

FreeCT_ICD: Free, Open-Source MBIR Reconstruction Software for Diagnostic CT

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Outline

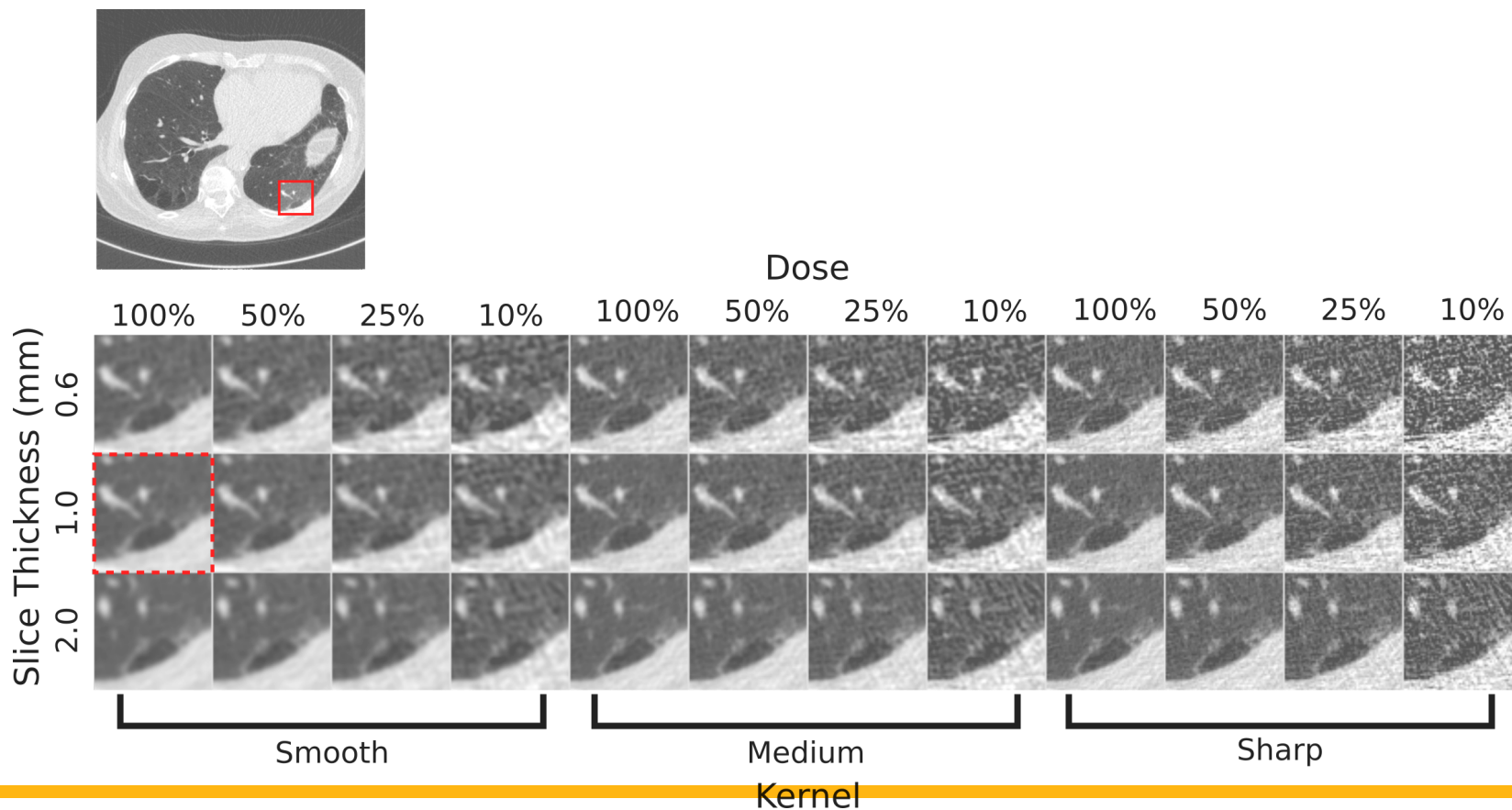
- 1) Background
- 2) FreeCT_ICD
 - Approach
 - Software
- 3) Results
- 4) Conclusions

Background

- Needed large-scale imaging datasets for quantitative imaging research

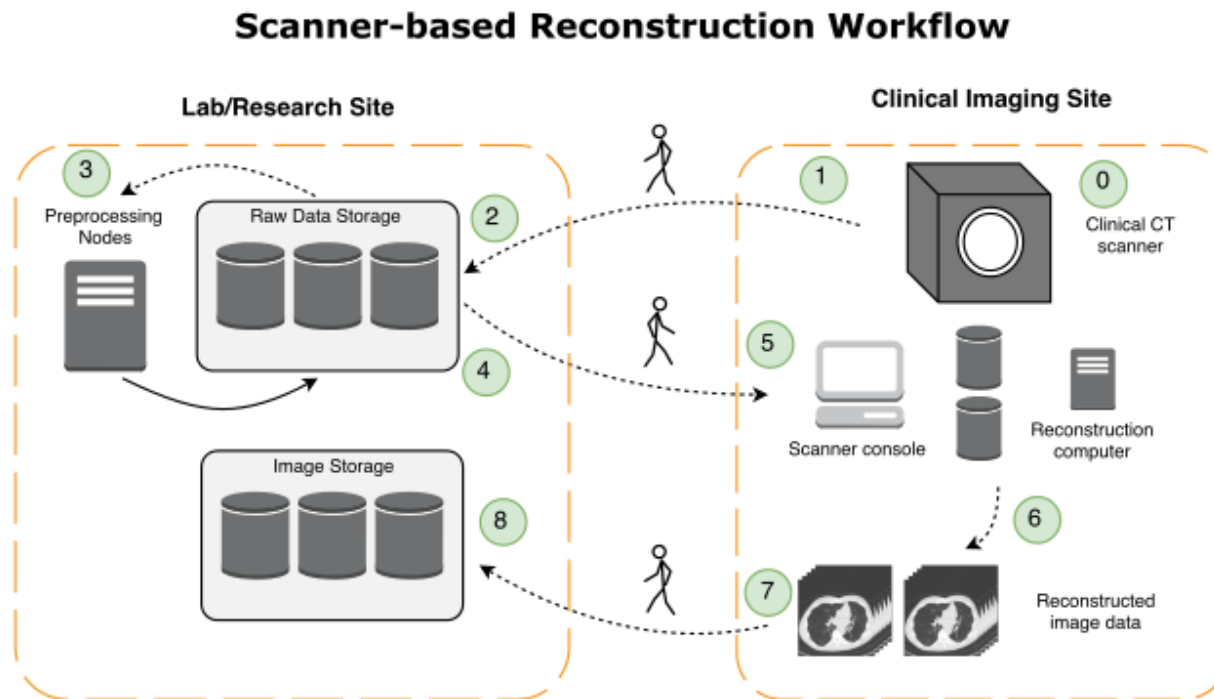
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- Problematic:
 - Not-automatable, time-consuming
 - Easy to make mistakes
 - Not flexible

Background

- 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).

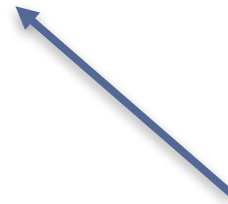
Background

- While wFBP is an important tool...



Background

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



(First question from reviewers!)

FreeCT

Today's Goal

- Introduce FreeCT_ICD software
 - Iterative “companion” to FreeCT_wFBP
 - Briefly touch on algorithmic/implementation details
 - Software details



Algorithm Details

- **System Geometry**
 - 3rd 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

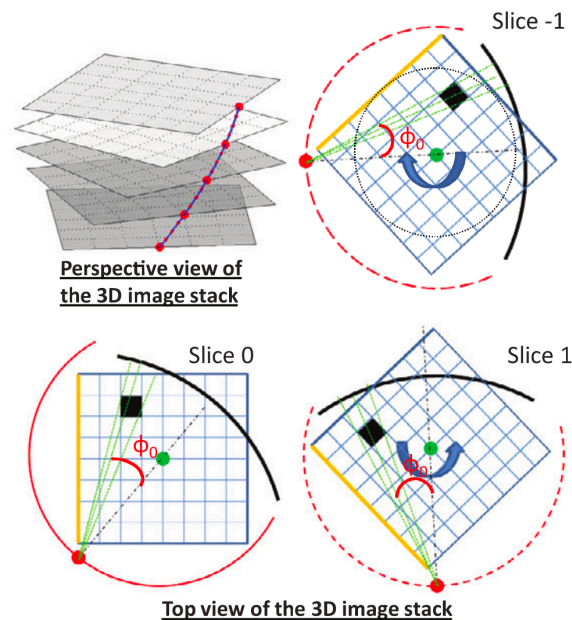
$$\hat{\mathbf{x}} = \arg \min_{\mathbf{x}} \left\{ \frac{1}{2} (\mathbf{y} - \mathbf{Ax})^T \mathbf{D} (\mathbf{y} - \mathbf{Ax}) + U(\mathbf{x}) \right\},$$

- 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**



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

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

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

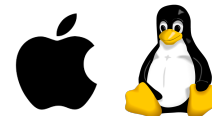
$$\psi(t) = \frac{1}{2}t^2$$

- 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”



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

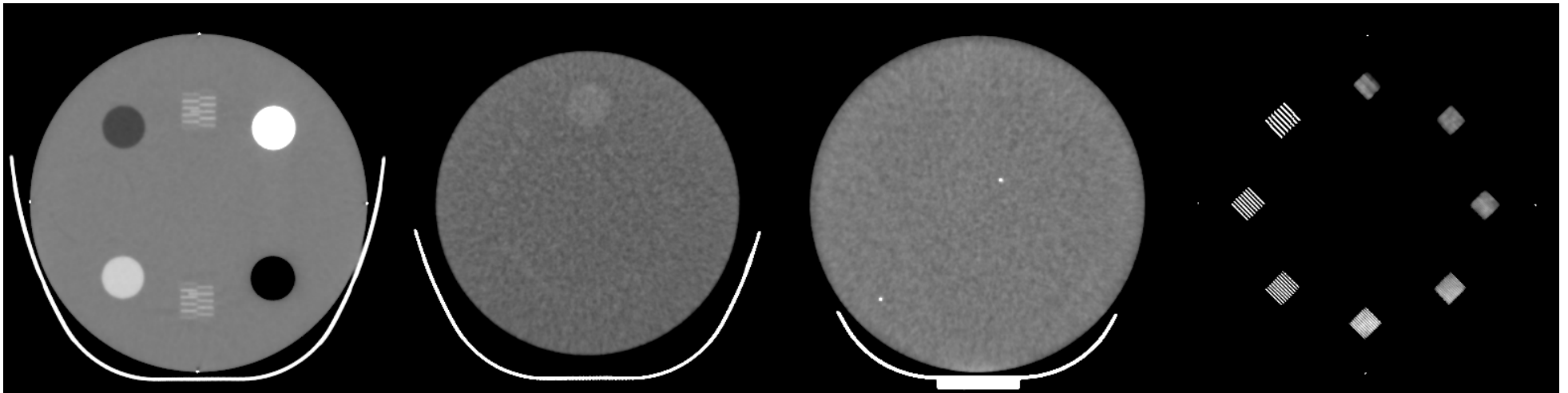
Some images... finally!

Results

- ACR Phantom Reconstructions

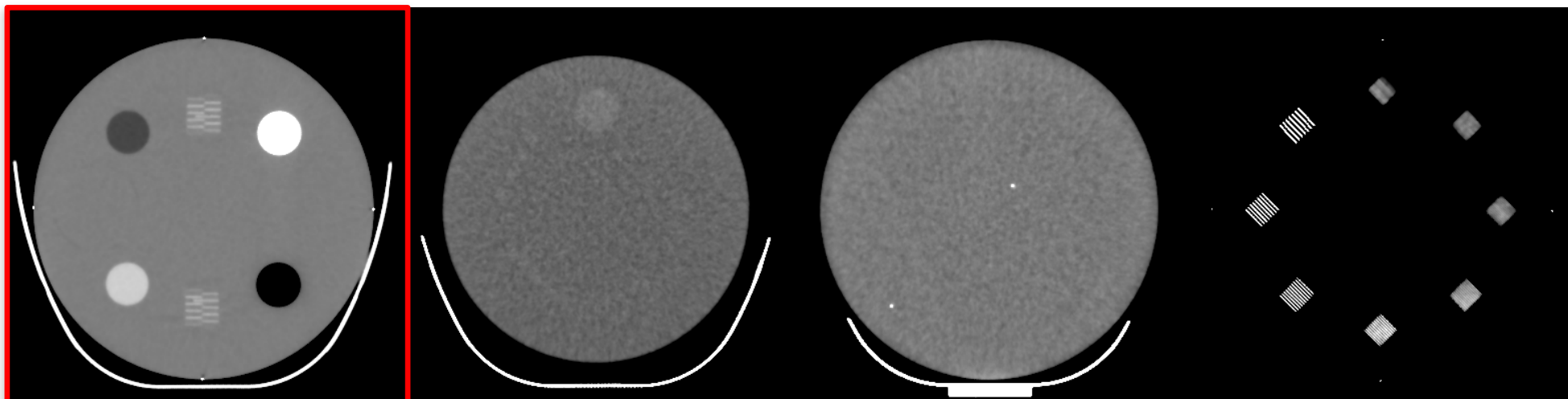
| Scan | ACR Phantom |
|----------------------------------|-------------------|
| <i>Acquisition Parameters</i> | |
| Collimation | 16 x 1.2mm |
| Pitch | 1.0 |
| Flying focal spot | Off |
| Rotation time [s] | 0.33 |
| <i>Reconstruction Parameters</i> | |
| wFBP initialization | yes |
| Voxel grid Dimensions | 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 |
| Matrix size [GB] | 8.5 |

Results



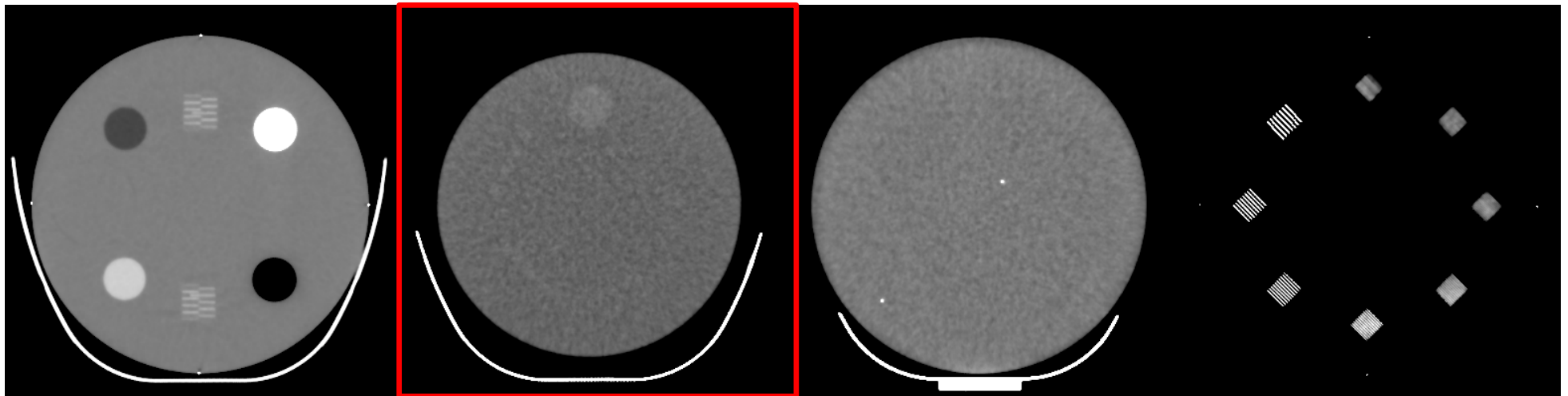
ACR CT Accreditation Phantom/Protocols

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 |

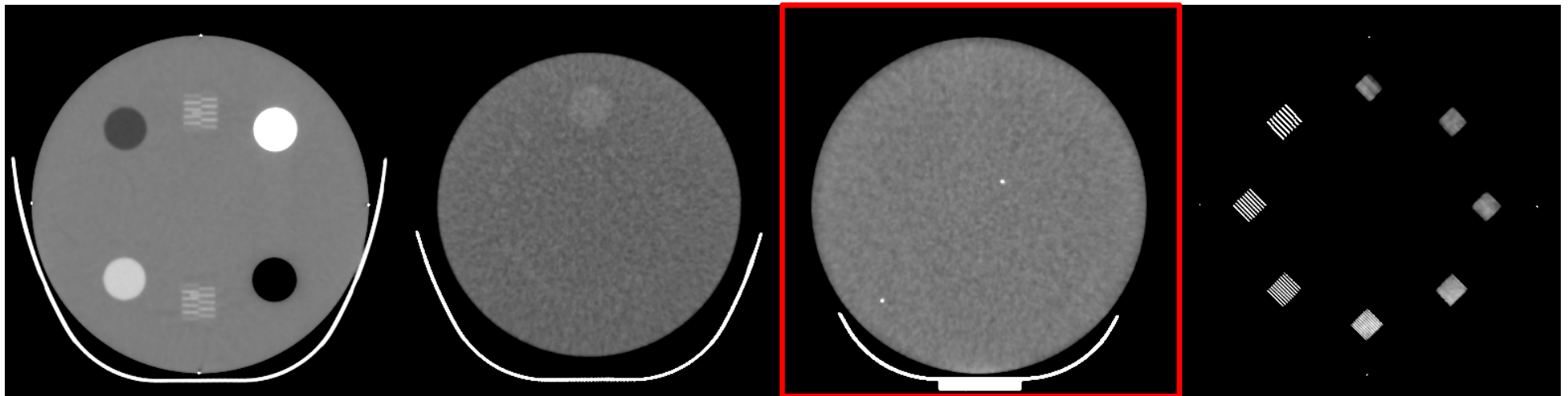
Results



CNR: 3.83 (> 1.0 req.)

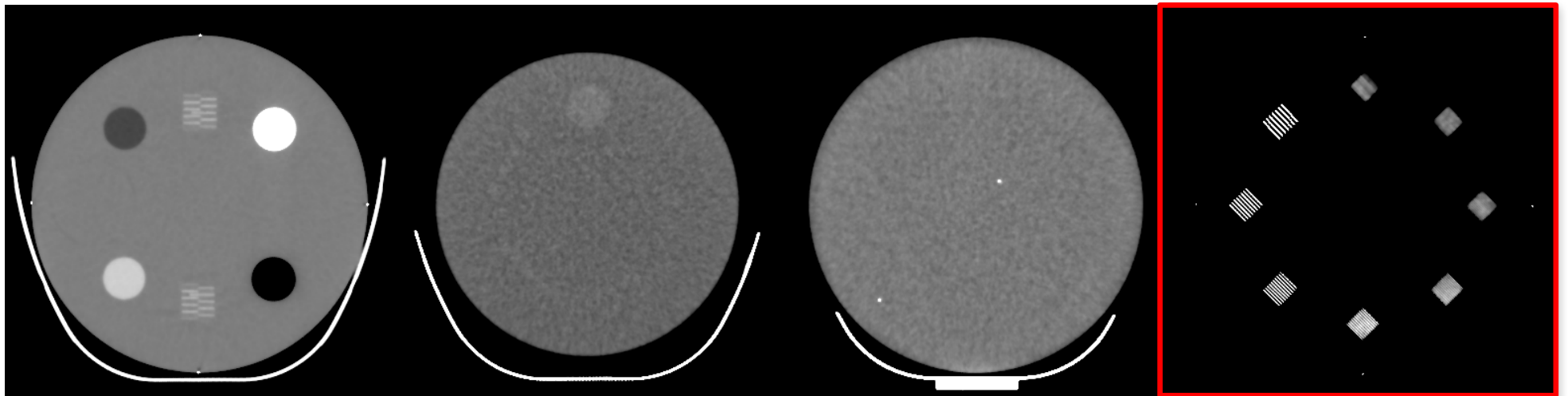
wFBP: 2.21

Results

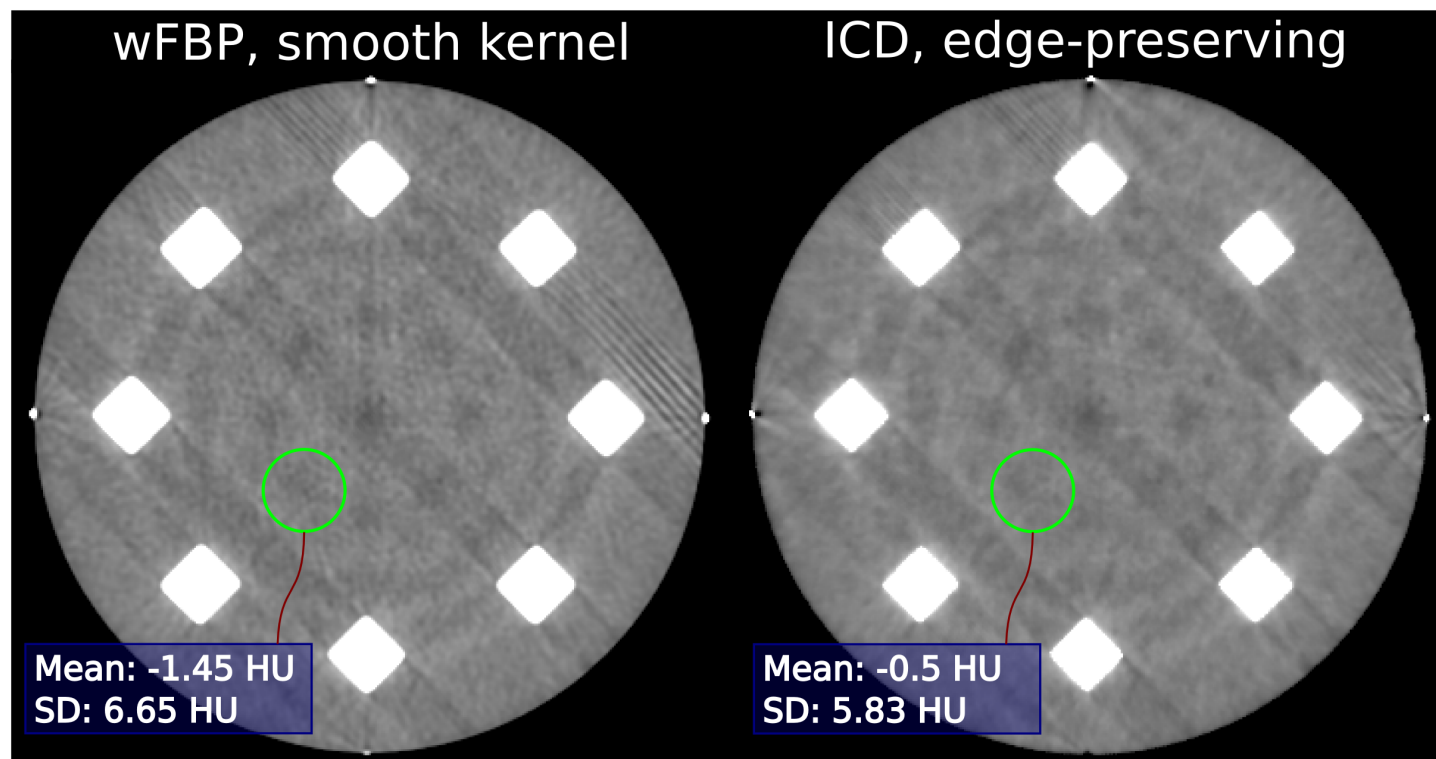


Uniformity: 0.9HU (< 5.0HU req.)

Results

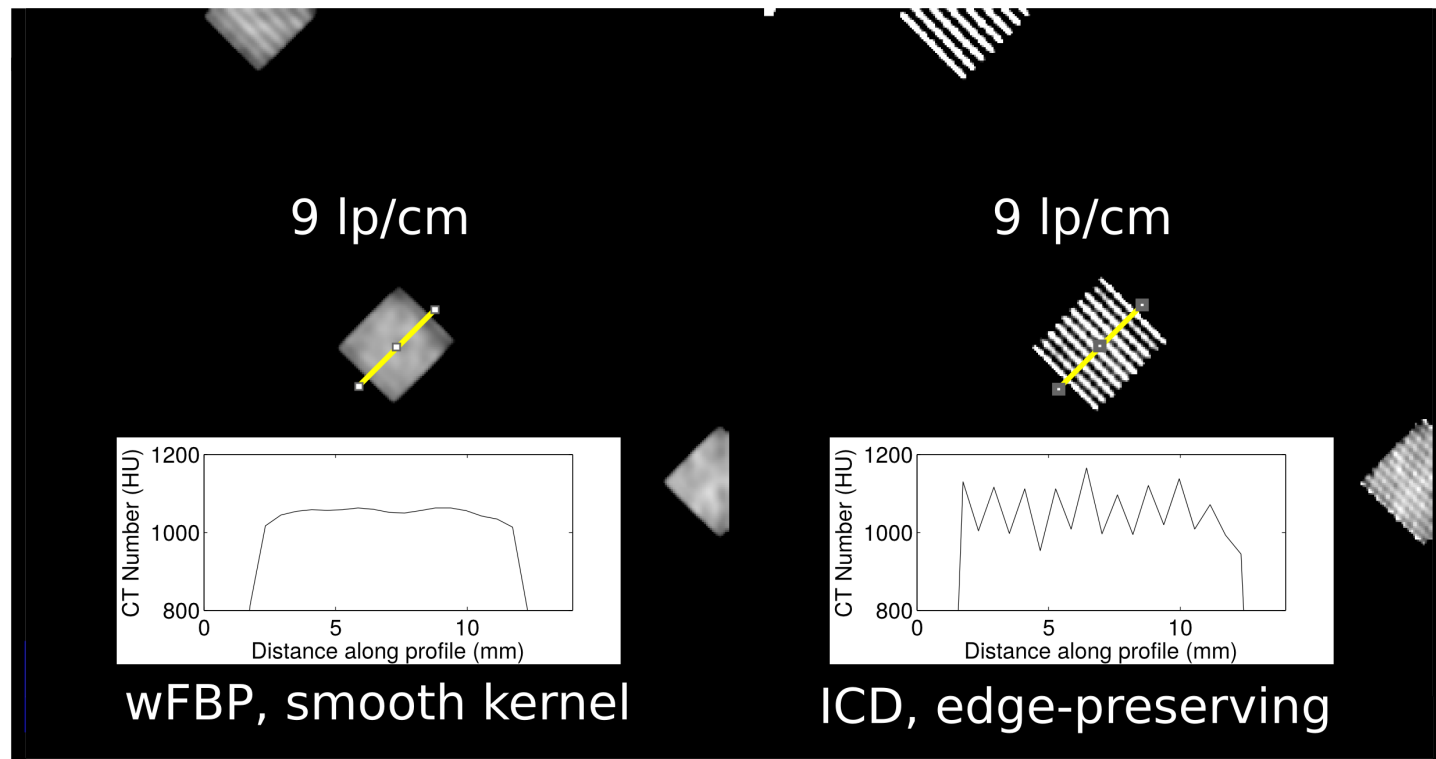


Results



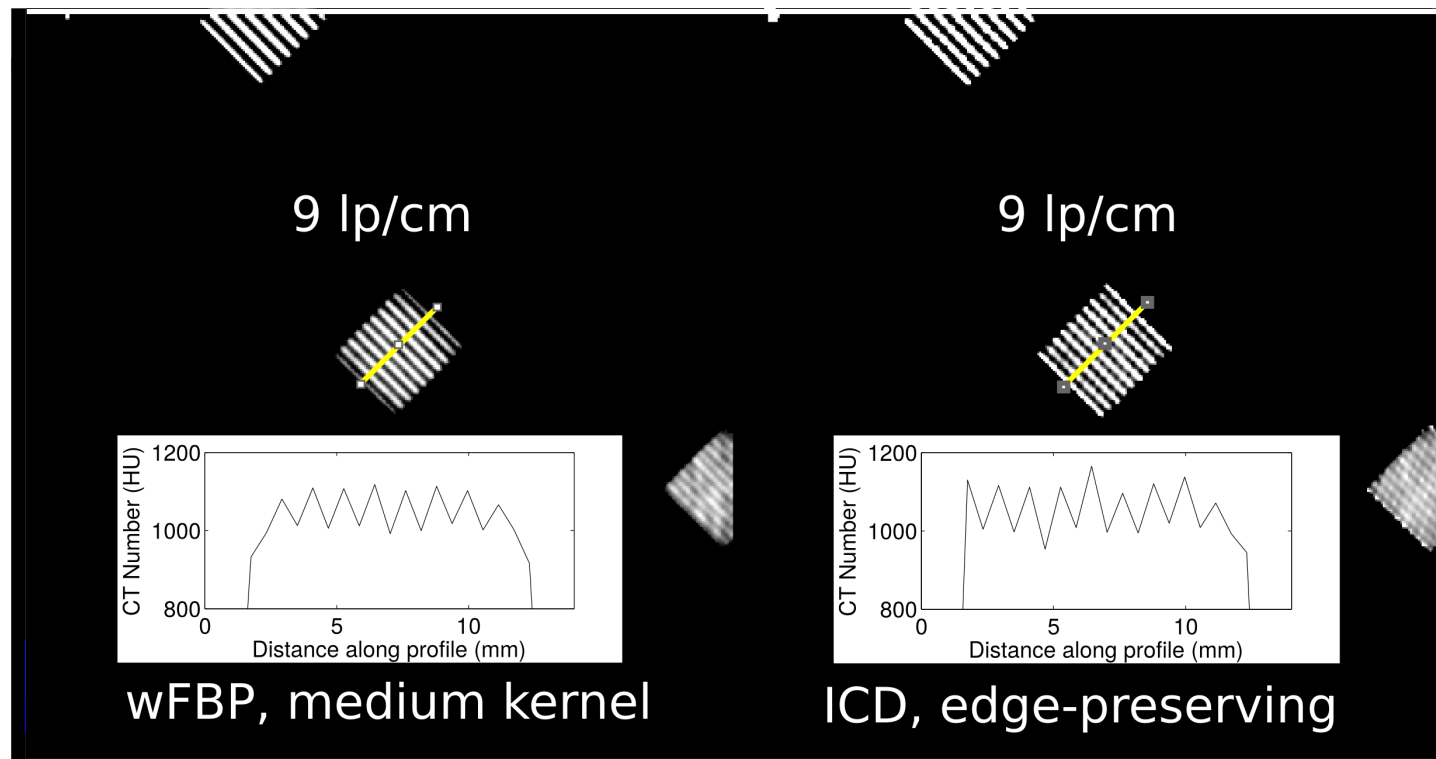
W/L: 200/0 HU

Results



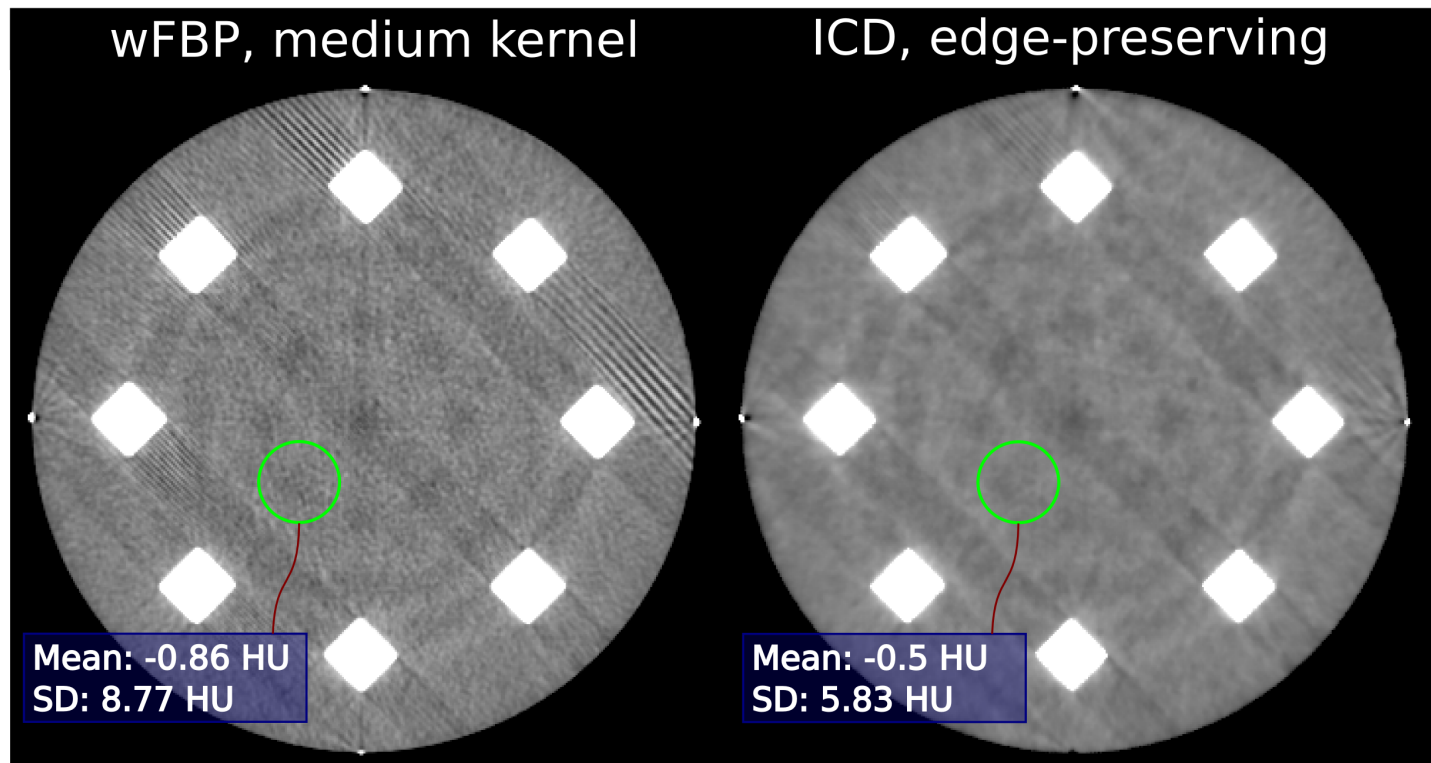
W/L: 100/1050 HU

Results



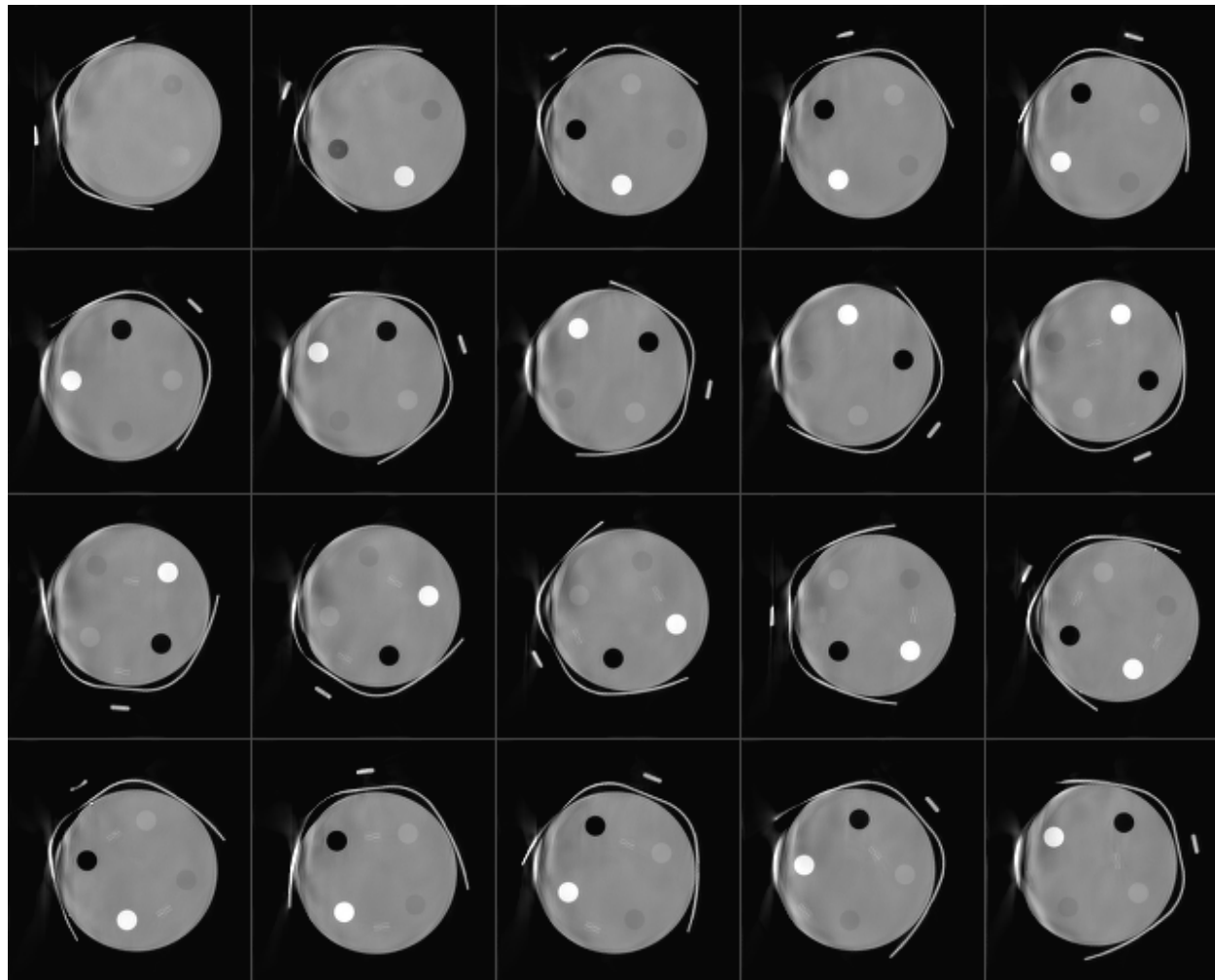
W/L: 100/1050 HU

Results



W/L: 200/0 HU

Results

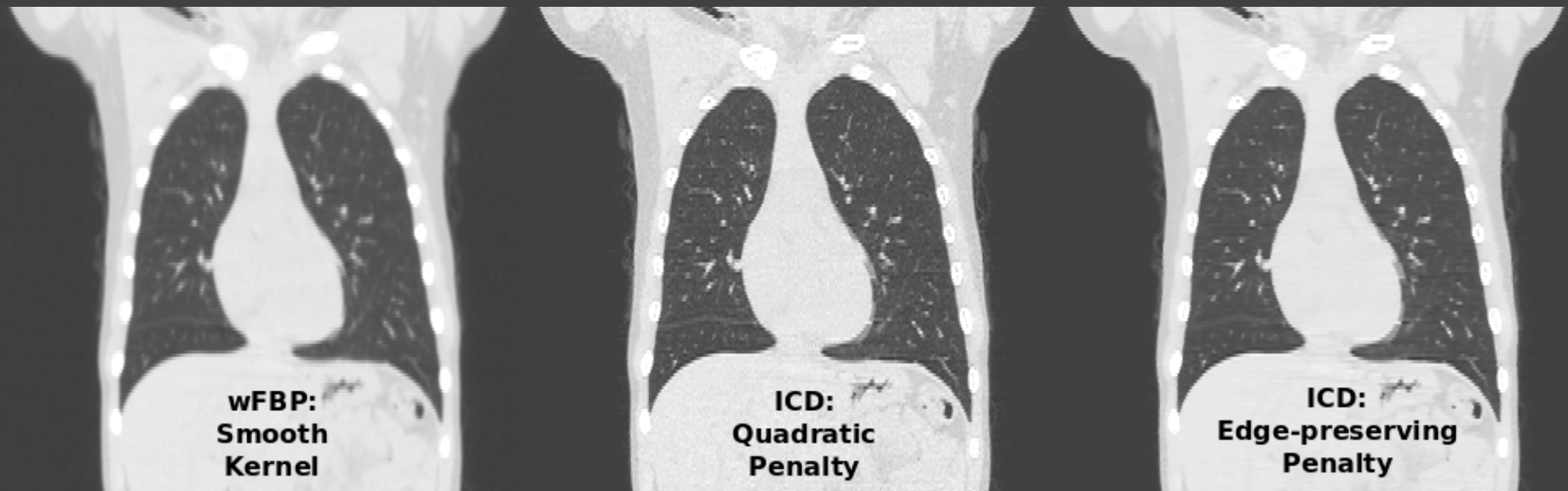


Results

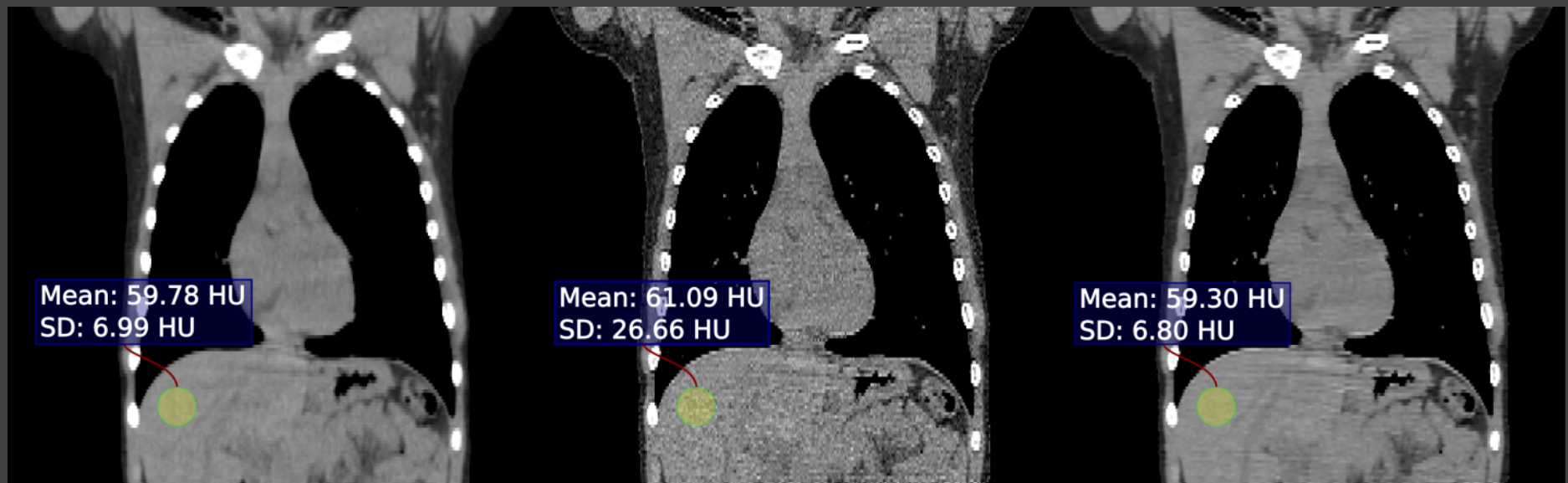
- Pediatric chest

| 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 x 1.5 |
| FOV radius [mm] | 500 |
| Edge-preserving parameter | 0.005 |
| Penalty term parameter | 0.1 |
| Iterations | 50 |
| Matrix size [GB] | 14.6 |

Results



Results

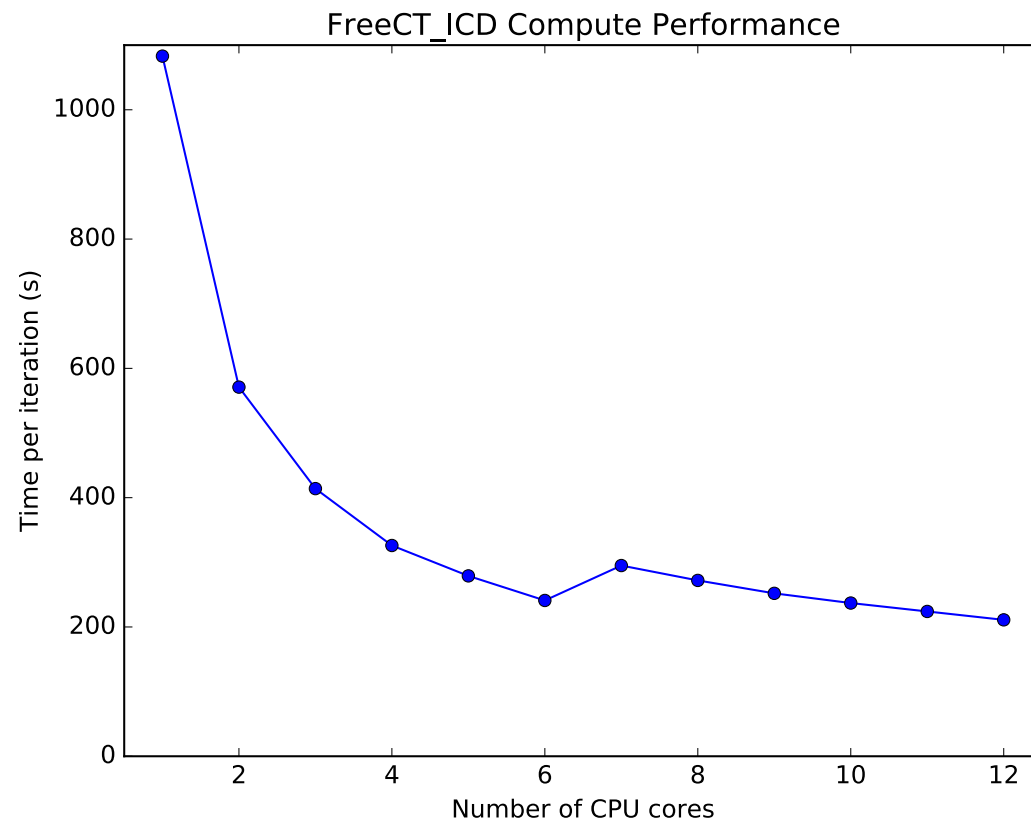


Results

- 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
- 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
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Thank you! Questions?