

# FreeCT\_ICD: Free, Open-Source MBIR Reconstruction Software for Diagnostic CT

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# Outline

1) Background

2) FreeCT\_ICD

- Approach
- Software

3) Results

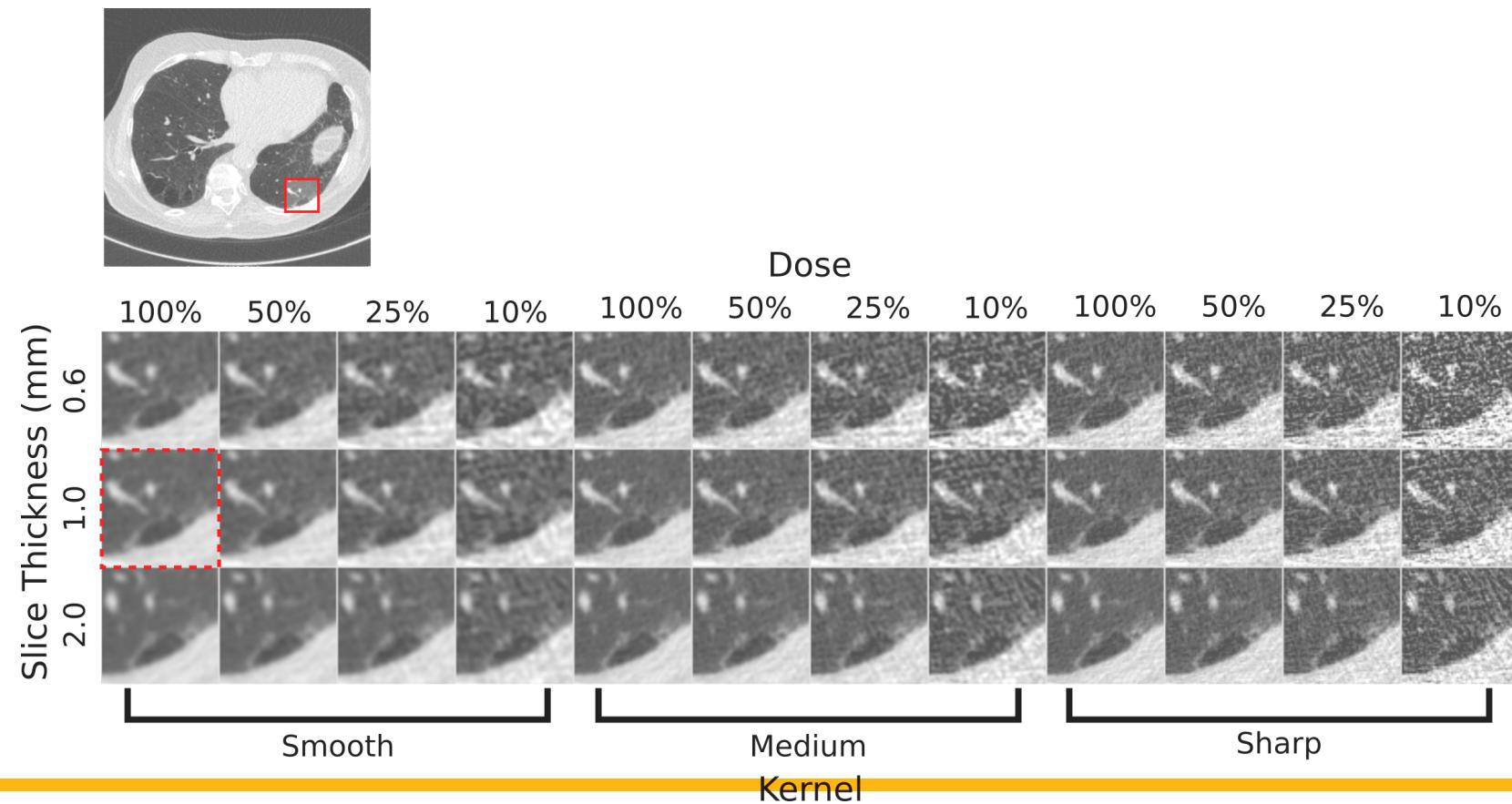
4) Conclusions

# Background

- Needed large-scale imaging datasets for quantitative imaging research

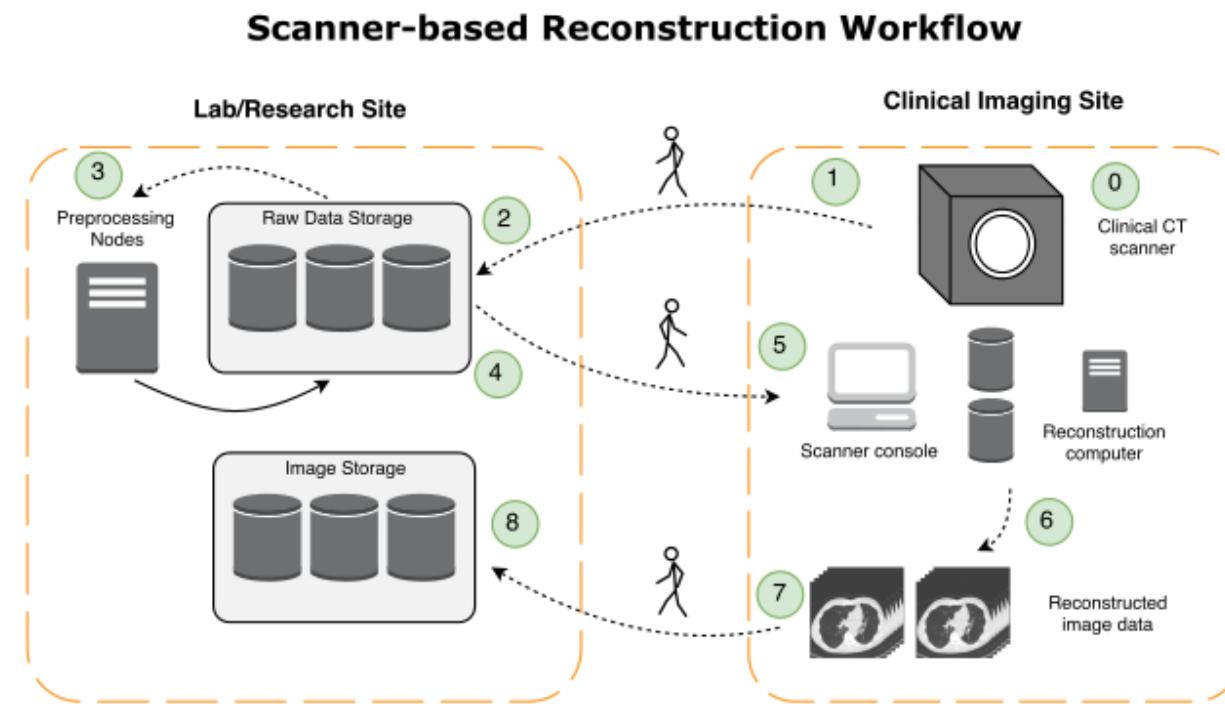
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- Needed large-scale imaging datasets for quantitative imaging research
- Problematic:
  - Not-automatable, time-consuming
  - Easy to make mistakes
  - Not flexible
- Existing “off-line” tools are not geared towards reconstruction of diagnostic data

# Background

- FreeCT\_wFBP (Hoffman et al. 2016)
  - Weighted filtered backprojection (Stierstorfer et al. 2004)
  - Support for reconstructing full diagnostic CT scans without needing access to the scanner
  - **Reduced time for large scale imaging dataset generation from weeks/months to <1 day**



J. Hoffman, S. Young, F. Noo, and M. McNitt-Gray, Technical Note: FreeCT\_wFBP: A robust, efficient, open-source implementation of weighted filtered backprojection for helical, fan-beam CT, Med. Phys. **43**(3), 1411–1420 (2016).

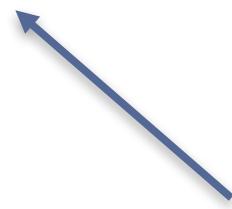
# Background

- While wFBP is an important tool...

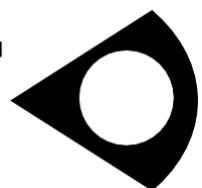


# Background

- While wFBP is an important tool...
- “Where’s iterative?”



(First question from reviewers!)

FreeCT The logo for FreeCT, featuring the word "FreeCT" in a bold, black, sans-serif font followed by a black speaker-like icon consisting of a circle with a triangle inside.

# Today's Goal

- Introduce FreeCT\_ICD software
  - Iterative “companion” to FreeCT\_wFBP
  - Briefly touch on algorithmic/implementation details
  - Software details



# Algorithm Details

- **System Geometry**
  - 3<sup>rd</sup> generation helical CT (most clinical diagnostic scanners)
  - Support for flying focal spots
    - In-plane available, Z & Z+in-plane under development
  - Little things:
    - Quarter detector offset

# Algorithm Details

- Implemented the approach of Thibault et al. 2007

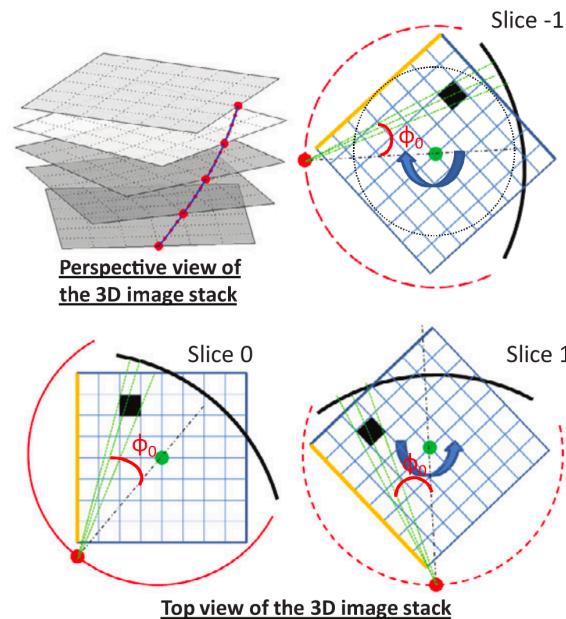
$$\hat{\mathbf{x}} = \arg \min_{\mathbf{x}} \left\{ \frac{1}{2} (\mathbf{y} - \mathbf{Ax})^T \mathbf{D} (\mathbf{y} - \mathbf{Ax}) + U(\mathbf{x}) \right\},$$

- Optimize with iterative coordinate descent (ICD)
  - Good convergence properties (few iterations)

J.-B. Thibault, K.D. Sauer, C. a Bouman, and J. Hsieh, A three-dimensional statistical approach to improved image quality for multislice helical CT., Med. Phys. **34**(2007), 4526–4544 (2007).

# Algorithm Details

- Utilizes a **stored system matrix** approach
  - Exploit symmetry using **rotating slices**



$$\Delta\theta = \frac{2\pi l}{N_{2\pi}}$$

$$d = \frac{lh}{N_{2\pi}}$$

Source: Xu & Tsui 2012

J. Xu and B.M.W. Tsui, Iterative image reconstruction in helical cone-beam x-ray CT using a stored system matrix approach., Phys. Med. Biol. **57**(11), 3477–97 (2012).

# Algorithm Details

- **System Matrix**
  - Joseph's method ... \*
  - (\*) Adapted in our particular case w/ an extra interpolation step that further reduces matrix size
    - No observed loss in image quality
  - Typical matrix sizes are ~10 GB for a Siemens Definition AS 64

# Algorithm Details

- **Penalties**
  - 2D support at present (w/ plans for 3D)
  - Quadratic penalty

$$\psi(t) = \frac{1}{2}t^2$$

- Edge-preserving (Fair potential)

$$\psi(t; \delta) = \delta^2 \left( \left| \frac{t}{\delta} \right| - \log \left( 1 + \left| \frac{t}{\delta} \right| \right) \right)$$

# Software Details

- Implemented in C++
- Minimal external dependencies (Easy to compile)
  - Boost (<https://www.boost.org/>)
  - Yaml-cpp (<https://github.com/jbeder/yaml-cpp>)
  - OpenMP (<https://www.openmp.org/>)
- Platform independent
  - Tested on Linux and MacOS
  - Windows should be supported, not tested
- GNU GPL v2.0
  - “Free to do whatever you want with the code as long as you share your changes”



Windows 10

# Software Details

- Key features
  - Modularity
    - Easily-modified penalty functions (code structure)
    - “Swappable” system matrices
  - Supports scans from clinical scanners
    - Also does simulated data
  - Ability to initialize with FreeCT\_wFBP
    - Non-trivial when using clinical data and rotating slice geometry
    - Automatically configured based on inputs to FreeCT\_ICD

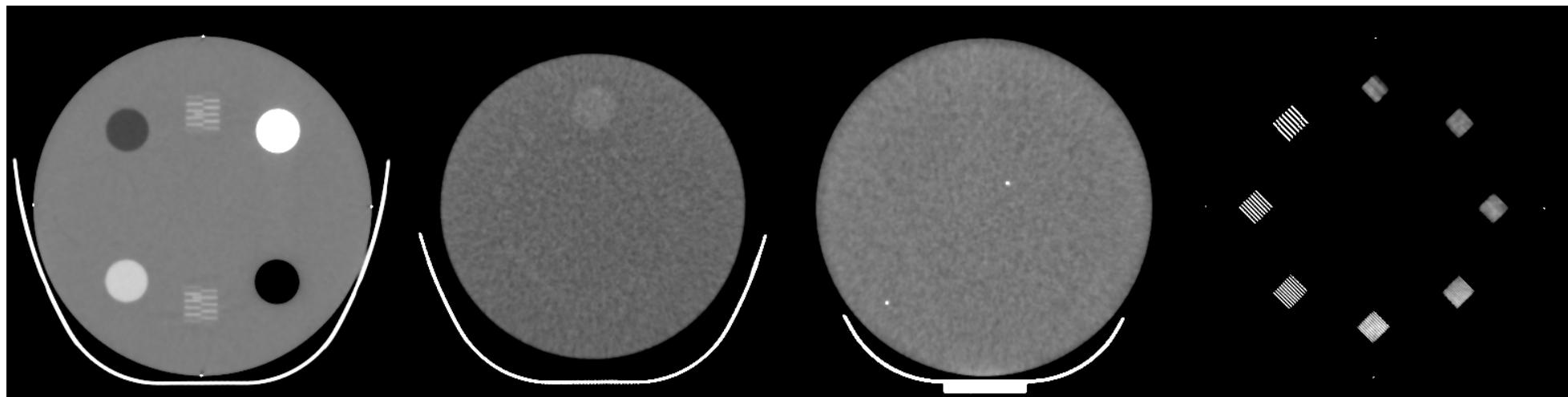
# Some images... finally!

# Results

- ACR Phantom Reconstructions

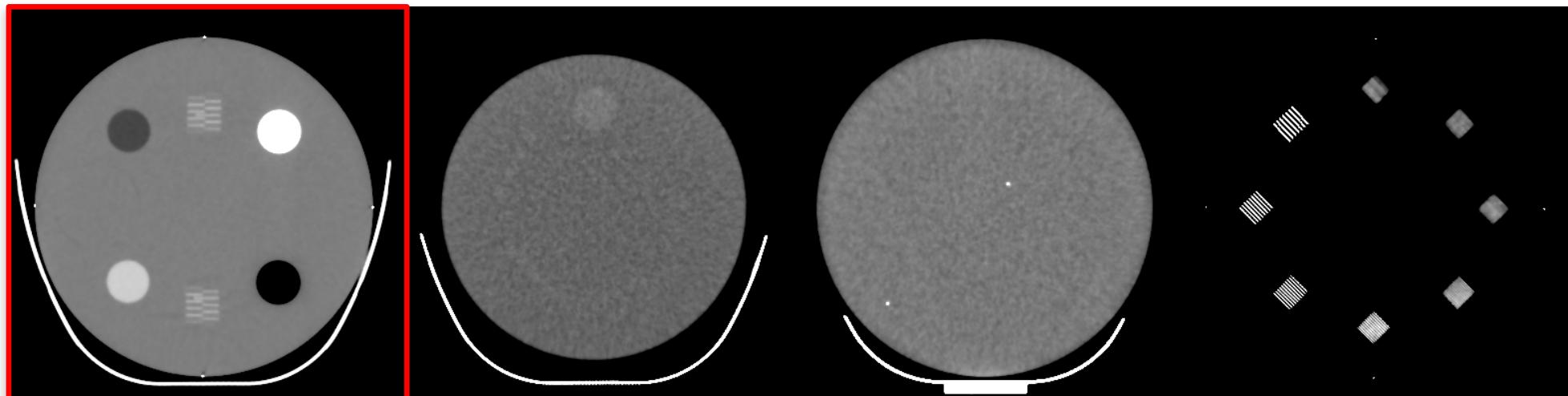
Scan	ACR Phantom
<i>Acquisition Parameters</i>	
Collimation	16 x 1.2mm
Pitch	1.0
Flying focal spot	Off
Rotation time [s]	0.33
<i>Reconstruction Parameters</i>	
wFBP initialization	yes
Voxel grid Dimensions	512 x 512 x 132
Voxel size [mm]	0.58 x 0.58 x 1.5
FOV radius [mm]	300
Edge-preserving parameter	0.005
Penalty term parameter	0.1
Iterations	50
<b>Matrix size [GB]</b>	<b>8.5</b>

# Results



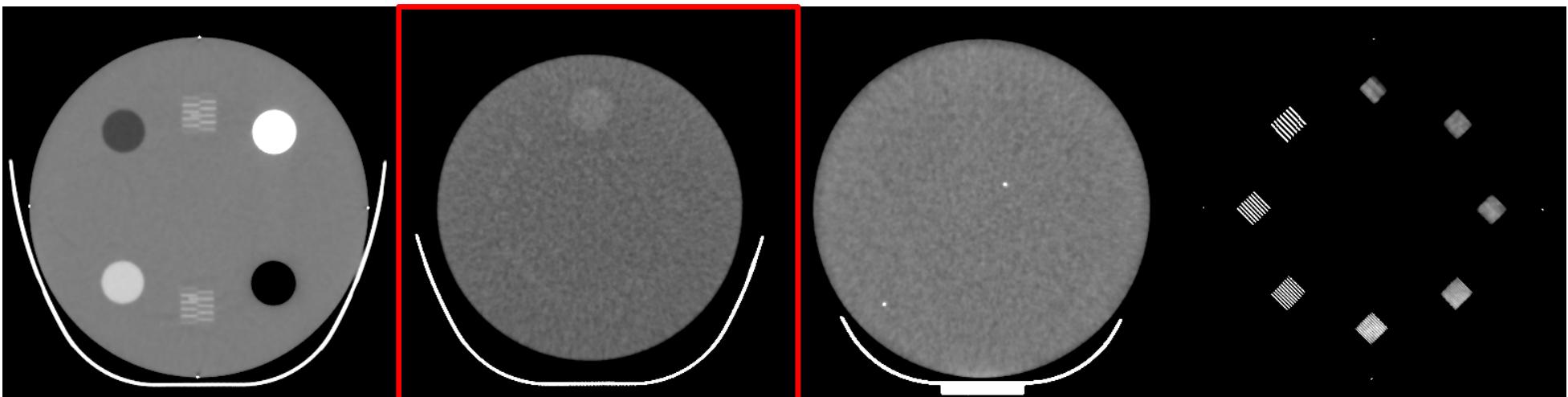
ACR CT Accreditation Phantom/Protocols

# Results



Material	Acceptable range [HU]	FreeCT_ICD [HU]
Polyethylene	-107 to -84	-89
Bone	850 to 970	864
Water	-7 to 7	-2
Acrylic	110 to 135	123
Air	-1005 to -970	-988

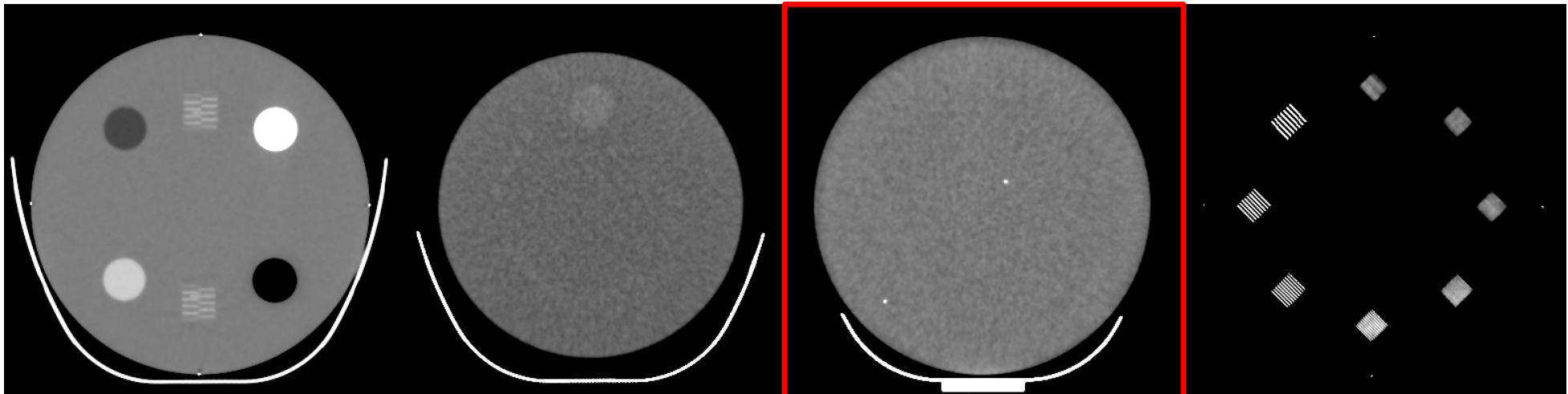
# Results



CNR: 3.83 ( > 1.0 req.)

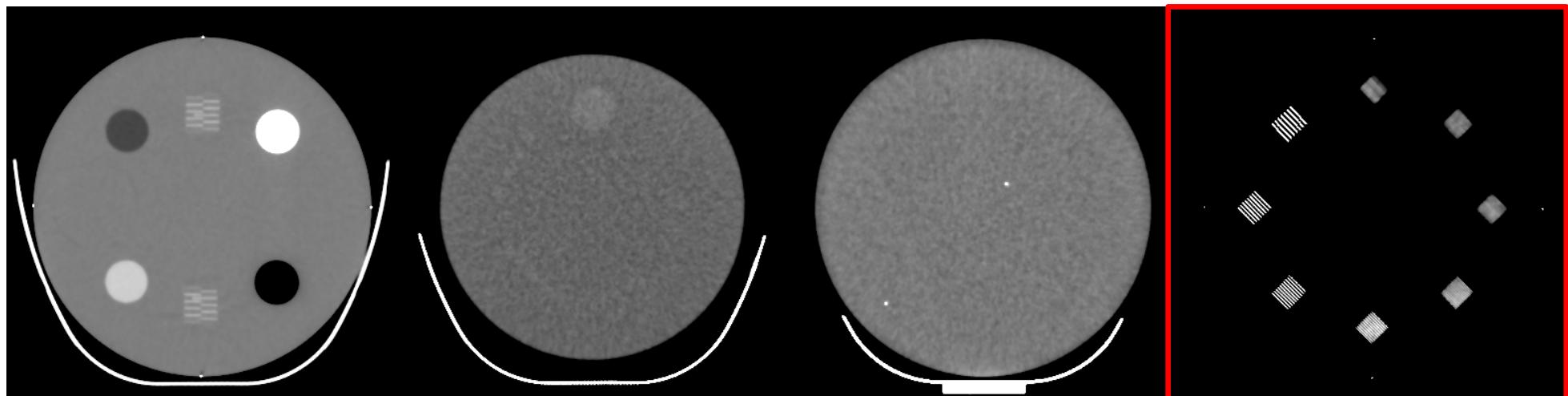
wFBP: 2.21

# Results

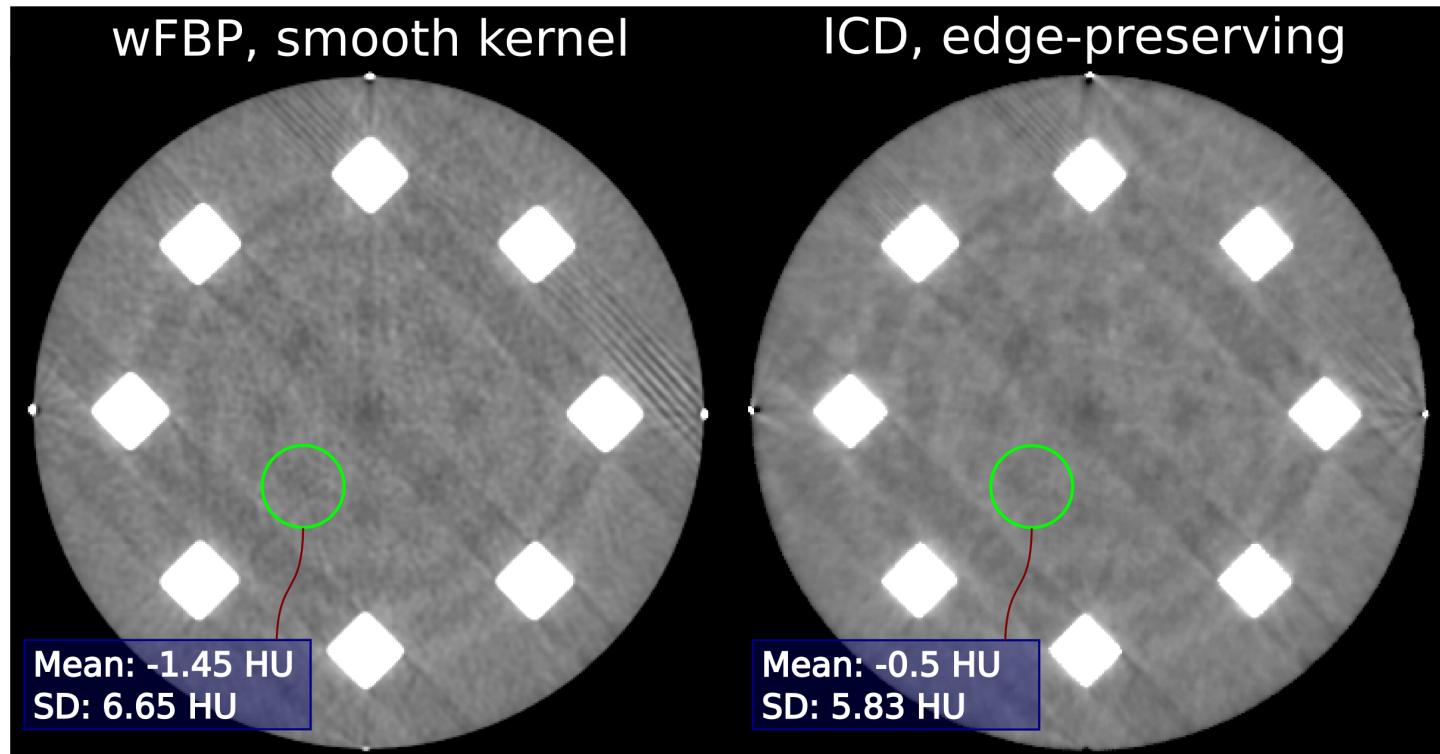


Uniformity: 0.9HU (< 5.0HU req.)

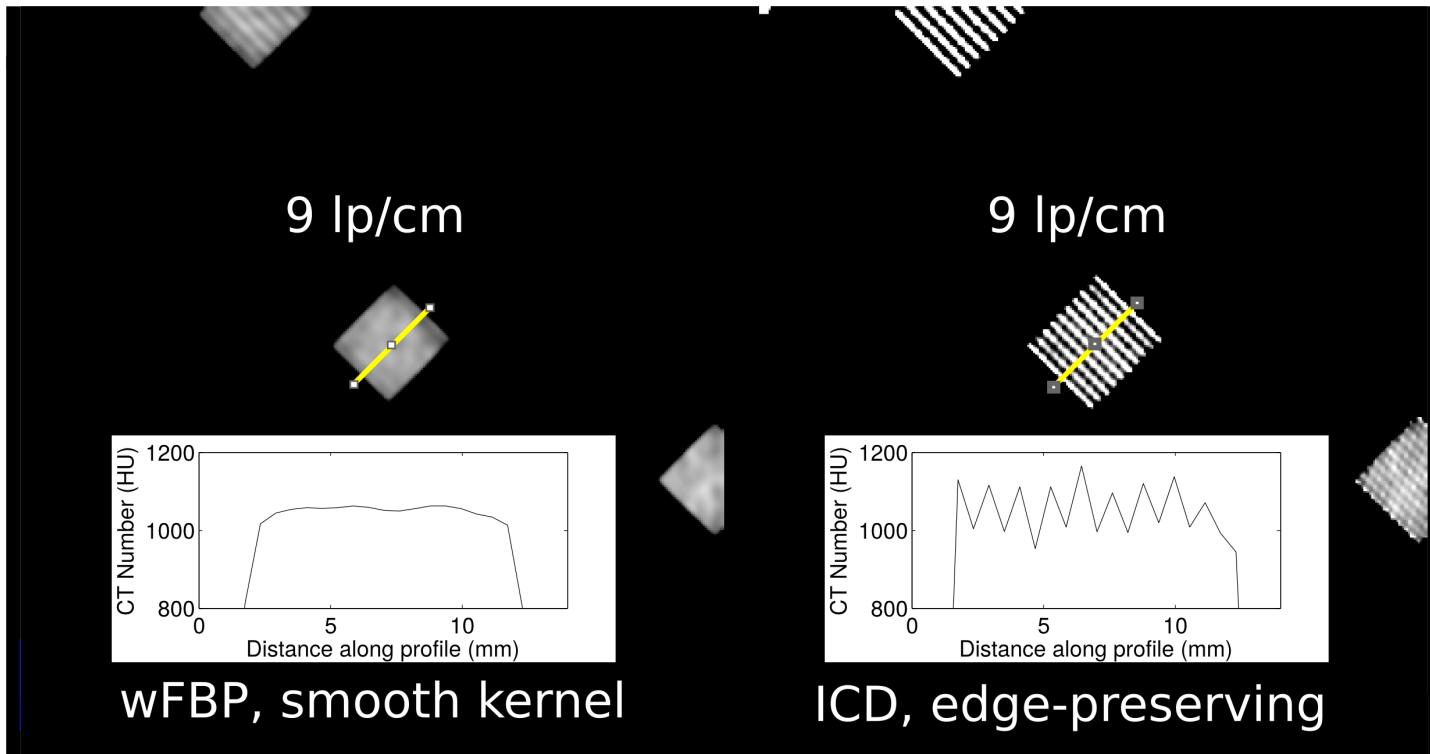
# Results



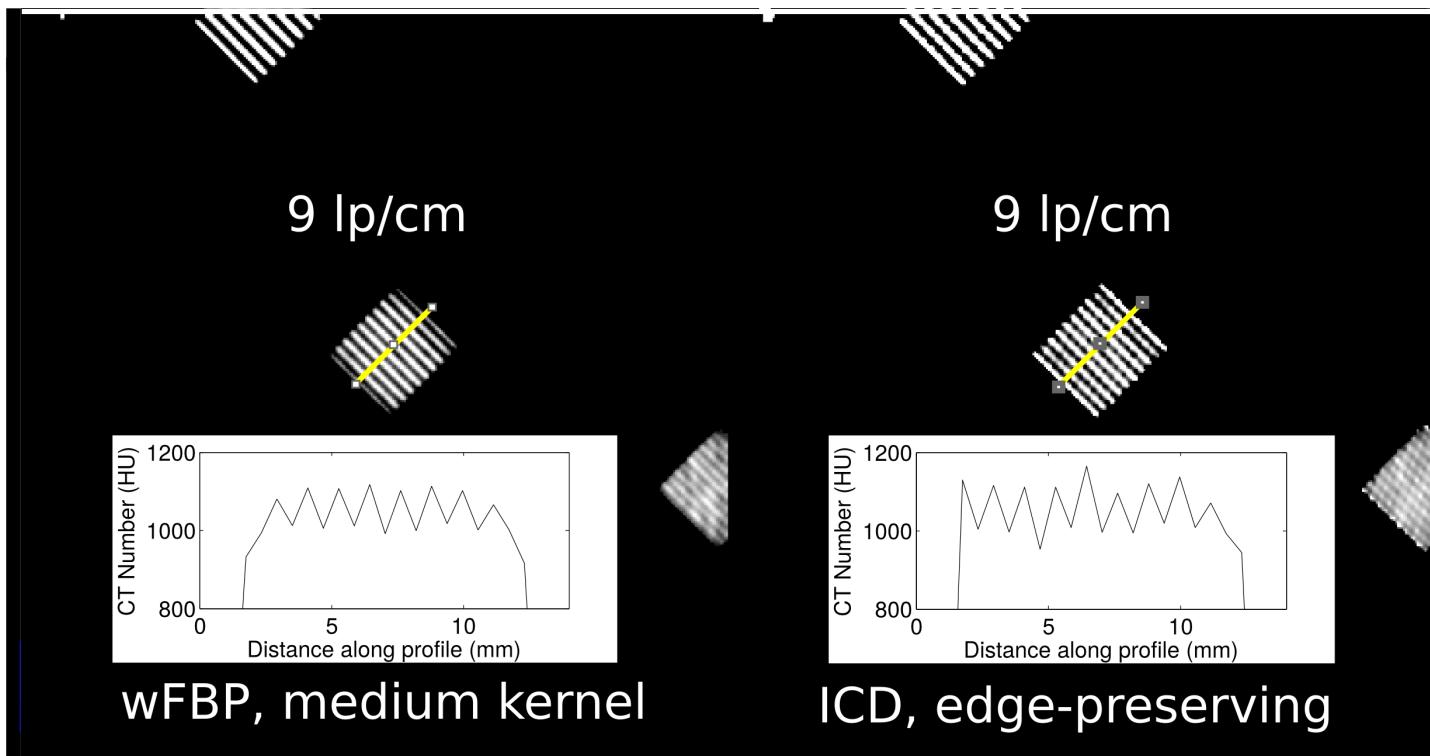
# Results



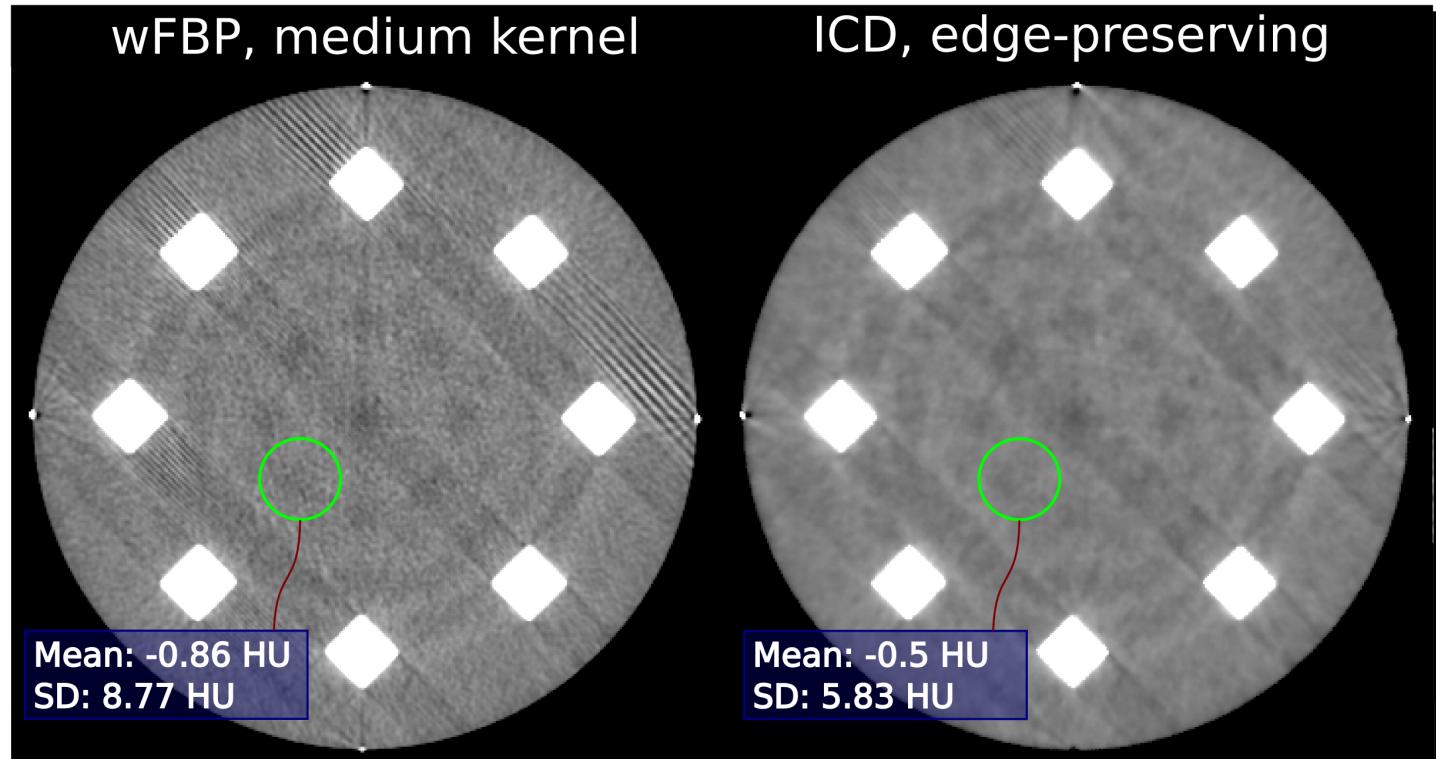
# Results



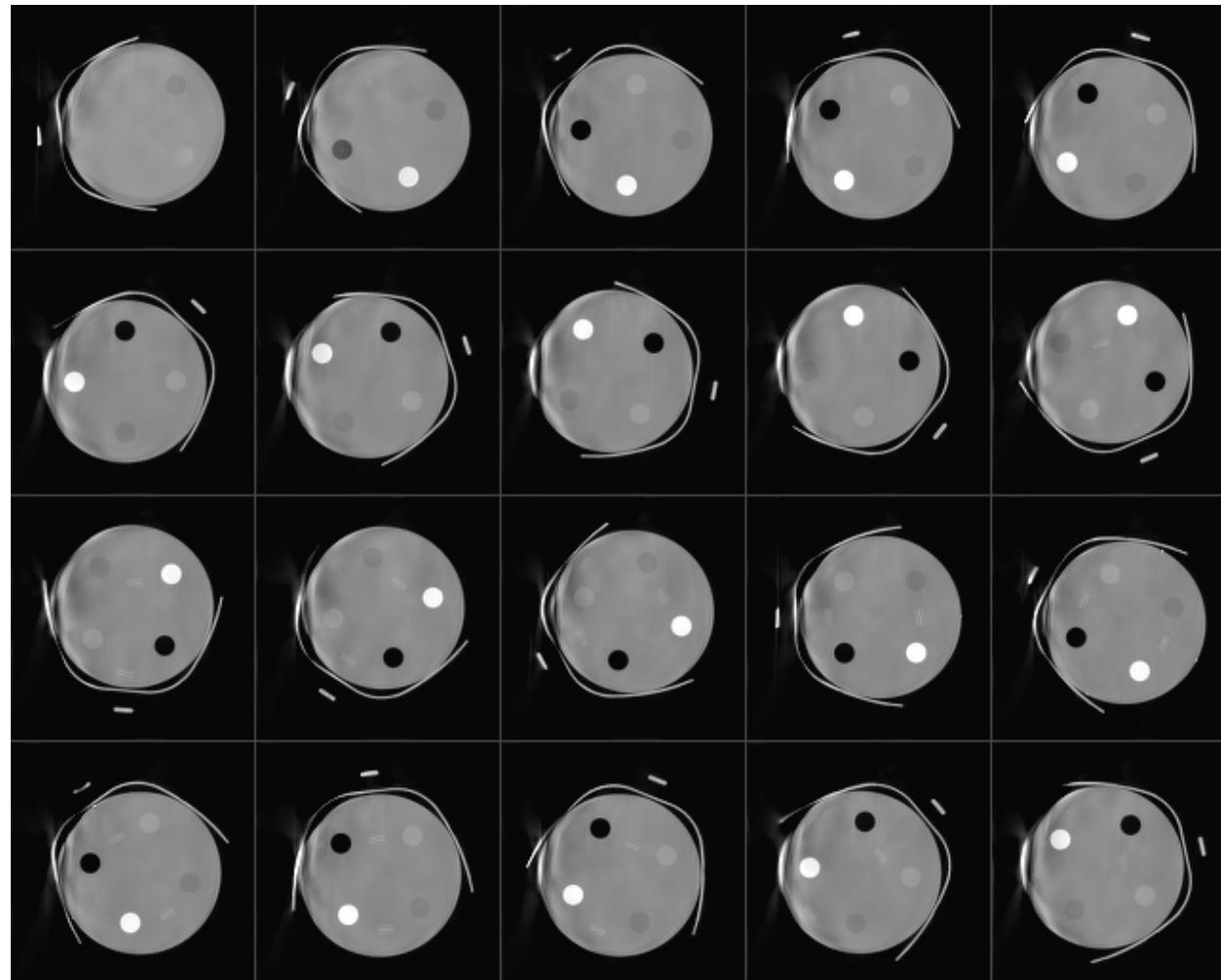
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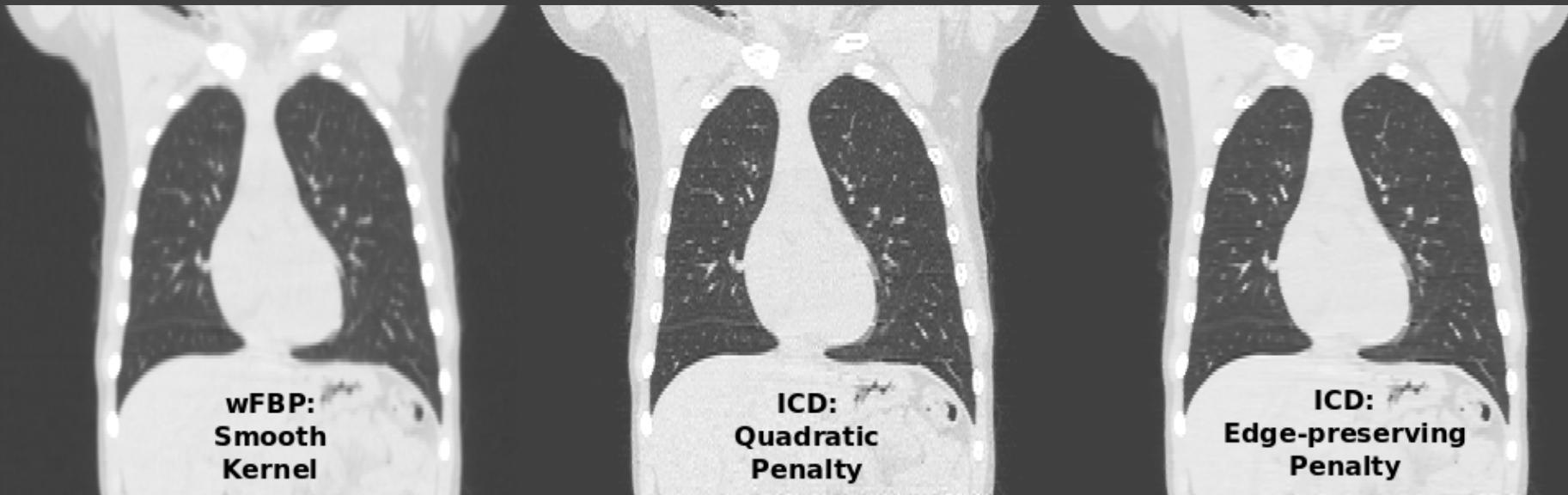


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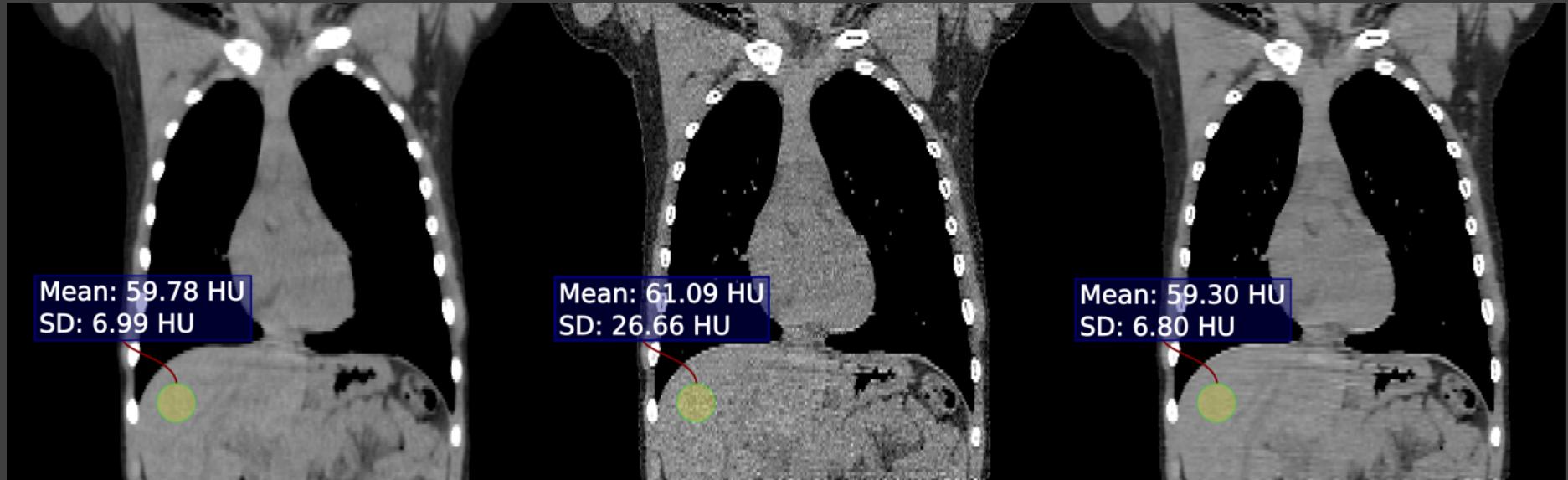
- Pediatric chest

Scan	Pediatric Chest
<i>Acquisition Parameters</i>	
Collimation	16 x 1.2mm
Pitch	1.0
Flying focal spot	Off
Rotation time [s]	0.33
<i>Reconstruction Parameters</i>	
wFBP initialization	yes
Voxel grid Dimensions	512 x 512 x 163
Voxel size [mm]	0.98 x 0.98 x1.5
FOV radius [mm]	500
Edge-preserving parameter	0.005
Penalty term parameter	0.1
Iterations	50
<b>Matrix size [GB]</b>	<b>14.6</b>

# Results



# Results

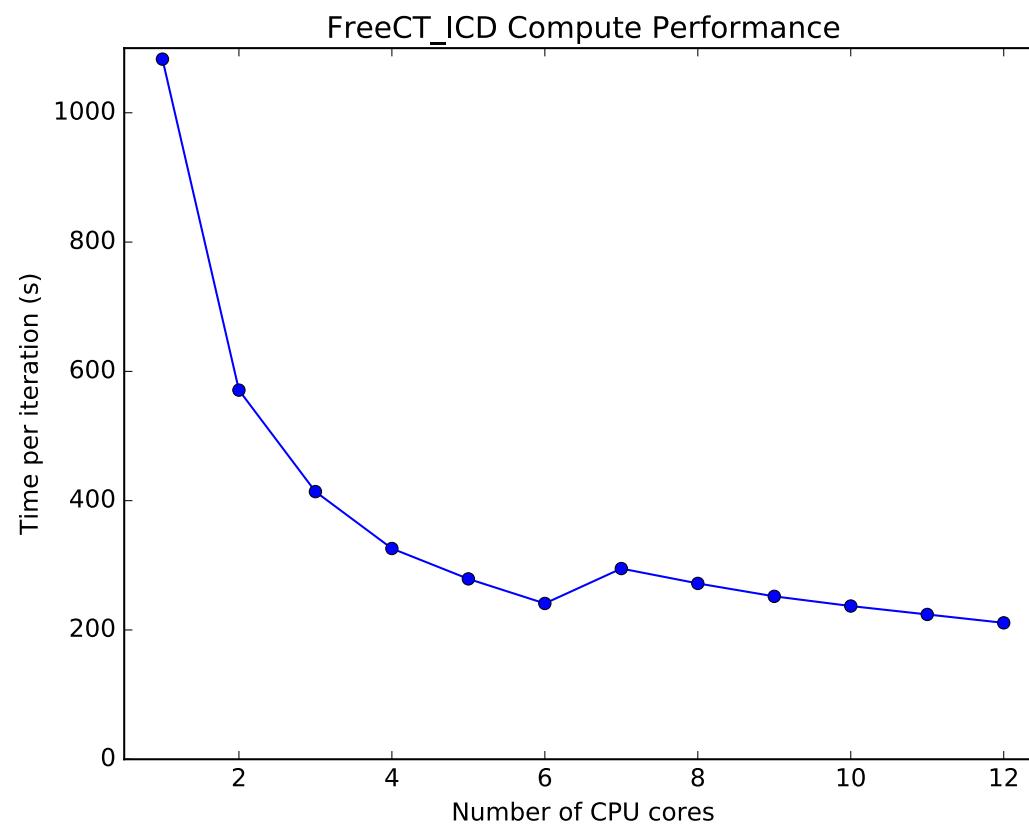


# Results

- Execution time/speed
  - ACR Dataset (512x512x120, 16x1.2mm, 11 cores)
    - ~20 minutes per iteration
    - With wFBP initializer, ~2-3 hours per reconstruction

# Results

- Execution time/speed



# Limitations and Future Work

- 2D Regularizer/Penalty
  - Working on an extension to 3D, math for rotation between slices
- A little slow
  - Working on a GPU version, theoretically ~1 min/iteration
- Raw data availability
- Documentation (incoming)

# Conclusions

- Free, open-source MBIR reconstruction software
  - GNU GPL v2.0
- Focus on modularity, low-dependency design
- Reconstructs clinical data

# Final Thoughts

- Available today!
  - [https://github.com/FreeCT/FreeCT\\_ICD](https://github.com/FreeCT/FreeCT_ICD)
- Feedback helps make it better, let us know if you're using it or have any thoughts, contributions
  - jmhoffman@mednet.ucla.edu
  - freect.project@gmail.com

# Acknowledgements

- Stefano Young
- Frédéric Noo
- Scott Hsieh

# Thank you! Questions?

