

Improved MRI reconstruction and denoising using SVD-based low-rank approximation



Davi Marco Lyra-Leite
João Paulo C. Lustosa da Costa
João Luiz A. de Carvalho
Universidade de Brasília

davi@ieee.org, jpdacosta@unb.br, joaoluiz@pgea.unb.br




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Summary

- Motivation
- Magnetic Resonance Imaging (MRI)
- SVD
- Problem
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- Methods and Results
- Conclusion
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Motivation

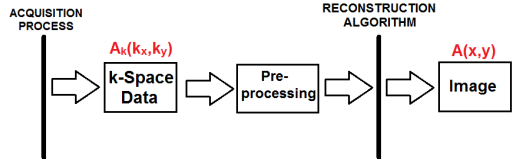
- Reconstruction of multi-dimensional magnetic resonance imaging (MRI) data
 - computationally demanding task
- Therefore, data compression can be applied:
 - to reduce reconstruction complexity and memory requirements,
 - and for denoising.

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Magnetic Resonance Imaging

- Magnetic Resonance Imaging acquisition and reconstruction process



The flowchart shows the process starting with the ACQUISITION PROCESS, which produces $A_x(k_x, k_y)$ (k-Space Data). This data then goes through Pre-processing, and finally the RECONSTRUCTION ALGORITHM, which produces the final Image $A(x, y)$.

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Magnetic Resonance Imaging


- The Fourier Transform of the image

$$A(k_x, k_y) = \int_x \int_y A(x, y) e^{-j2\pi(k_x x + k_y y)} dx dy$$
- The Fourier coordinates

$$k_x(t) = \frac{\gamma}{2\pi} \int_0^t G_x(\tau) d\tau$$

$$k_y(t) = \frac{\gamma}{2\pi} \int_0^t G_y(\tau) d\tau$$

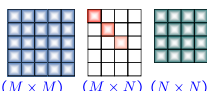
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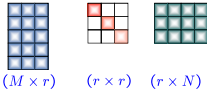
Singular Value Decomposition

$X \in \mathbb{C}^{M \times N}$, $\text{rank}(X) = r$

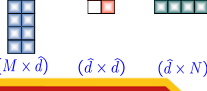
"Full SVD"
 $X = U \cdot \Sigma \cdot V^H$




"Economy size SVD"
 $X = U_s \cdot \Sigma_s \cdot V_s^H$



Low-rank approximation
 $X \approx U_s \cdot \Sigma_s \cdot V_s^H$



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Singular Value Decomposition

- Decomposition into a sum of rank one matrices:

$$X = U \cdot \Sigma \cdot V^H$$

$(M \times N) = (M \times r) \cdot (r \times r) \cdot (r \times N)$
 $= \sum_{n=1}^r \sigma_n \cdot \mathbf{u}_n \cdot \mathbf{v}_n^H$

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Proposal

- Apply a low-rank approximation singular value decomposition to MRI data in order to:
 - remove noise components
 - verified through the Signal to Error Ratio
 - reduce the storage memory
 - with a high quality level

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Akaike Information Criterion

- Akaike Information Criterion
 - based on the difference between the maximum likelihood function and the free parameters.
- We shall find m that minimizes:

$$AIC(m) = -\ln(\mathcal{L}(m)) + \kappa(m)$$
 where
 - $\ln(\mathcal{L}(m))$ is the loglikelihood function.
 - $\kappa(m)$ are the free parameters.

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Akaike Information Criterion

- The Loglikelihood as function of the eigenvalues proposed by Wax and Kailath.

Eigenvalues profile

$d = 3$

$AIC(m) = -\ln(\mathcal{L}(m)) + \kappa(m)$

$\hat{d} + 1 = 4$

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Results

Original ground-truth image

Denoised in Image Domain via AIC/SVD

$\hat{d} = 30$

Denoised in Frequency Domain via AIC/SVD

$\hat{d} = 30$

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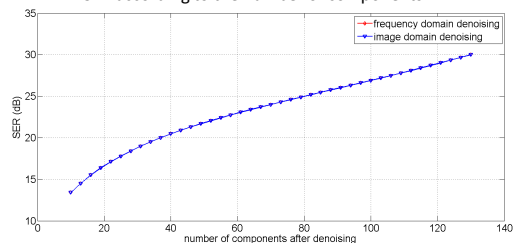
Results

SVD profile in frequency domain

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Results

SER according to the number of components



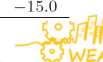
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Magnetic Resonance Imaging

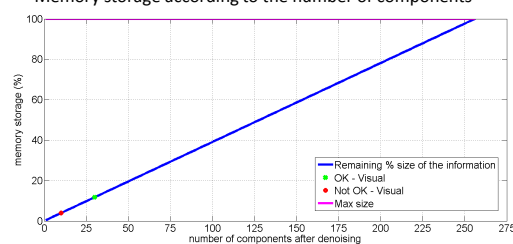
Number of Components	SER (dB)		RMSE (dB)
	frequency domain	image domain	
10	13.4	13.4	-6.7
22	17.1	17.1	-8.6
31	19.0	19.0	-9.5
43	20.9	20.9	-10.5
55	22.4	22.4	-11.2
67	23.7	23.7	-11.9
79	24.9	24.9	-12.5
88	25.8	25.8	-12.9
97	26.7	26.7	-13.3
109	27.9	27.9	-13.9
118	28.8	28.8	-14.4
124	29.4	29.4	-14.7
130	30.1	30.1	-15.0

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Results

Memory storage according to the number of components



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Conclusion

- We propose a SVD-based low rank approximation for MRI data.
- Our approach:
 - can be applied for denoising;
 - can reduce significantly the memory storage;
 - can be applied in image and frequency domain.

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Future Works

- Extension for the low-rank approximation HOSVD
 - The MRI data is usually three dimensional
 - two dimensions related to image and one related to time

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Thank you !

<http://www.pgea.unb.br/~lasp/>

jpdacosta@unb.br

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