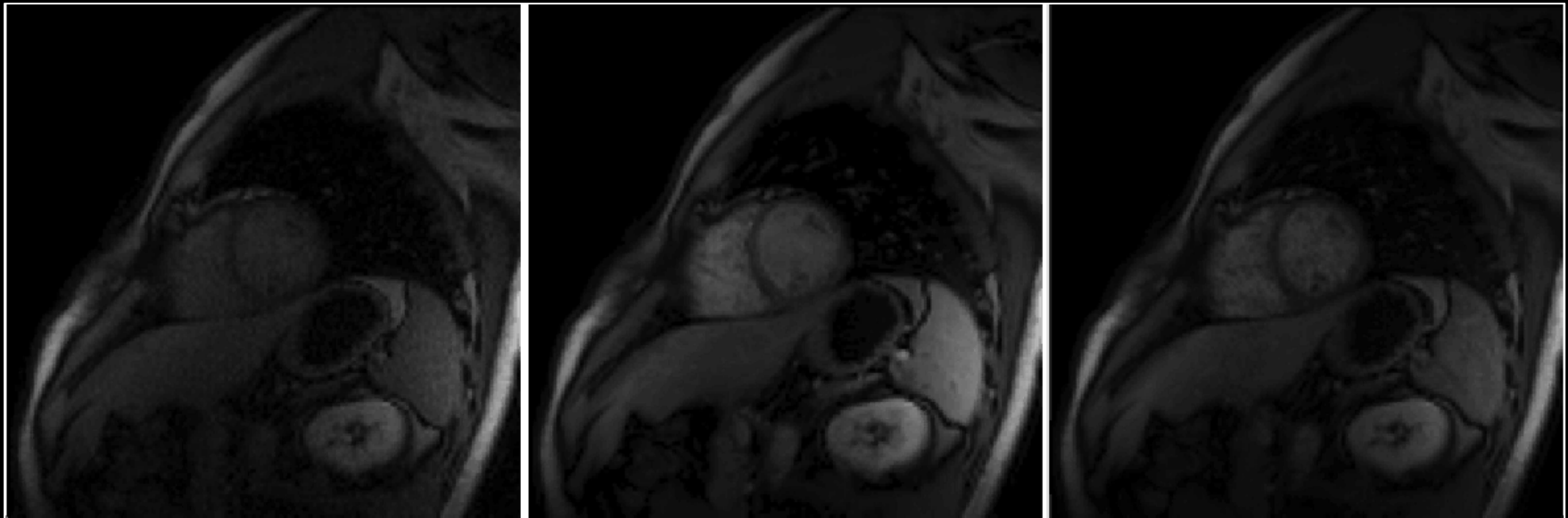
# L+S decomposition of dynamic MRI data

- Increased compressibility

No compression

X30 S-only compression

X30 L+S compression



# L+S reconstruction of undersampled k-t data

- Find L and S to

$$\begin{aligned} &\text{minimize } \|L\|_* + \lambda \|TS\|_1 \\ &\text{subject to } E(L + S) = d \end{aligned}$$

---

L: low-rank component

S: sparse component

T: temporal sparsifying transform

E: encoding operator (Fourier transform + coil sensitivities)

d: undersampled k-t data

# L+S reconstruction of undersampled k-t data

- Incoherence between

- Undersampled k-t and L

- Undersampled k-t and TS

} Remove aliasing

- L and TS

} Background/dynamic separation

# L+S reconstruction of undersampled k-t data

- Proximal gradient algorithm (iterative soft-thresholding)

$$[L, S] = \min_{L, S} \|E(L + S) - d\|_2^2 + \lambda_L \|L\|_* + \lambda_S \|TS\|_1$$

**initial solution:**  $M_0 = E^*d, S_0 = 0$


**while** not converged **do:**

$$L_k = SVT_{\lambda_L}(M_{k-1} - S_{k-1})$$

$$S_k = T^{-1}(\Lambda_{\lambda_S}(T(M_{k-1} - L_{k-1})))$$

$$M_k = L_k + S_k - E^*(E(L_k + S_k) - d)$$

**end while**

Soft-thresholding:  $\Lambda_\lambda(x) = \frac{x}{|x|} \max(|x| - \lambda, 0)$  

Singular value thresholding:  $SVT_\lambda(M) = U\Lambda_\lambda(\Sigma)V^H$

# L+S reconstruction of cardiac cine

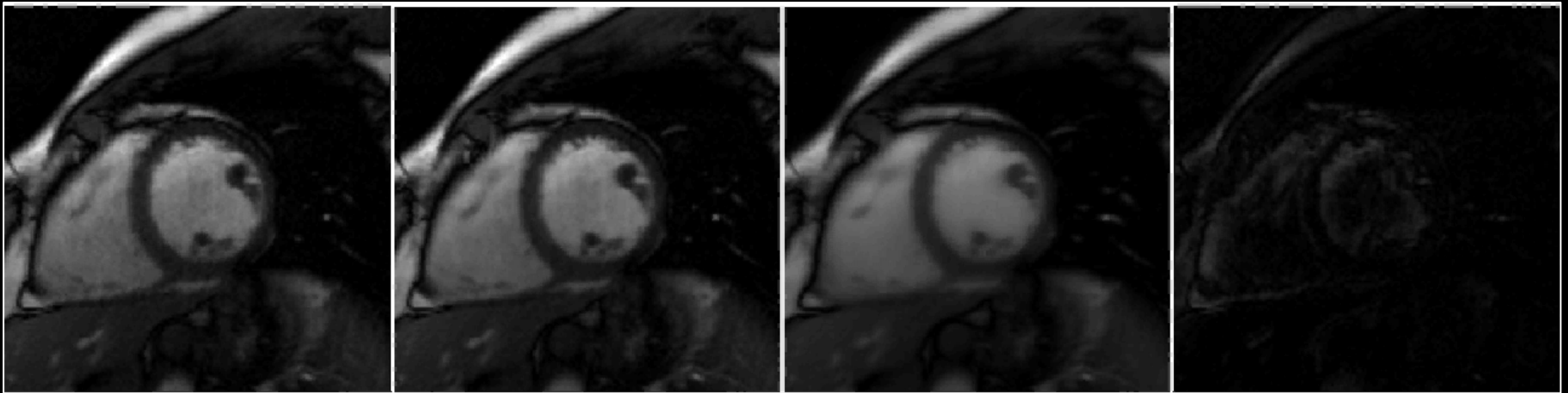
- 6-fold acceleration ( $k_y$ -t random undersampling )
- Temporal resolution = 40 ms
- Spatial resolution =  $1.3 \times 1.3 \times 3 \text{mm}^3$
- T: temporal FFT

CS

L+S

L

S

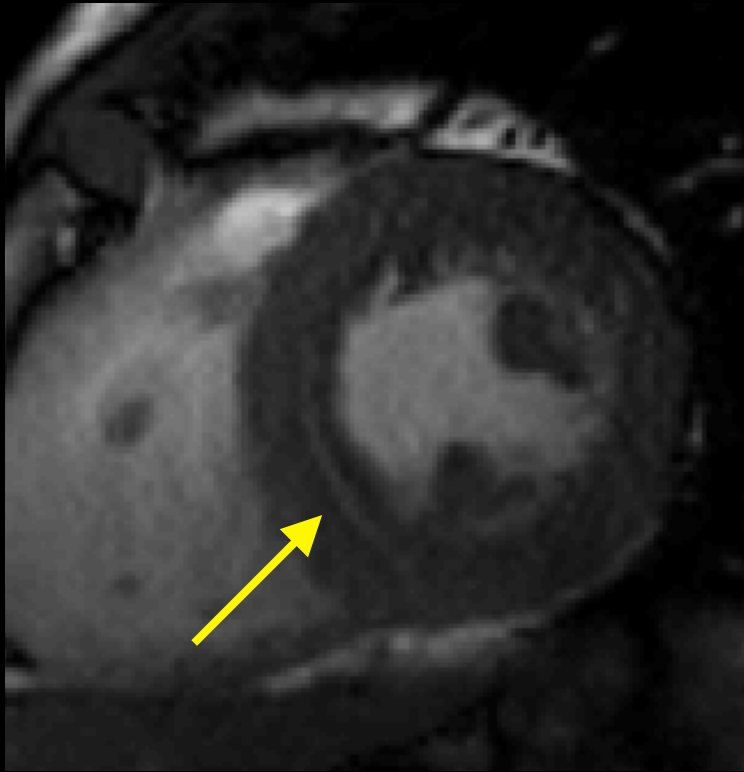


CS: standard compressed sensing using T



# L+S reconstruction of cardiac cine

CS



L+S



# L+S reconstruction of cardiac perfusion (patient)

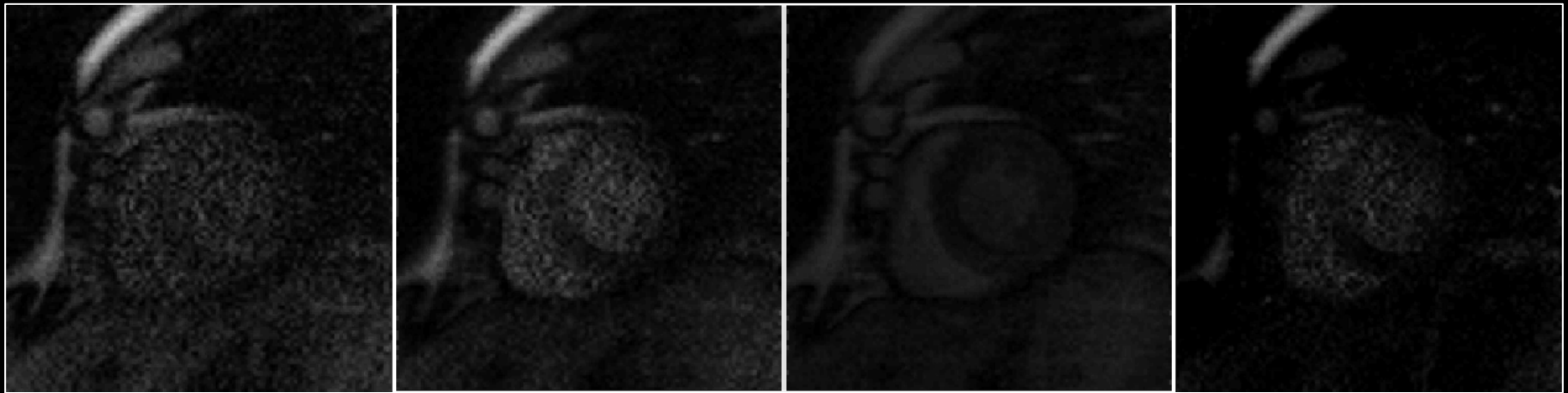
- 8-fold acceleration ( $k_y$ -t random undersampling)
- Temporal resolution = 60 ms
- Spatial resolution =  $1.7 \times 1.7 \times 3 \text{mm}^3$
- T: temporal FFT

CS

L+S

L

S



Patient with coronary artery disease

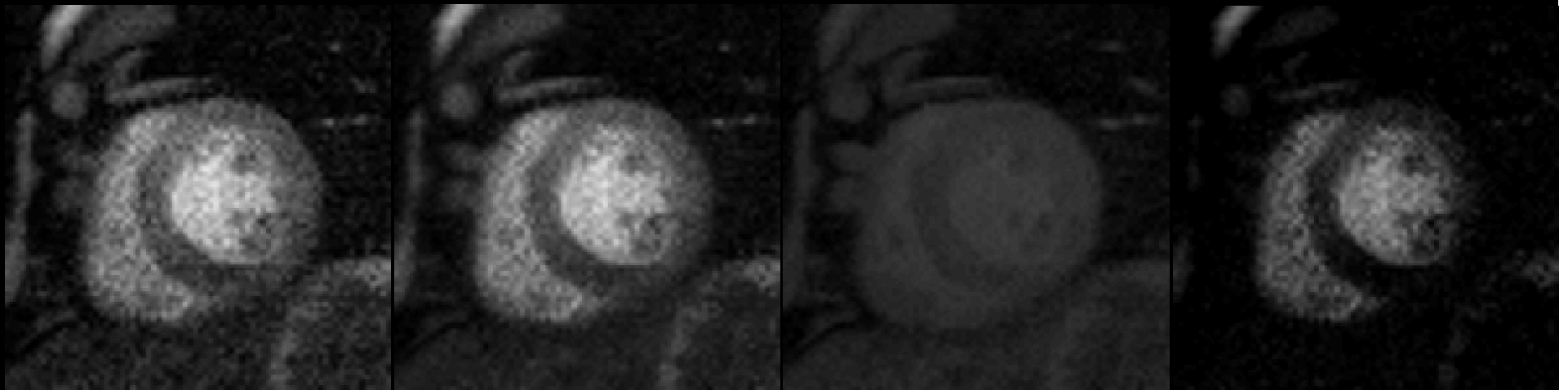
# L+S reconstruction of cardiac perfusion (patient)

CS

L+S

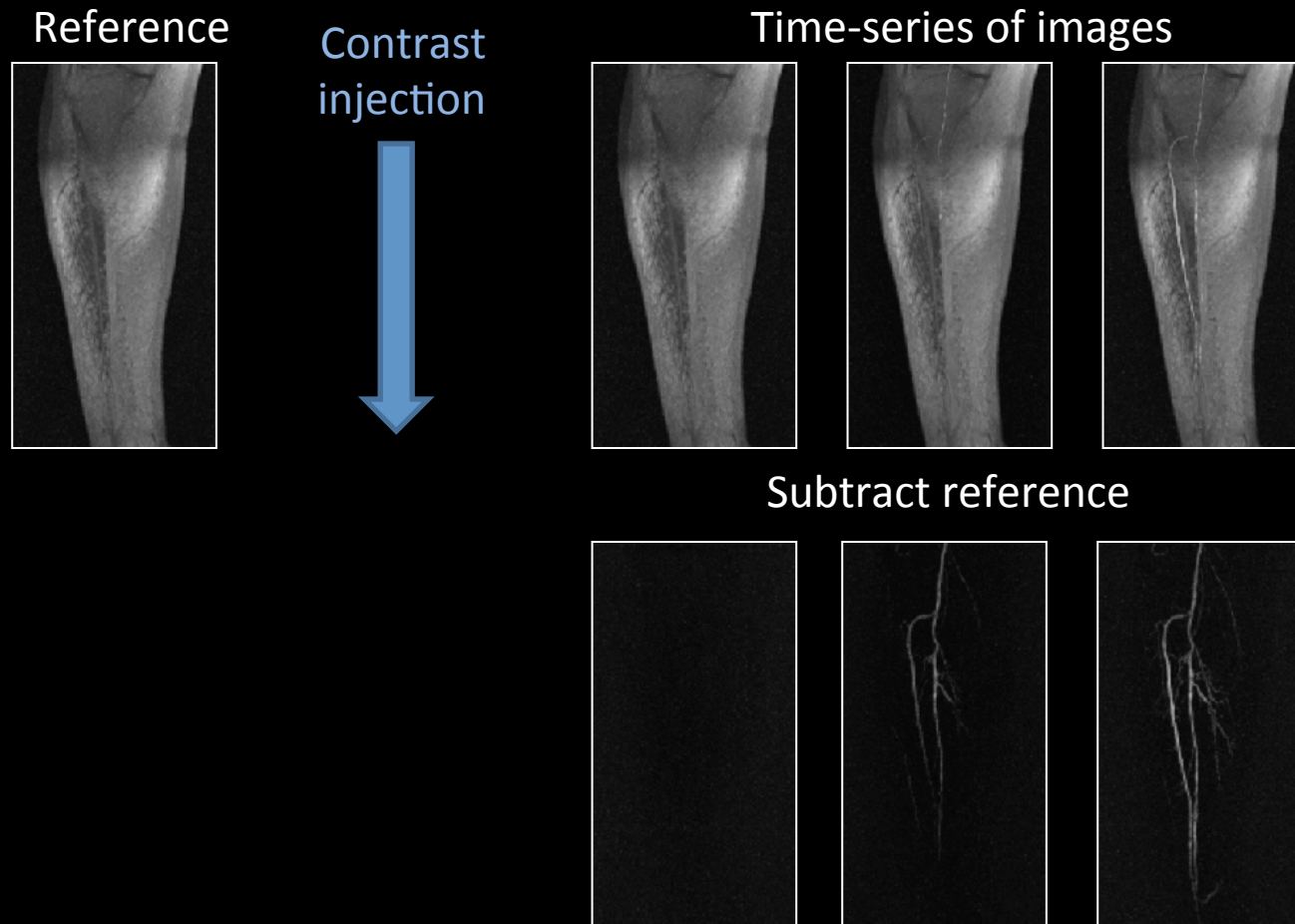
L

S



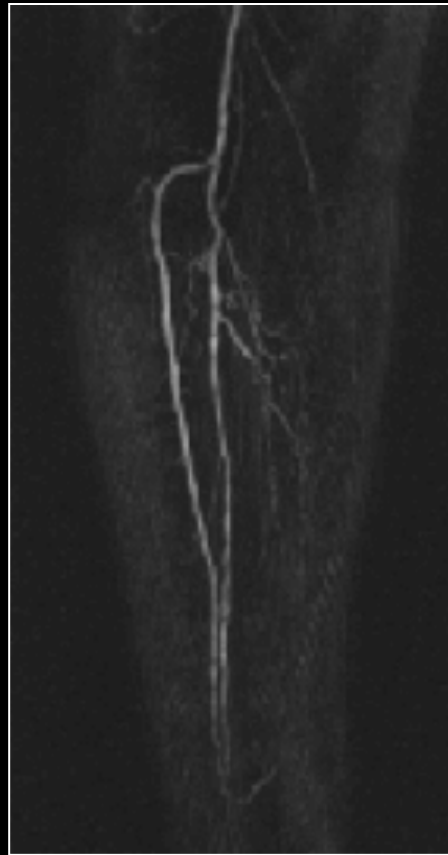
# L+S reconstruction of time-resolved MRA

- It requires background suppression



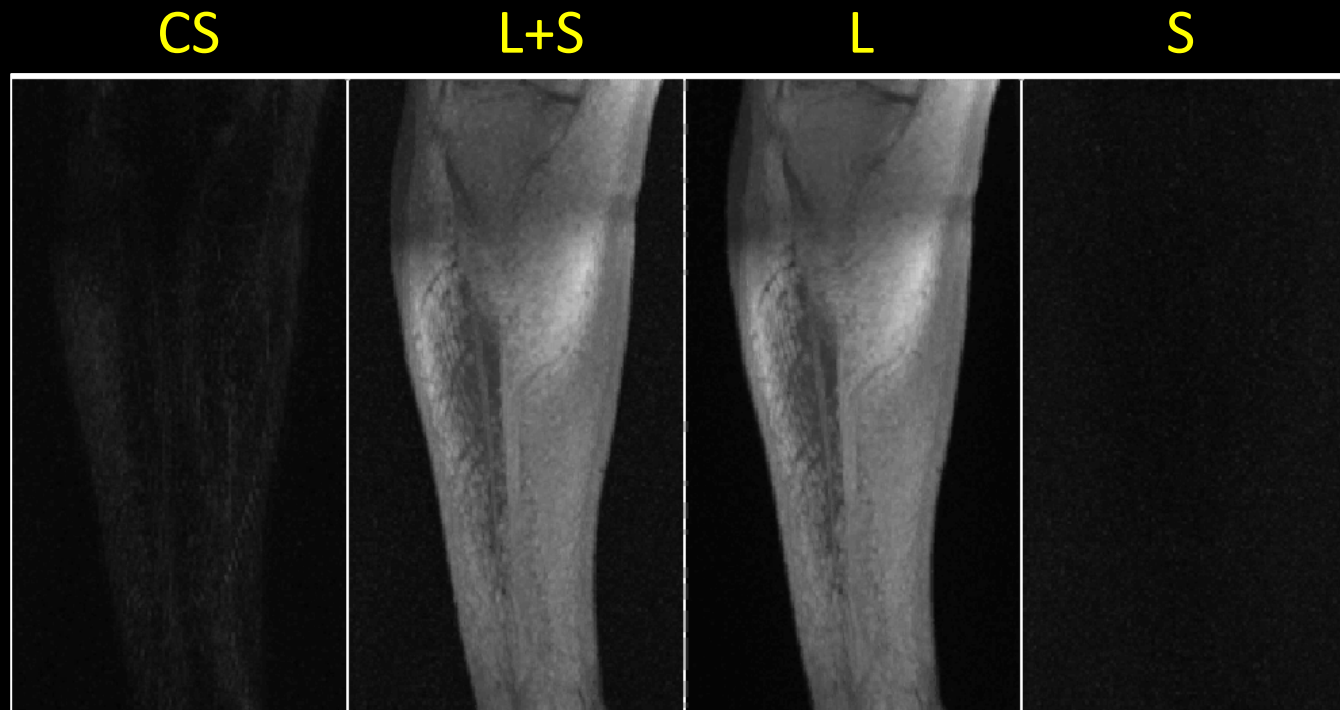
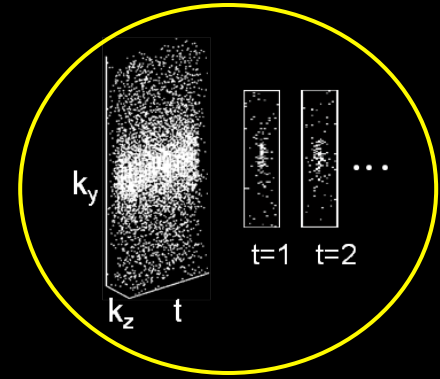
# L+S reconstruction of time-resolved MRA

- Inconsistencies between reference and dynamic data
  - Residual background
  - Incorrect angiogram



# L+S reconstruction of time-resolved MRA

- 7.5-fold acceleration
  - $k_y$ - $k_z$ - $t$  random undersampling
- T: identity (angiograms are already sparse)



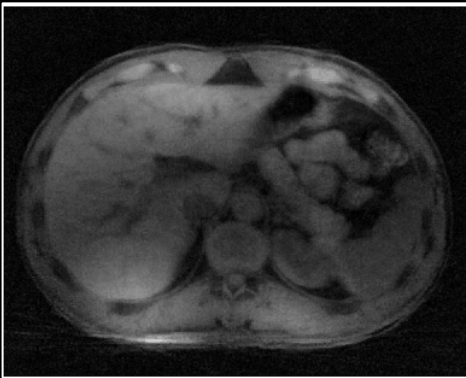
CS uses data subtraction as sparsifying transform

# L+S reconstruction of radial abdominal DCE-MRI

- Continuous golden-angle radial acquisition
- NUFFT (gridding and density compensation)
- Only 8 spokes/temporal frame
  - About 50 fold-acceleration
- T: temporal finite differences



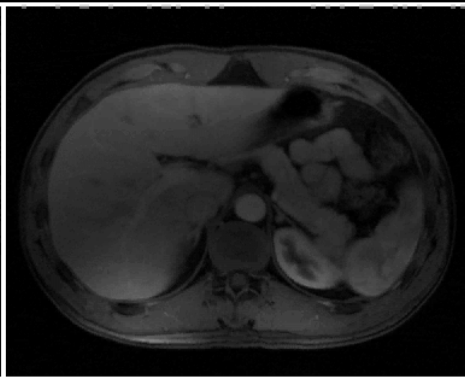
CS



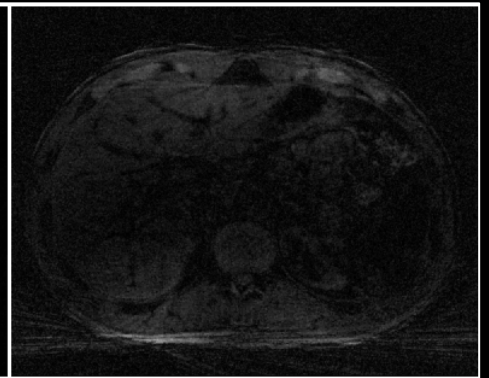
L+S



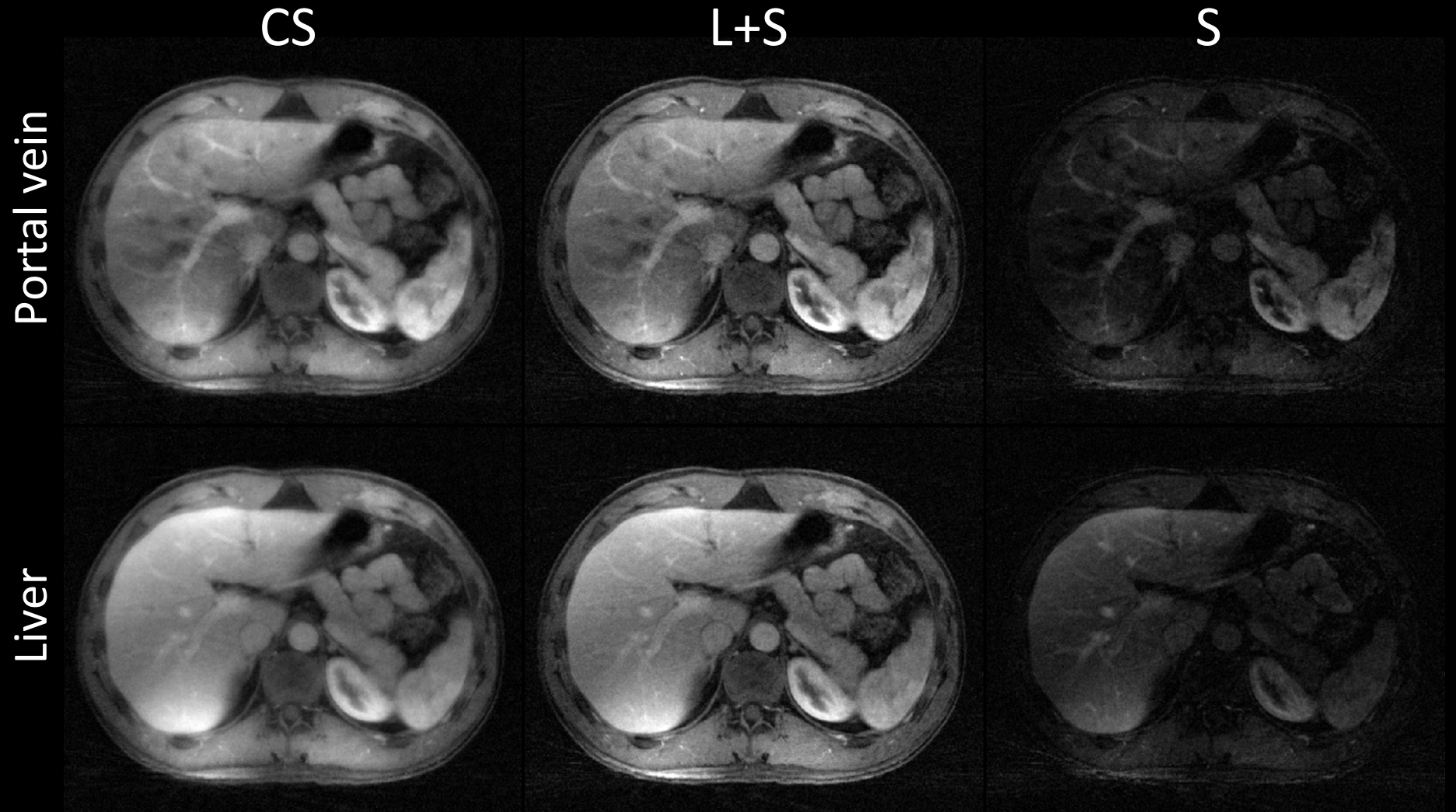
L



S



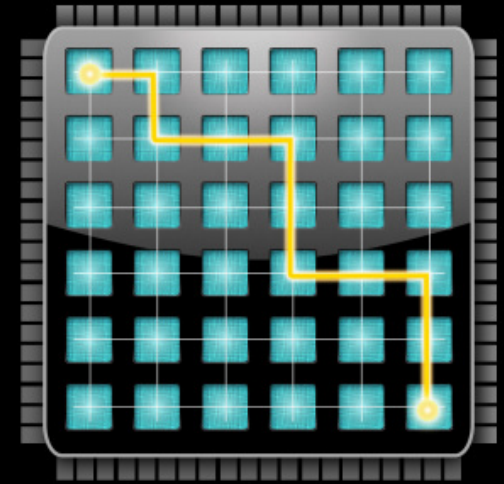
# L+S reconstruction of radial abdominal DCE-MRI





# L+S reconstruction using parallel computing

- Speed up the reconstruction process
- Parallelization
  - Separate reconstruction for each slice
  - Singular value soft-thresholding
  - Fourier transform (time and coil dimensions)
- 96-core computer, reconstructed matrix =  $384 \times 384$ , 40 slices, 12 coils, 48 time points
  - 5 seconds/slice
  - 4D reconstruction under 5 minutes



# Matlab code online

<http://cai2r.net/resources/software/l-s-reconstruction-matlab-code>

The screenshot displays the MATLAB R2014a environment. The main editor window shows a script named 'example1\_cardiac\_perf.m' with the following code:

```
1 % =====  
2 % L+S reconstruction of undersampled multicoil cardiac perfusion MRI  
3 %  
4 % Ricardo Otazo (2013)  
5 % =====  
6 clear all; close all;  
7 % load fully-sampled data  
8 load cardiac_perf_R0.mat;  
9 [nx,ny,nt,nc]=size(kdata);  
10 % L+S reconstruction =====  
11 param.E=mat_xyt(kdata(:,:,1)~=0,b1);  
12 param.d=kdata;  
13 param.T=TempFFT(3);  
14 param.lambda_L=0.01;  
15 param.lambda_S=0.01;  
16 param.nite=50;  
17 param.tol=0.0025;  
18 [L,S] = lps_1st(param);  
19 LplusS=L+S;  
20  
21 % display 4 frames  
22 LplusSd=LplusS(33:96,33:96,2);LplusSd=cat(2,LplusSd,LplusS(33:96,33:96,2));  
23 Ld=L(33:96,33:96,2);Ld=cat(2,Ld,L(33:96,33:96,8));Ld=cat(2,Ld,L(33:96,33:96,8));  
24 Sd=S(33:96,33:96,2);Sd=cat(2,Sd,S(33:96,33:96,8));Sd=cat(2,Sd,S(33:96,33:96,8));  
25  
26 figure;  
27 subplot(3,1,1),imshow(abs(LplusSd),[0,1]);ylabel('L+S')  
28 subplot(3,1,2),imshow(abs(Ld),[0,1]);ylabel('L')  
29 subplot(3,1,3),imshow(abs(Sd),[0,1]);ylabel('S')
```

The Command Window shows the iterative optimization process:

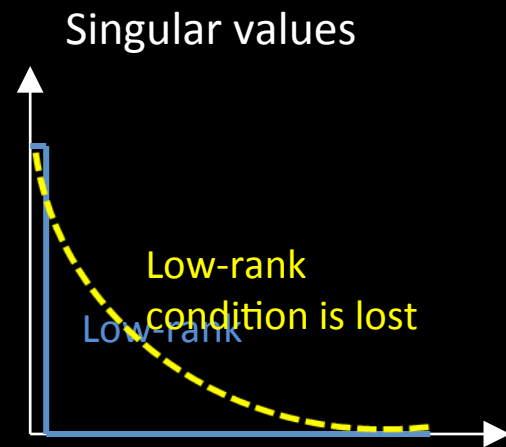
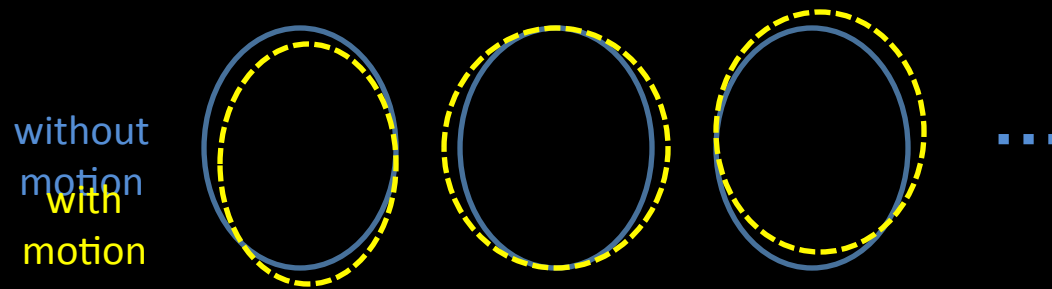
```
ite: 6 , cost: 461.2069263 , update: 0.0356563  
ite: 7 , cost: 453.4043943 , update: 0.0297233  
ite: 8 , cost: 438.2088763 , update: 0.0277863  
ite: 9 , cost: 434.2531293 , update: 0.0241883  
ite: 10 , cost: 426.5837813 , update: 0.0224943  
ite: 11 , cost: 424.6683963 , update: 0.0199863  
ite: 12 , cost: 420.6374183 , update: 0.0184293  
ite: 13 , cost: 419.4549653 , update: 0.0165163  
ite: 14 , cost: 417.2602423 , update: 0.0151253  
ite: 15 , cost: 416.4325233 , update: 0.0135903  
ite: 16 , cost: 414.7147753 , update: 0.0123223  
ite: 17 , cost: 413.8326793 , update: 0.0111493  
ite: 18 , cost: 412.9599133 , update: 0.0102313  
ite: 19 , cost: 412.6501283 , update: 0.0092703  
ite: 20 , cost: 411.9505403 , update: 0.0084843  
ite: 21 , cost: 411.5335613 , update: 0.0077253  
ite: 22 , cost: 410.9702293 , update: 0.0071183  
ite: 23 , cost: 410.8246953 , update: 0.0065523  
ite: 24 , cost: 410.6749213 , update: 0.0060293
```

The Figure window, titled 'Figure 1', displays three rows of MRI slices. The top row is labeled 'L+S', the middle row is labeled 'L', and the bottom row is labeled 'S'. Each row shows four slices of a cardiac MRI scan, demonstrating the reconstruction results.

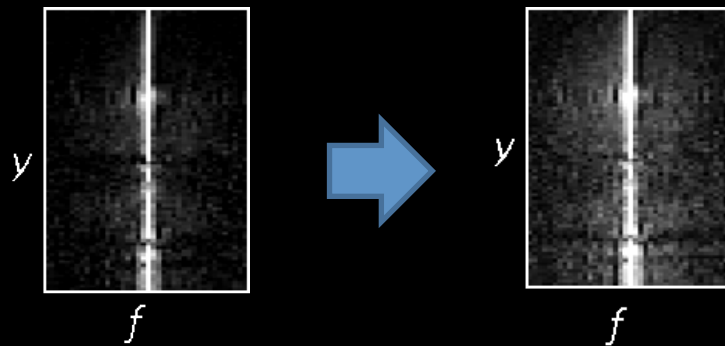
# Motion-guided L+S reconstruction

# Inter-frame motion issues

- Low-rank plus sparse model breaks down



Sparse domain (T)



Motion reduces sparsity

# Inter-frame motion issues

- Motion model into L+S decomposition

$$W = [ W_1 \quad W_2 \quad W_3 \quad \dots ]$$

(frame-by-frame warping operator)

$$L+S = W(M)$$

**low-rank + sparse model is back!**

# Motion-guided L+S reconstruction

- L + S reconstruction with self motion estimation and compensation
  - Find  $L$ ,  $S$  and  $W$  to

$$\begin{aligned} & \text{minimize } \|L\|_* + \lambda \|TS\|_1 \\ & \text{subject to } EW(L + S) = d \end{aligned}$$

---

L: low-rank component

S: sparse component

T: temporal sparsifying transform

W: warping operator

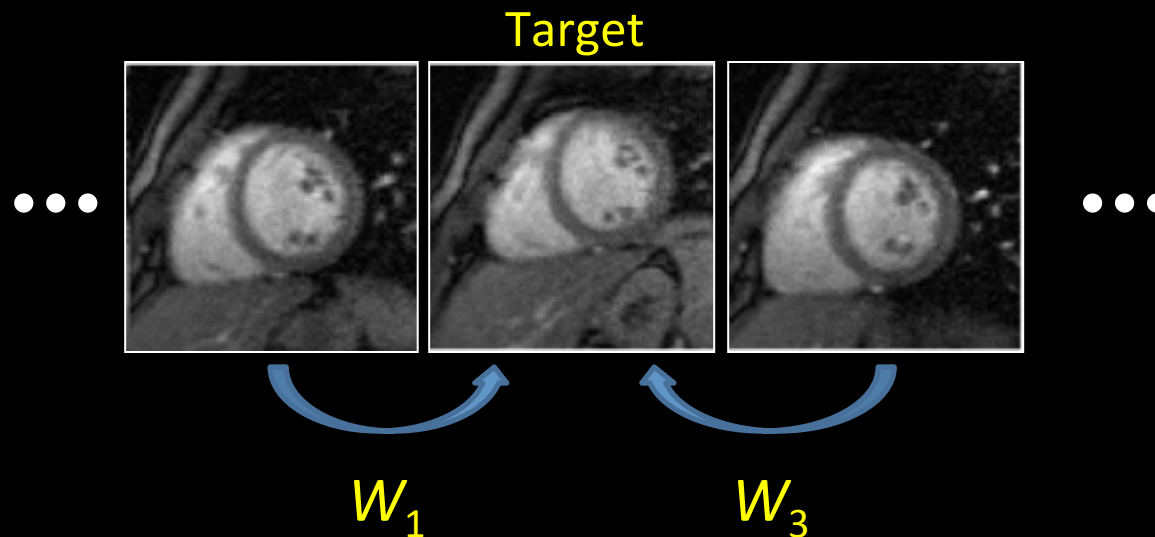
E: encoding operator (Fourier transform + coil sensitivities)

d: undersampled k-t data

# Motion-guided L+S reconstruction

- L+S model as a tool for image-series registration
  - Different from standard image registration

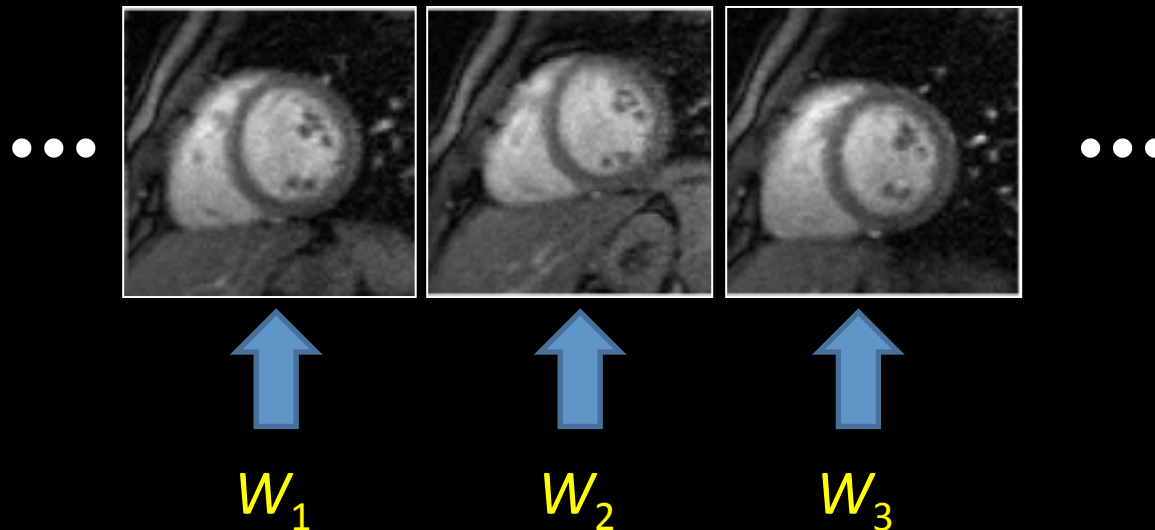
## Standard registration



# Motion-guided L+S reconstruction

- L+S model as a tool for image-series registration
  - Matrix rank-sparsity as a measure of image similarity
  - Does not require similar intensity between frames

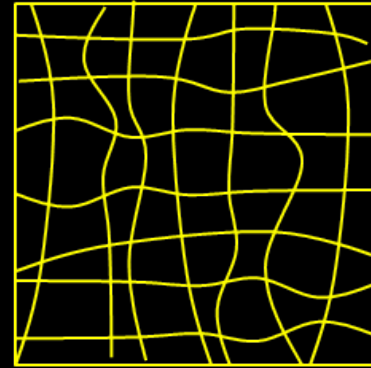
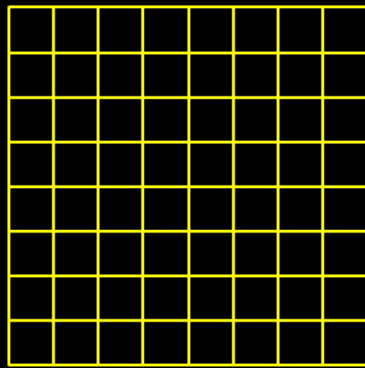
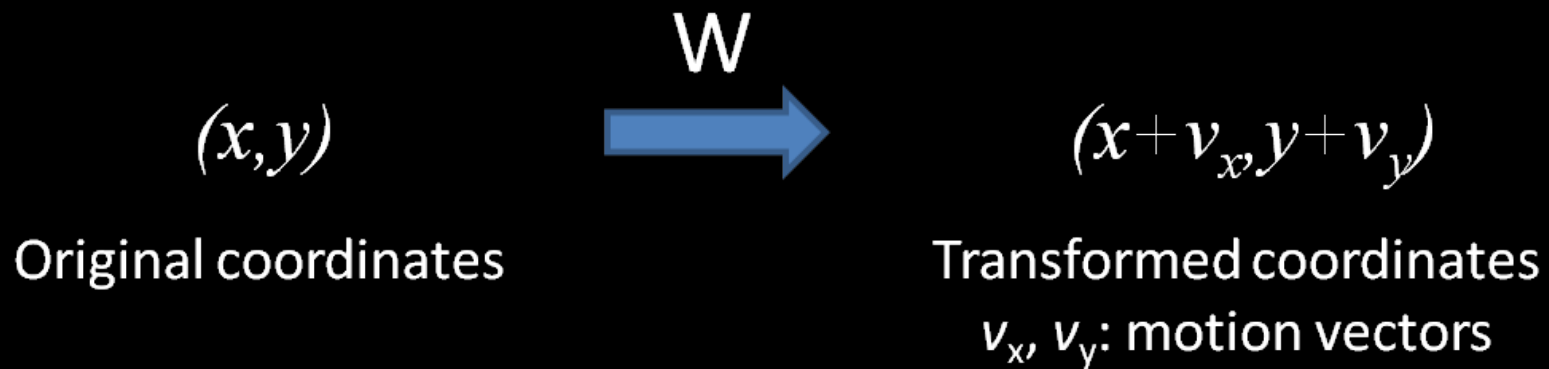
## L+S registration





# Warping operator

- Regridding in the image domain



(non-rigid motion)

# Motion vectors

- Linear approximation of the warping operator
  - Optical flow
  - Pixel-by-pixel motion information

$$W(M) = M + \frac{\partial M}{\partial x} v_x + \frac{\partial M}{\partial y} v_y$$

Gradients of M

Motion vectors

# Motion-guided L+S reconstruction

- L + S reconstruction with self motion estimation and compensation
  - Find  $L$ ,  $S$  and  $v$  to

$$\begin{aligned} & \text{minimize } \|L\|_* + \lambda \|TS\|_1 \\ & \text{subject to } E(L + S + G_{L+S}v) = d \end{aligned}$$

---

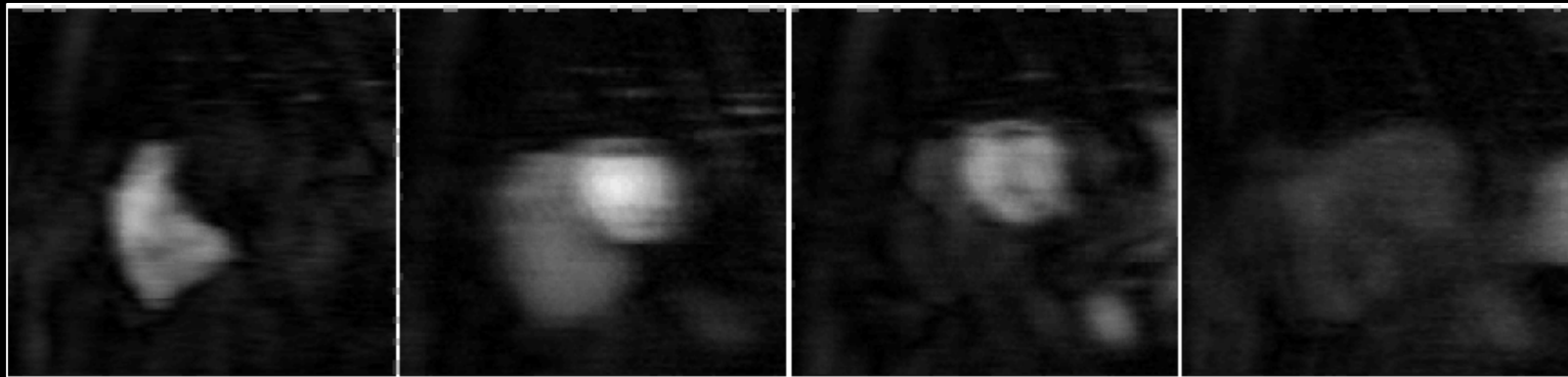
$G_{L+S}$  : gradient of L+S  
 $v$  : motion vectors

# Motion-guided L+S reconstruction

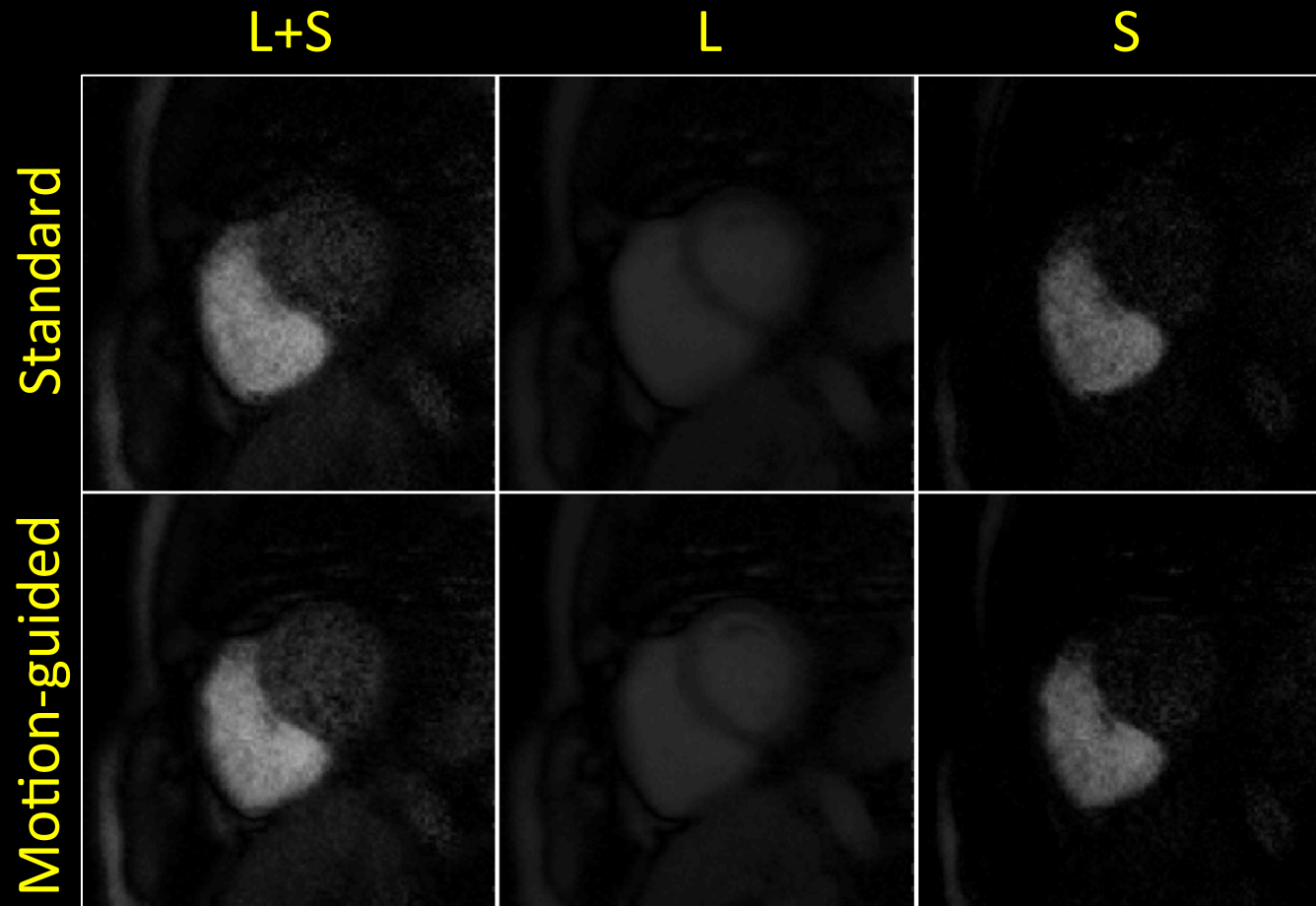
- Initial solution (coil-combined inverse FFT)



- Iterations



# Free-breathing cardiac perfusion



Siemens 3T scanner

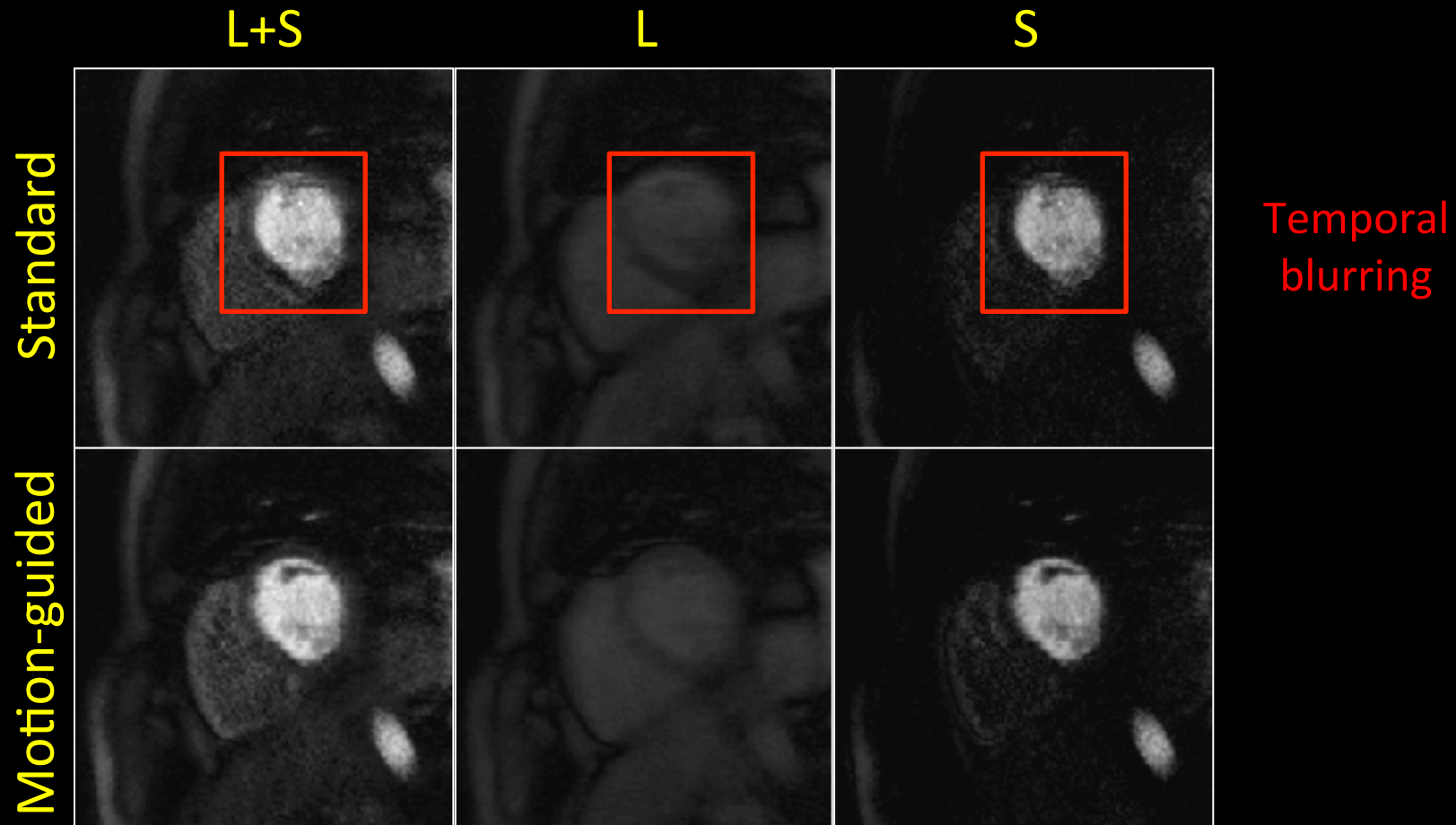
TurboFLASH sequence with 8-fold acceleration ( $k_y$ -t random undersampling)

Temporal resolution = 60ms

Spatial resolution =  $1.7 \times 1.7$  mm<sup>2</sup>

T: temporal FFT

# Free-breathing cardiac perfusion



Siemens 3T scanner

TurboFLASH sequence with 8-fold acceleration ( $k_y$ -t random undersampling)

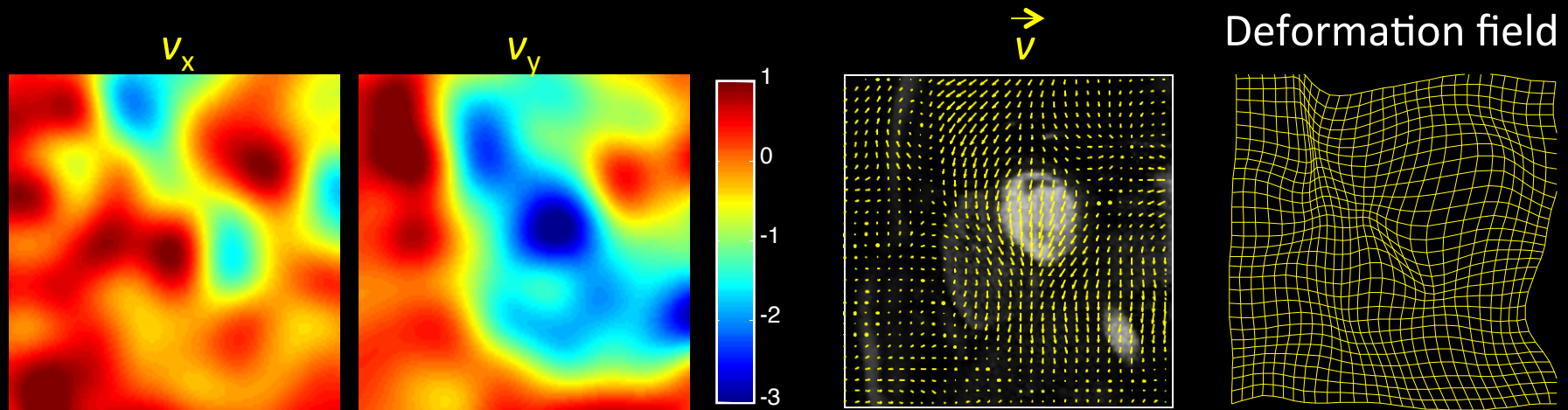
Temporal resolution = 60ms

Spatial resolution =  $1.7 \times 1.7$  mm<sup>2</sup>

T: temporal FFT

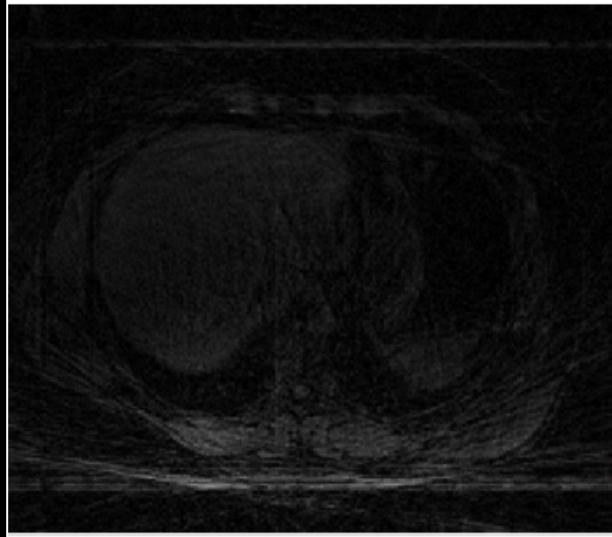
# Free-breathing cardiac perfusion

- Motion vectors
  - non-rigid motion estimation

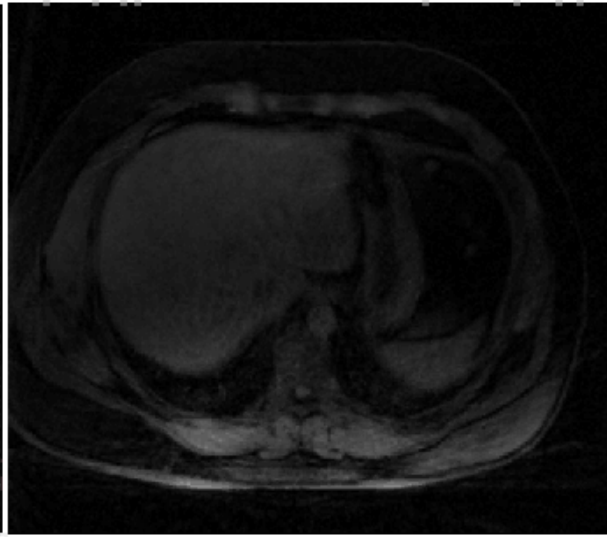


# Free-breathing DCE-MRI of the liver

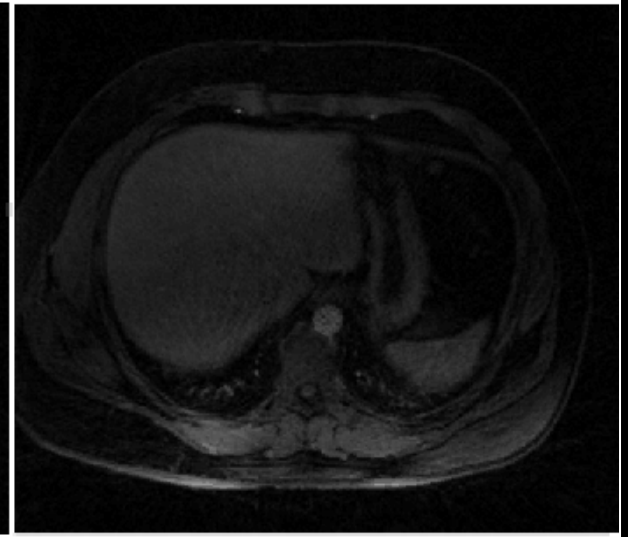
NUFFT



Standard L+S



Motion-guided L+S

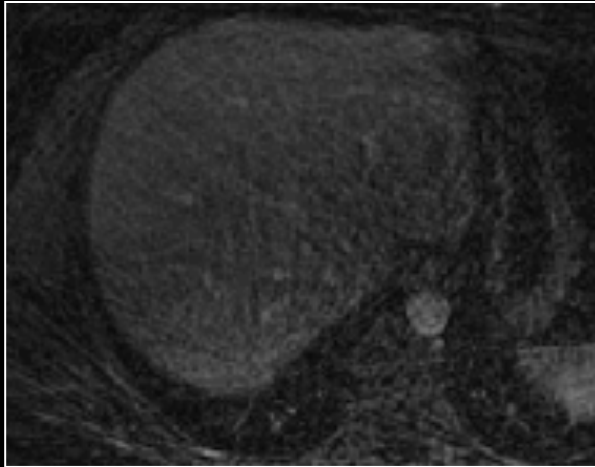


Siemens 3T scanner  
Radial VIBE sequence  
20 spokes/frame (12.8-fold acceleration)  
T: temporal finite differences

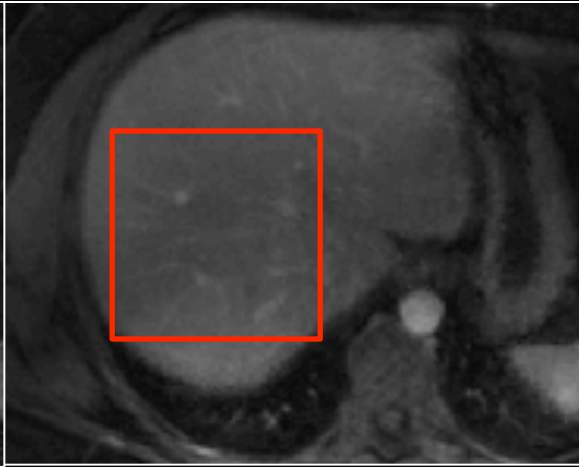


# Free-breathing DCE-MRI of the liver

NUFFT

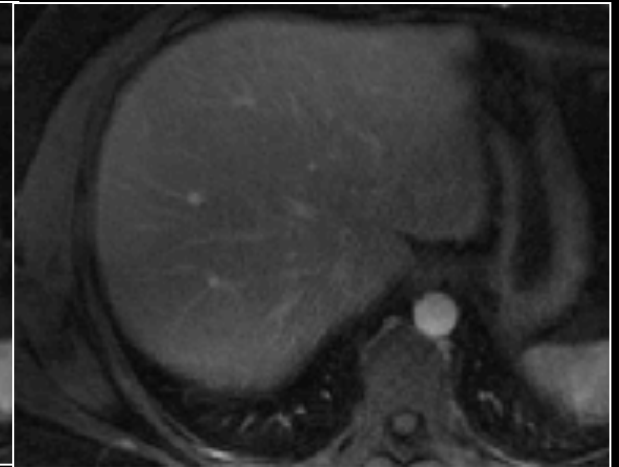


Standard L+S



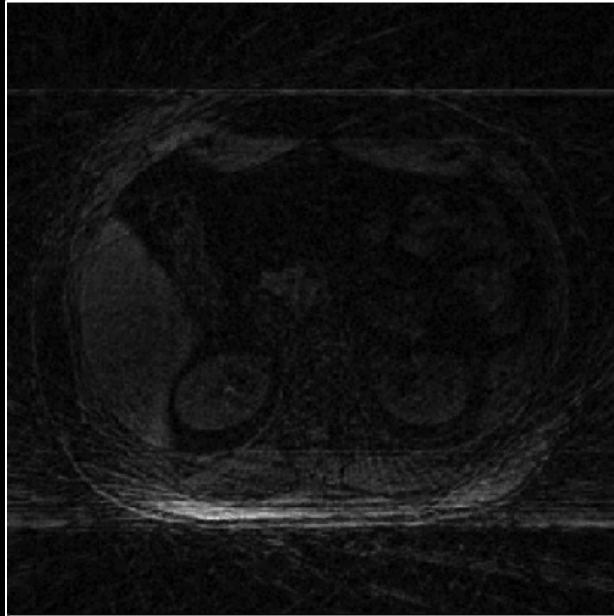
Temporal  
blurring

Motion-guided L+S

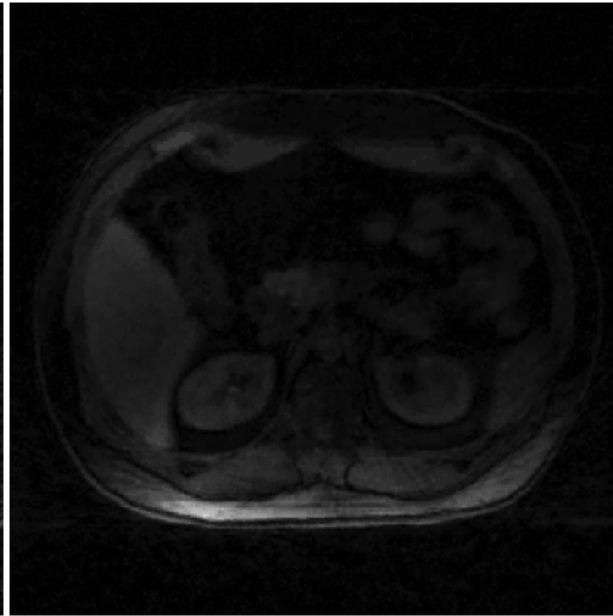


# Free-breathing DCE-MRI of the kidneys

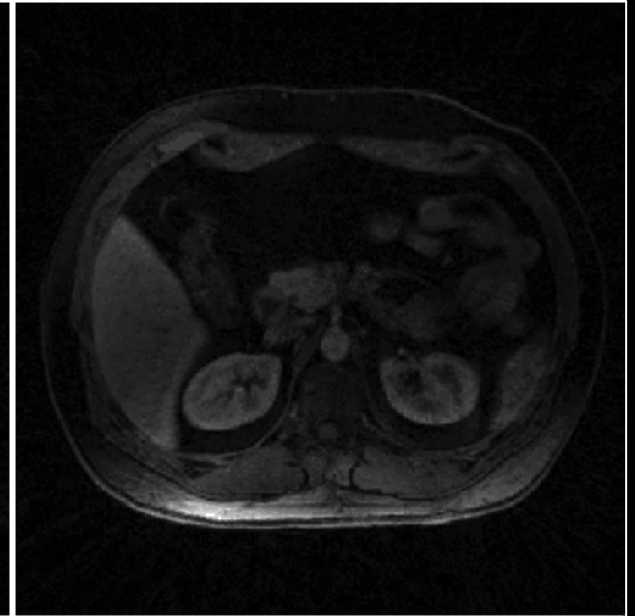
NUFFT



Standard L+S



Motion-guided L+S



Siemens 3T scanner  
Radial VIBE sequence  
20 spokes/frame (12.8-fold acceleration)  
T: temporal finite differences

# Free-breathing DCE-MRI of the kidneys

NUFFT



Standard L+S



Motion-guided L+S



# Summary

- L+S reconstruction
  - Robust PCA for accelerated dynamic MRI
  - Higher performance than standard compressed sensing
  - Background/dynamic separation
- Motion-guided L+S
  - Self-discovery of motion
  - Undersampled data only
- Lots of clinical applications

# Acknowledgments

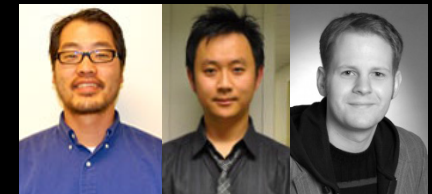
- Emmanuel Candes



- Daniel Sodickson



- Daniel Kim, Li Feng, Tobias Block – data acquisition



- Thomas Koesters – motion models



- Jack Poulson – parallel computing



- Leon Axel and Hersh Chandarana – clinical interpretation





Thank you!