

Relaxed linearized algorithms for faster X-ray CT image reconstruction

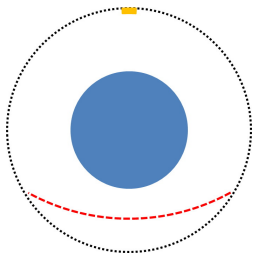
Hung Nien and Jeffrey A. Fessler



University of Michigan, Ann Arbor

The 13th Fully 3D Meeting – June 2, 2015

Statistical image reconstruction for X-ray CT

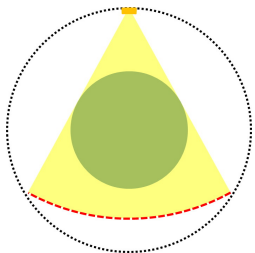


$$\mathbf{y} = \mathbf{A} \mathbf{x} + \boldsymbol{\varepsilon}$$

The diagram illustrates the linear model for X-ray CT reconstruction. The vector \mathbf{y} (red squares) represents the measured data. The matrix \mathbf{A} (yellow squares) represents the system matrix. The vector \mathbf{x} (blue squares) represents the unknown image. The vector $\boldsymbol{\varepsilon}$ (green squares) represents the noise. The equation is shown as $\mathbf{y} = \mathbf{A} \mathbf{x} + \boldsymbol{\varepsilon}$.



Statistical image reconstruction for X-ray CT

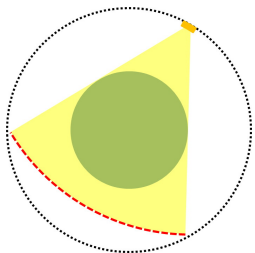


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Statistical image reconstruction for X-ray CT

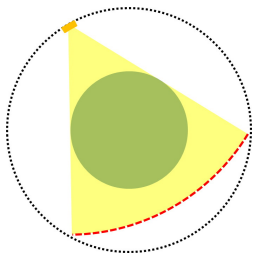


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The diagram shows the linear model for X-ray CT reconstruction. \mathbf{y} is a vertical vector of red squares, \mathbf{A} is a matrix of yellow squares, \mathbf{x} is a vertical vector of blue squares, and $\boldsymbol{\varepsilon}$ is a vertical vector of green squares. The matrix \mathbf{A} is sparse, with a gap in the middle indicated by vertical dots.



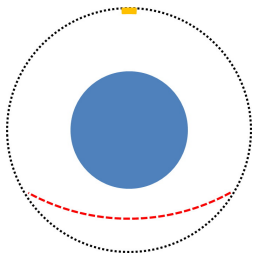
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Statistical image reconstruction for X-ray CT

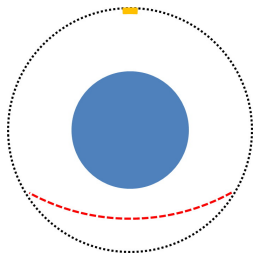


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Statistical image reconstruction for X-ray CT



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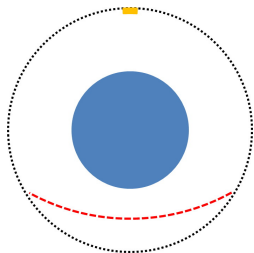
The diagram illustrates the linear model for X-ray CT reconstruction. On the left, a vertical vector \mathbf{y} is shown with red squares. This is equal to the product of a matrix \mathbf{A} (a grid of yellow squares) and a vector \mathbf{x} (a vertical vector of blue squares). The result is then added to a noise vector $\boldsymbol{\varepsilon}$ (a vertical vector of green squares).

Statistical image reconstruction (SIR)

$$\hat{\mathbf{x}} \in \arg \min_{\mathbf{x}} \left\{ \Psi_{\text{PWLS}}(\mathbf{x}) \triangleq \frac{1}{2} \|\mathbf{y} - \mathbf{A}\mathbf{x}\|_{\mathbf{W}}^2 + R(\mathbf{x}) + \iota_{\Omega}(\mathbf{x}) \right\}$$



Statistical image reconstruction for X-ray CT



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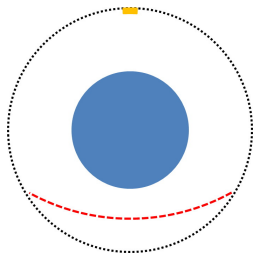
The diagram illustrates the linear model for X-ray CT reconstruction. \mathbf{y} is a vertical vector of red squares representing the measured projection data. \mathbf{A} is a large yellow grid representing the system matrix. \mathbf{x} is a vertical vector of blue squares representing the unknown image pixels. $\boldsymbol{\varepsilon}$ is a vertical vector of green squares representing the noise. The equation shows that the measured data \mathbf{y} is equal to the product of the system matrix \mathbf{A} and the image \mathbf{x} , plus the noise $\boldsymbol{\varepsilon}$.

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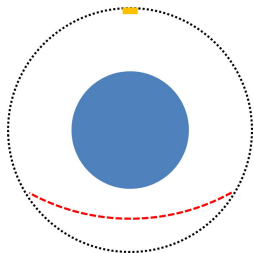
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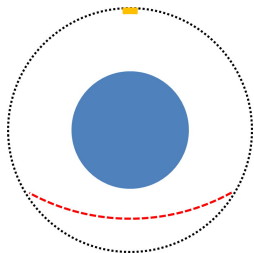
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First-order algorithms with ordered-subsets

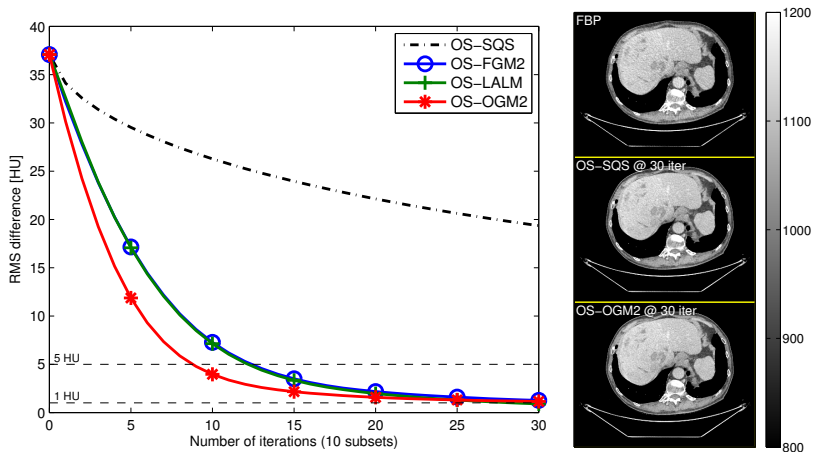


Figure: Chest: Existing first-order algorithms with ordered-subsets (OS).

First-order algorithms with ordered-subsets

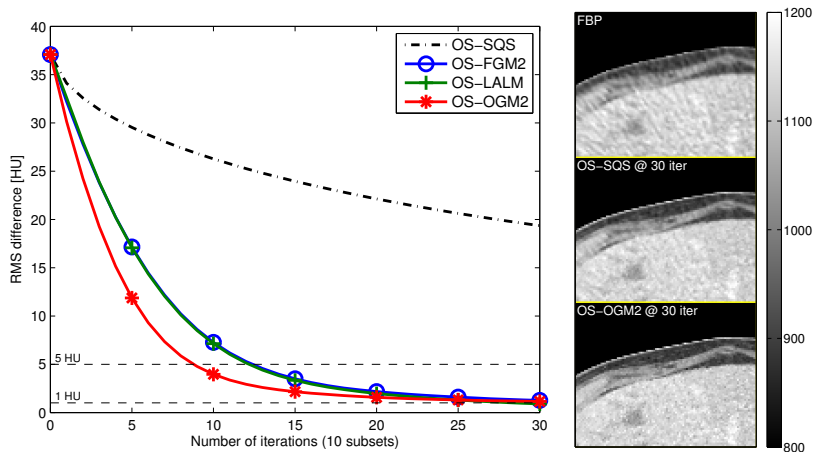


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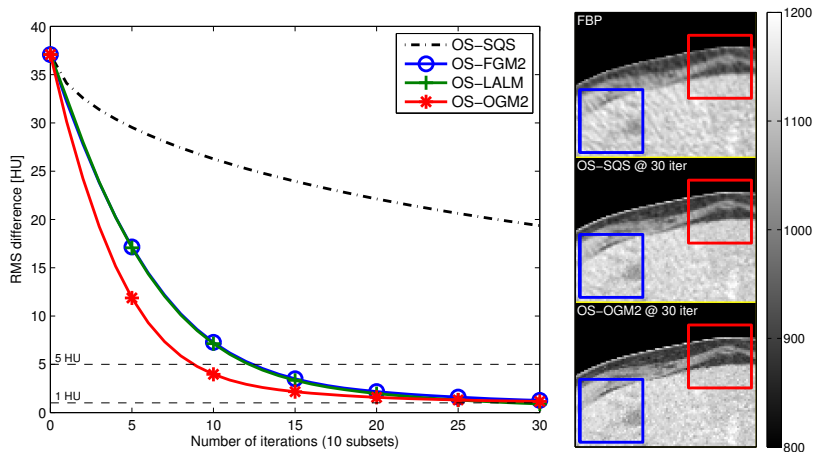


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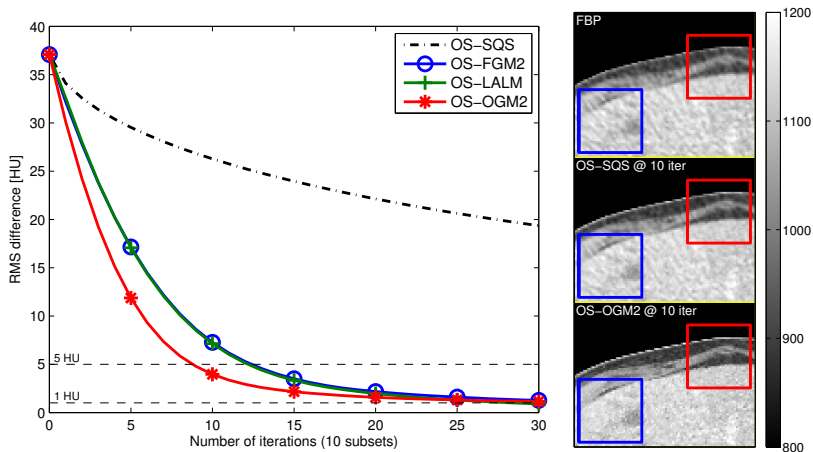


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First-order algorithms with ordered-subsets

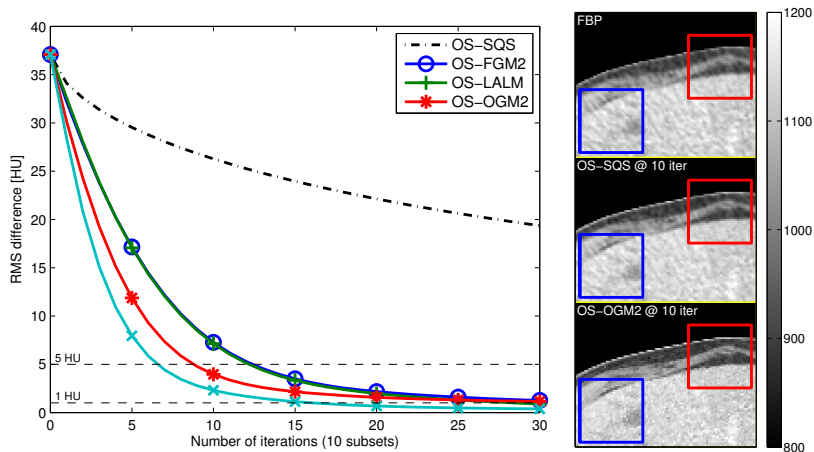


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Equality-constrained composite minimization

Consider an equality-constrained minimization problem:

$$(\hat{\mathbf{x}}, \hat{\mathbf{u}}) \in \arg \min_{\mathbf{x}, \mathbf{u}} \left\{ \frac{1}{2} \|\mathbf{u} - \mathbf{y}\|_2^2 + h(\mathbf{x}) \right\} \text{ s.t. } \mathbf{u} = \mathbf{Ax}, \quad (1)$$

where h are closed and proper convex functions.

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In particular, the **quadratic loss function** penalizes the difference between the linear model \mathbf{Ax} and noisy measurement \mathbf{y} , and h is a **regularization term** that introduces the prior knowledge of \mathbf{x} to the reconstruction. For example,

$$(\hat{\mathbf{x}}, \hat{\mathbf{u}}) \in \arg \min_{\mathbf{x}, \mathbf{u}} \left\{ \frac{1}{2} \|\mathbf{u} - \mathbf{W}^{1/2} \mathbf{y}\|_2^2 + (R + \iota_{\Omega})(\mathbf{x}) \right\} \text{ s.t. } \mathbf{u} = \mathbf{W}^{1/2} \mathbf{Ax}$$

represents an **X-ray CT image reconstruction** problem.

Standard AL method

The standard AL method finds a saddle-point of the augmented Lagrangian (AL) of (1):

$$\mathcal{L}_A(\mathbf{x}, \mathbf{u}, \mathbf{d}; \rho) \triangleq \frac{1}{2} \|\mathbf{u} - \mathbf{y}\|_2^2 + h(\mathbf{x}) + \frac{\rho}{2} \|\mathbf{Ax} - \mathbf{u} - \mathbf{d}\|_2^2$$

in an alternating direction manner:

$$\begin{cases} \mathbf{x}^{(k+1)} \in \arg \min_{\mathbf{x}} \left\{ h(\mathbf{x}) + \frac{\rho}{2} \|\mathbf{Ax} - \mathbf{u}^{(k)} - \mathbf{d}^{(k)}\|_2^2 \right\} \\ \mathbf{u}^{(k+1)} \in \arg \min_{\mathbf{u}} \left\{ \frac{1}{2} \|\mathbf{u} - \mathbf{y}\|_2^2 + \frac{\rho}{2} \|\mathbf{Ax}^{(k+1)} - \mathbf{u} - \mathbf{d}^{(k)}\|_2^2 \right\} \\ \mathbf{d}^{(k+1)} = \mathbf{d}^{(k)} - \mathbf{Ax}^{(k+1)} + \mathbf{u}^{(k+1)}, \end{cases}$$

where \mathbf{d} is the scaled Lagrange multiplier of \mathbf{u} , and $\rho > 0$ is the AL penalty parameter.

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The linearized AL method adds an additional **G-proximity term** to the **x**-subproblem in the **standard AL method**:

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where $\mathbf{G} \triangleq \mathbf{D}_L - \mathbf{A}'\mathbf{A}$, and \mathbf{D}_L is a diagonal majorizing matrix of $\mathbf{A}'\mathbf{A}$.

[Zhang et al., J. Sci. Comput., 2011]

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where $\mathbf{G} \triangleq \mathbf{D}_L - \mathbf{A}'\mathbf{A}$, and \mathbf{D}_L is a diagonal majorizing matrix of $\mathbf{A}'\mathbf{A}$. Here, we **linearize** the algorithm in a sense that the Hessian matrix of the “augmented” quadratic AL term is **diagonal**.

[Zhang et al., J. Sci. Comput., 2011]

[HN and Fessler, IEEE Trans. Med. Imag., 2015]

Linearized AL algorithm for X-ray CT

Algorithm: OS-LALM for CT reconstruction.

Input: $K \geq 1$, $M \geq 1$, and an initial (FBP) image \mathbf{x} .

set $\rho = 1$, $\zeta = \mathbf{g} = M\nabla L_M(\mathbf{x})$

for $k = 1, 2, \dots, K$ **do**

for $m = 1, 2, \dots, M$ **do**

$\mathbf{s} = \rho\zeta + (1 - \rho)\mathbf{g}$

$\mathbf{x}^+ = [\mathbf{x} - (\rho\mathbf{D}_L + \mathbf{D}_R)^{-1}(\mathbf{s} + \nabla R(\mathbf{x}))]_{\Omega}$

$\zeta^+ = M\nabla L_m(\mathbf{x}^+)$

$\mathbf{g}^+ = \frac{\rho}{\rho+1}\zeta^+ + \frac{1}{\rho+1}\mathbf{g}$

 decrease ρ gradually

end

end

Output: The final image \mathbf{x} .

Relaxed AL method

The relaxed AL method accelerates the **standard AL method** with **under- or over-relaxation**:

$$\begin{cases} \mathbf{x}^{(k+1)} \in \arg \min_{\mathbf{x}} \left\{ h(\mathbf{x}) + \frac{\rho}{2} \|\mathbf{Ax} - \mathbf{u}^{(k)} - \mathbf{d}^{(k)}\|_2^2 \right\} \\ \mathbf{u}^{(k+1)} \in \arg \min_{\mathbf{u}} \left\{ \frac{1}{2} \|\mathbf{u} - \mathbf{y}\|_2^2 + \frac{\rho}{2} \|\mathbf{r}_{\mathbf{u},\alpha}^{(k+1)} - \mathbf{u} - \mathbf{d}^{(k)}\|_2^2 \right\} \\ \mathbf{d}^{(k+1)} = \mathbf{d}^{(k)} - \mathbf{r}_{\mathbf{u},\alpha}^{(k+1)} + \mathbf{u}^{(k+1)}, \end{cases}$$

where

$$\mathbf{r}_{\mathbf{u},\alpha}^{(k+1)} \triangleq \alpha \mathbf{Ax}^{(k+1)} + (1 - \alpha) \mathbf{u}^{(k)}$$

is the relaxation variable of \mathbf{u} , and $0 < \alpha < 2$ is the relaxation parameter.

[Eckstein and Bertsekas, Math. Prog., 1992]

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is the relaxation variable of \mathbf{u} , and $0 < \alpha < 2$ is the relaxation parameter. When $\alpha = 1$, it reverts to the **standard AL method**.

[Eckstein and Bertsekas, Math. Prog., 1992]

Implicit linearization via redundant variable-splitting

Consider an equivalent equality-constrained minimization problem with a **redundant equality constraint**:

$$(\hat{\mathbf{x}}, \hat{\mathbf{u}}, \hat{\mathbf{v}}) \in \arg \min_{\mathbf{x}, \mathbf{u}, \mathbf{v}} \left\{ \frac{1}{2} \|\mathbf{u} - \mathbf{y}\|_2^2 + h(\mathbf{x}) \right\} \text{ s.t. } \begin{cases} \mathbf{u} = \mathbf{A}\mathbf{x} \\ \mathbf{v} = \mathbf{G}^{1/2}\mathbf{x}. \end{cases}$$

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Suppose we use the same AL penalty parameter ρ for both equality constraints in AL methods. The quadratic AL term

$$\frac{\rho}{2} \|\mathbf{A}\mathbf{x} - \mathbf{u}^{(k)} - \mathbf{d}^{(k)}\|_2^2 + \frac{\rho}{2} \|\mathbf{G}^{1/2}\mathbf{x} - \mathbf{v}^{(k)} - \mathbf{e}^{(k)}\|_2^2$$

has a **diagonal** Hessian matrix $\mathbf{H}_\rho \triangleq \rho\mathbf{A}'\mathbf{A} + \rho\mathbf{G} = \rho\mathbf{D}_L$, leading to a **separable** quadratic AL term.

[HN and Fessler, IEEE Trans. Med. Imag., 2015]

Proposed relaxed linearized AL method

The proposed relaxed linearized AL method solves the equivalent minimization problem with a **redundant equality constraint** using the **relaxed AL method**:

$$\begin{cases} \mathbf{x}^{(k+1)} \in \arg \min_{\mathbf{x}} \left\{ h(\mathbf{x}) + \frac{\rho}{2} \|\mathbf{Ax} - \mathbf{u}^{(k)} - \mathbf{d}^{(k)}\|_2^2 \right. \\ \quad \left. + \frac{\rho}{2} \|\mathbf{G}^{1/2}\mathbf{x} - \mathbf{v}^{(k)} - \mathbf{e}^{(k)}\|_2^2 \right\} \\ \mathbf{u}^{(k+1)} \in \arg \min_{\mathbf{u}} \left\{ \frac{1}{2} \|\mathbf{u} - \mathbf{y}\|_2^2 + \frac{\rho}{2} \|\mathbf{r}_{\mathbf{u},\alpha}^{(k+1)} - \mathbf{u} - \mathbf{d}^{(k)}\|_2^2 \right\} \\ \mathbf{d}^{(k+1)} = \mathbf{d}^{(k)} - \mathbf{r}_{\mathbf{u},\alpha}^{(k+1)} + \mathbf{u}^{(k+1)} \\ \mathbf{v}^{(k+1)} = \mathbf{r}_{\mathbf{v},\alpha}^{(k+1)} - \mathbf{e}^{(k)} \\ \mathbf{e}^{(k+1)} = \mathbf{e}^{(k)} - \mathbf{r}_{\mathbf{v},\alpha}^{(k+1)} + \mathbf{v}^{(k+1)}. \end{cases}$$

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When $\alpha = 1$, the proposed method reverts to the **linearized AL method**.

Proposed relaxed linearized AL method (cont'd)

The proposed relaxed linearized AL method further simplifies as follows:

$$\left\{ \begin{array}{l} \boldsymbol{\gamma}^{(k+1)} = (\rho - 1) \mathbf{g}^{(k)} + \rho \mathbf{h}^{(k)} \\ \mathbf{x}^{(k+1)} \in \arg \min_{\mathbf{x}} \left\{ h(\mathbf{x}) + \frac{1}{2} \|\mathbf{x} - (\rho \mathbf{D}_L)^{-1} \boldsymbol{\gamma}^{(k+1)}\|_{\rho \mathbf{D}_L}^2 \right\} \\ \boldsymbol{\zeta}^{(k+1)} = \nabla L(\mathbf{x}^{(k+1)}) \triangleq \mathbf{A}' (\mathbf{A} \mathbf{x}^{(k+1)} - \mathbf{y}) \\ \mathbf{g}^{(k+1)} = \frac{\rho}{\rho+1} (\alpha \boldsymbol{\zeta}^{(k+1)} + (1 - \alpha) \mathbf{g}^{(k)}) + \frac{1}{\rho+1} \mathbf{g}^{(k)} \\ \mathbf{h}^{(k+1)} = \alpha (\mathbf{D}_L \mathbf{x}^{(k+1)} - \boldsymbol{\zeta}^{(k+1)}) + (1 - \alpha) \mathbf{h}^{(k)}, \end{array} \right.$$

where $L(\mathbf{x}) \triangleq (1/2) \|\mathbf{A} \mathbf{x} - \mathbf{y}\|_2^2$ denotes the **quadratic data-fidelity term**, and $\mathbf{g}^{(k)} \triangleq \mathbf{A}' (\mathbf{u}^{(k)} - \mathbf{y})$ denotes the **split gradient** of L .

Proposed relaxed linearized AL method (cont'd)

The proposed relaxed linearized AL method further simplifies as follows:

$$\left\{ \begin{array}{l} \boldsymbol{\gamma}^{(k+1)} = (\rho - 1) \mathbf{g}^{(k)} + \rho \mathbf{h}^{(k)} \\ \mathbf{x}^{(k+1)} \in \arg \min_{\mathbf{x}} \left\{ h(\mathbf{x}) + \frac{1}{2} \|\mathbf{x} - (\rho \mathbf{D}_L)^{-1} \boldsymbol{\gamma}^{(k+1)}\|_{\rho \mathbf{D}_L}^2 \right\} \\ \boldsymbol{\zeta}^{(k+1)} = \nabla L(\mathbf{x}^{(k+1)}) \triangleq \mathbf{A}' (\mathbf{A} \mathbf{x}^{(k+1)} - \mathbf{y}) \\ \mathbf{g}^{(k+1)} = \frac{\rho}{\rho+1} (\alpha \boldsymbol{\zeta}^{(k+1)} + (1 - \alpha) \mathbf{g}^{(k)}) + \frac{1}{\rho+1} \mathbf{g}^{(k)} \\ \mathbf{h}^{(k+1)} = \alpha (\mathbf{D}_L \mathbf{x}^{(k+1)} - \boldsymbol{\zeta}^{(k+1)}) + (1 - \alpha) \mathbf{h}^{(k)}, \end{array} \right.$$

where $L(\mathbf{x}) \triangleq (1/2) \|\mathbf{A} \mathbf{x} - \mathbf{y}\|_2^2$ denotes the **quadratic data-fidelity term**, and $\mathbf{g}^{(k)} \triangleq \mathbf{A}' (\mathbf{u}^{(k)} - \mathbf{y})$ denotes the **split gradient** of L .

Relaxed OS-LALM for faster CT reconstruction

To solve X-ray CT image reconstruction problem:

$$\hat{\mathbf{x}} \in \arg \min_{\mathbf{x}} \left\{ \frac{1}{2} \|\mathbf{y} - \mathbf{A}\mathbf{x}\|_{\mathbf{W}}^2 + R(\mathbf{x}) + \iota_{\Omega}(\mathbf{x}) \right\}$$

using the proposed relaxed linearized AL method, we apply the following substitution:

$$\begin{cases} \mathbf{A} \leftarrow \mathbf{W}^{1/2} \mathbf{A} \\ \mathbf{y} \leftarrow \mathbf{W}^{1/2} \mathbf{y} \end{cases}$$

and set $h \triangleq R + \iota_{\Omega}$.

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The image update now is a **diagonally weighted denoising problem**. We solve it using a **projected gradient descent step** from $\mathbf{x}^{(k)}$. For speed-up, **ordered subsets (OS)** or incremental gradients are used.

Speed-up with decreasing continuation sequence

We also use a **continuation technique** to speed up convergence; that is, we **decrease** ρ gradually with iteration.

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Based on a second-order recursive system analysis, we use

$$\rho_i(\alpha) = \begin{cases} 1, & \text{if } i = 0 \\ \frac{\pi}{\alpha(i+1)} \sqrt{1 - \left(\frac{\pi}{2\alpha(i+1)}\right)^2}, & \text{otherwise.} \end{cases} \quad (2)$$

Therefore, we use a **faster-decreasing** continuation sequence in a **more over-relaxed** linearized AL method.

[HN and Fessler, IEEE Trans. Med. Imag., submitted]

Proposed relaxed linearized algorithm

Algorithm: Relaxed OS-LALM for CT reconstruction.

Input: $K \geq 1$, $M \geq 1$, $0 < \alpha < 2$, and an initial (FBP) image \mathbf{x} .

set $\rho = 1$, $\zeta = \mathbf{g} = M\nabla L_M(\mathbf{x})$, $\mathbf{h} = \mathbf{D}_L\mathbf{x} - \zeta$

for $k = 1, 2, \dots, K$ **do**

for $m = 1, 2, \dots, M$ **do**

$$\mathbf{s} = \rho(\mathbf{D}_L\mathbf{x} - \mathbf{h}) + (1 - \rho)\mathbf{g}$$

$$\mathbf{x}^+ = [\mathbf{x} - (\rho\mathbf{D}_L + \mathbf{D}_R)^{-1}(\mathbf{s} + \nabla R(\mathbf{x}))]_{\Omega}$$

$$\zeta = M\nabla L_m(\mathbf{x}^+)$$

$$\mathbf{g}^+ = \frac{\rho}{\rho+1}(\alpha\zeta + (1 - \alpha)\mathbf{g}) + \frac{1}{\rho+1}\mathbf{g}$$

$$\mathbf{h}^+ = \alpha(\mathbf{D}_L\mathbf{x}^+ - \zeta) + (1 - \alpha)\mathbf{h}$$

 decrease ρ using (2)

end

end

Output: The final image \mathbf{x} .

Chest region helical scan

We reconstruct a $600 \times 600 \times 222$ image from an $888 \times 64 \times 3611$ helical (pitch 1.0) CT scan.

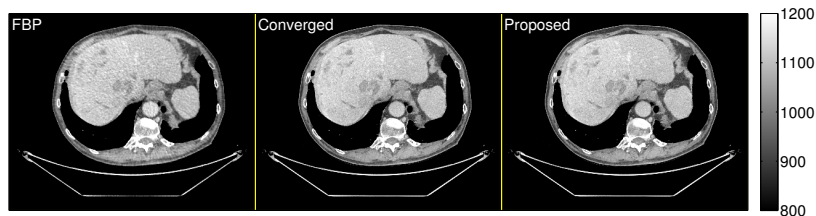


Figure: Chest: Cropped images of the initial FBP image $\mathbf{x}^{(0)}$ (left), the reference reconstruction \mathbf{x}^* (center), and the reconstructed image $\mathbf{x}^{(20)}$ using relaxed OS-LALM with 10 subsets after 20 iterations (right).

Chest region helical scan (cont'd)

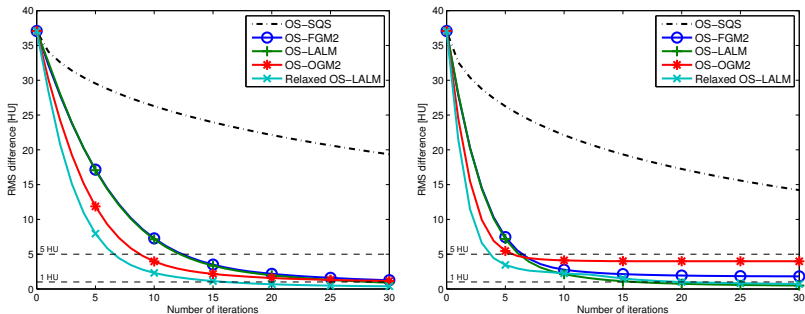


Figure: Chest: Convergence rate curves of different OS algorithms with 10 (left) and 20 (right) subsets.

Chest region helical scan (cont'd)

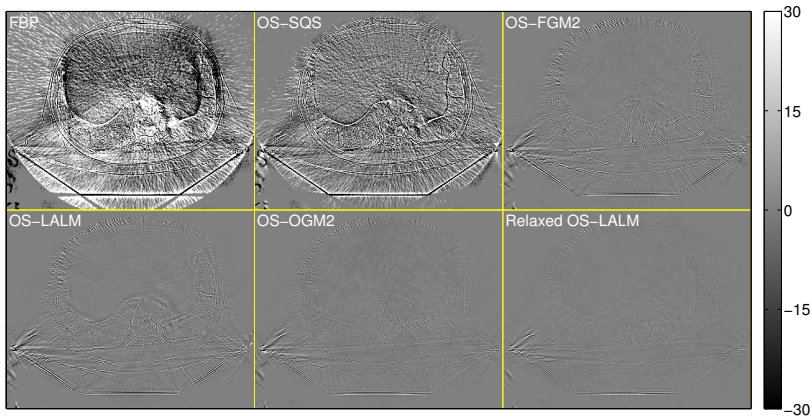


Figure: Chest: Difference images of the initial FBP image $\mathbf{x}^{(0)} - \mathbf{x}^*$ and the reconstructed image $\mathbf{x}^{(10)} - \mathbf{x}^*$ using OS algorithms with 10 subsets after 10 iterations.

Chest region helical scan (cont'd)

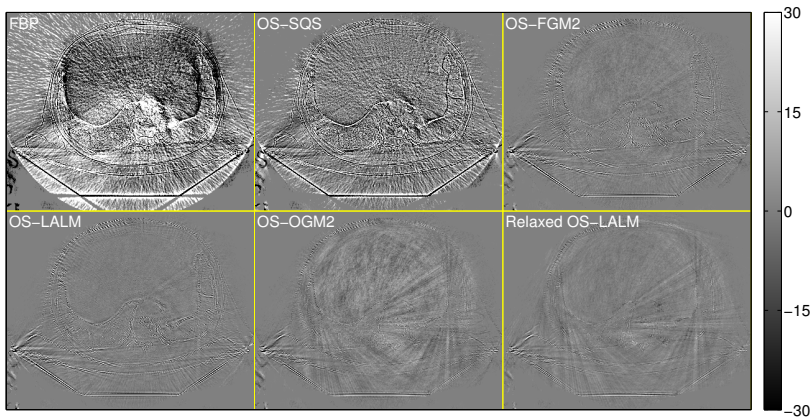


Figure: Chest: Difference images of the initial FBP image $\mathbf{x}^{(0)} - \mathbf{x}^*$ and the reconstructed image $\mathbf{x}^{(5)} - \mathbf{x}^*$ using OS algorithms with 20 subsets after 5 iterations.

Chest region helical scan (cont'd)

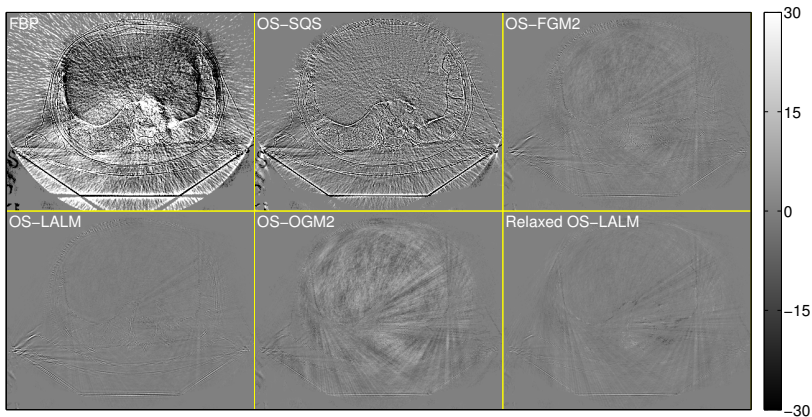


Figure: Chest: Difference images of the initial FBP image $\mathbf{x}^{(0)} - \mathbf{x}^*$ and the reconstructed image $\mathbf{x}^{(10)} - \mathbf{x}^*$ using OS algorithms with 20 subsets after 10 iterations.

Chest region helical scan (cont'd)

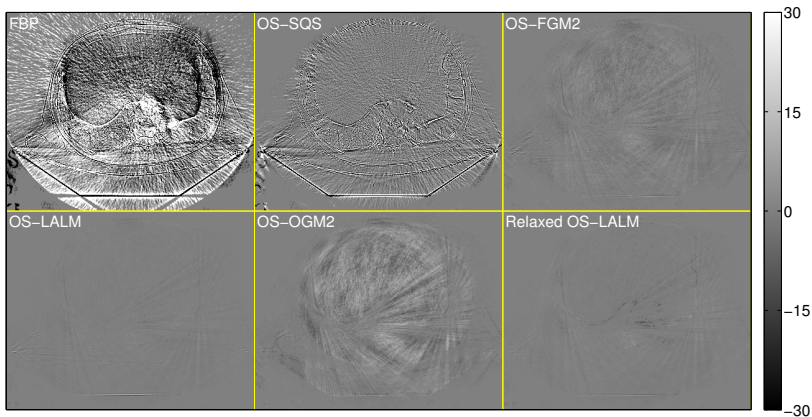


Figure: Chest: Difference images of the initial FBP image $\mathbf{x}^{(0)} - \mathbf{x}^*$ and the reconstructed image $\mathbf{x}^{(20)} - \mathbf{x}^*$ using OS algorithms with 20 subsets after 20 iterations. [More results](#)

Conclusions and future work

In summary,

- ▶ We proposed a **relaxed variant** of linearized AL methods for faster X-ray CT image reconstruction
- ▶ Experimental results showed that the proposed algorithm converges **α -fold faster** than its unrelaxed counterpart
- ▶ The speed-up means that one needs **fewer subsets** to reach an RMS difference criteria in a given number of iterations
- ▶ Empirically, the proposed algorithm is reasonably stable when we use **moderate numbers of subsets**

For future work,

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Acknowledgments

- ▶ Supported in part by NIH grant U01 EB-018753
- ▶ Equipment support from Intel Corporation
- ▶ Research support from GE Healthcare

Shoulder region helical scan

We reconstruct a $512 \times 512 \times 109$ image from an $888 \times 32 \times 7146$ helical (pitch 0.5) CT scan.

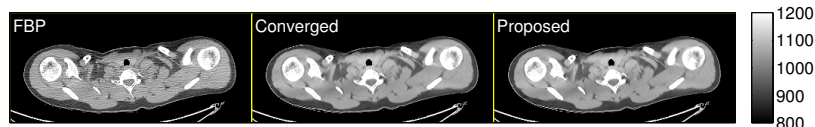


Figure: Shoulder: Cropped images of the initial FBP image $\mathbf{x}^{(0)}$ (left), the reference reconstruction \mathbf{x}^* (center), and the reconstructed image $\mathbf{x}^{(20)}$ using relaxed OS-LALM with 20 subsets after 20 iterations (right).

Shoulder region helical scan (cont'd)

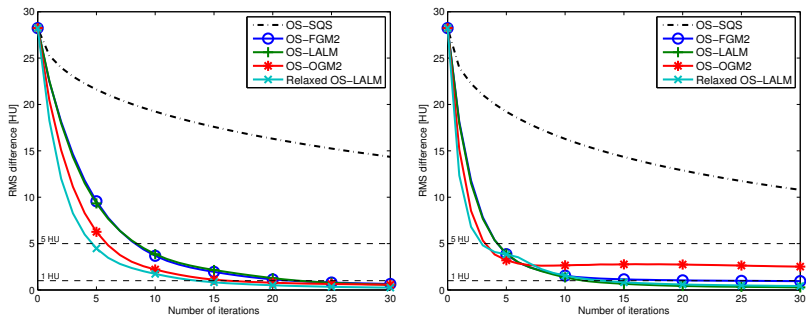


Figure: Shoulder: Convergence rate curves of different OS algorithms with 20 (left) and 40 (right) subsets.

Shoulder region helical scan (cont'd)

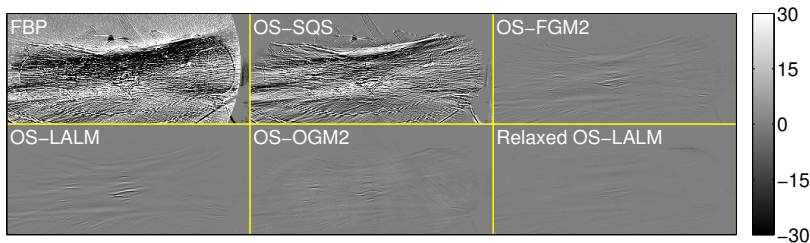


Figure: Shoulder: Difference images of the initial FBP image $\mathbf{x}^{(0)} - \mathbf{x}^*$ and the reconstructed image $\mathbf{x}^{(20)} - \mathbf{x}^*$ using OS algorithms with 20 subsets after 20 iterations.

Abdomen region helical scan

We reconstruct a $600 \times 600 \times 239$ image from an $888 \times 64 \times 3516$ helical (pitch 1.0) CT scan.



Figure: Abdomen: Cropped images of the initial FBP image $\mathbf{x}^{(0)}$ (left), the reference reconstruction \mathbf{x}^* (center), and the reconstructed image $\mathbf{x}^{(20)}$ using relaxed OS-LALM with 10 subsets after 20 iterations (right).

Abdomen region helical scan (cont'd)

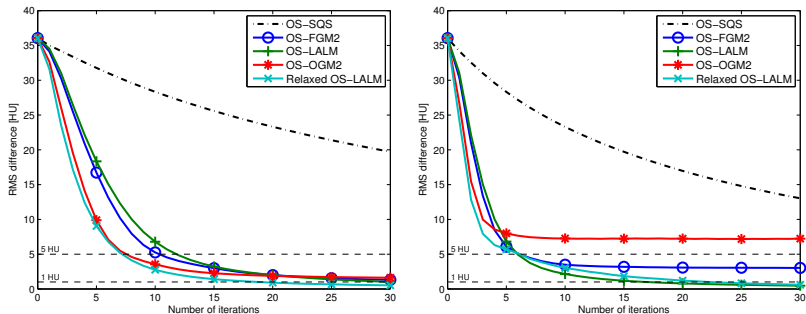


Figure: Abdomen: Convergence rate curves of different OS algorithms with 10 (left) and 20 (right) subsets.

Abdomen region helical scan (cont'd)

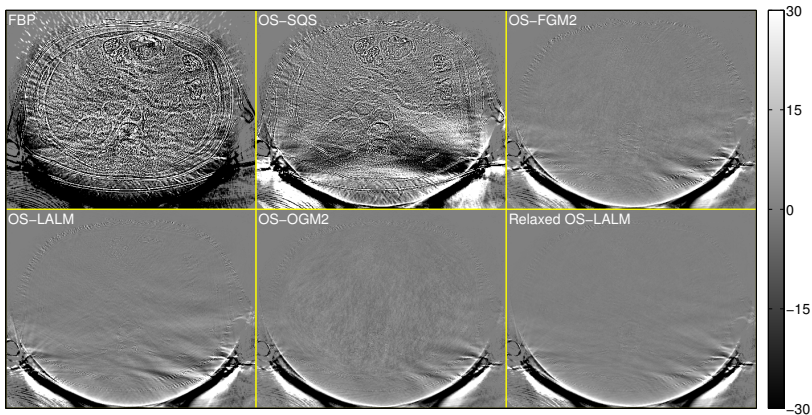


Figure: Abdomen: Difference images of the initial FBP image $\mathbf{x}^{(0)} - \mathbf{x}^*$ and the reconstructed image $\mathbf{x}^{(20)} - \mathbf{x}^*$ using OS algorithms with 10 subsets after 20 iterations. [Conclusions and future work](#)

Simple vs. proposed relaxed OS-LALM

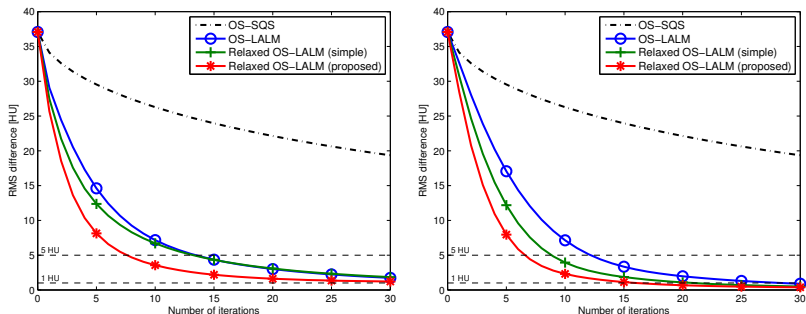


Figure: Chest: Convergence rate curves of different relaxed algorithms with a fixed AL parameter $\rho = 0.05$ (left) and the decreasing ρ (right).

Wide-cone axial scan

We reconstruct a $718 \times 718 \times 440$ image from an $888 \times 256 \times 984$ axial CT scan.

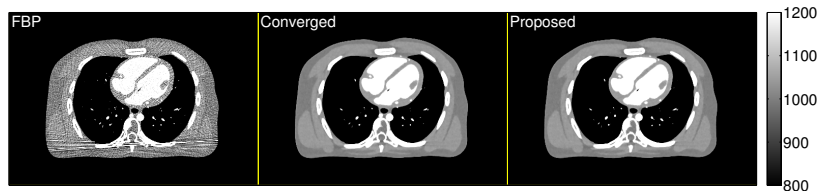


Figure: Wide-cone: Cropped images of the initial FBP image $\mathbf{x}^{(0)}$ (left), the reference reconstruction \mathbf{x}^* (center), and the reconstructed image $\mathbf{x}^{(20)}$ using relaxed OS-LALM with 24 subsets after 20 iterations (right).

Wide-cone axial scan (cont'd)

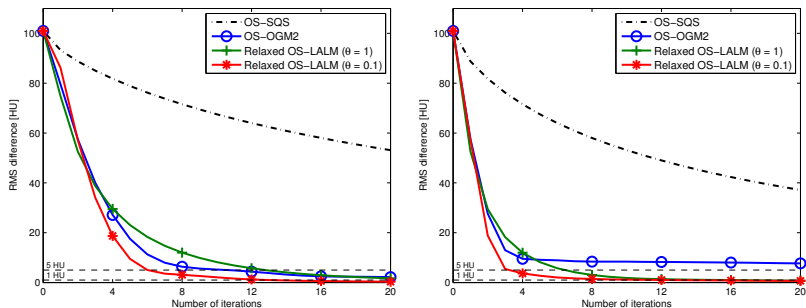


Figure: Wide-cone: Convergence rate curves of different OS algorithms with 12 (left) and 24 (right) subsets.