

Deep Learning-Based Detector Row Upsampling to Reduce Windmill Artifacts in Diagnostic Spiral CT

Jan Magonov^{1,2,3}, Julien Erath^{1,2,3}, Joscha Maier^{1,3}, Eric Fournié²,
Karl Stierstorfer², and Marc Kachelrieß^{1,3}

¹German Cancer Research Center (DKFZ), Heidelberg, Germany

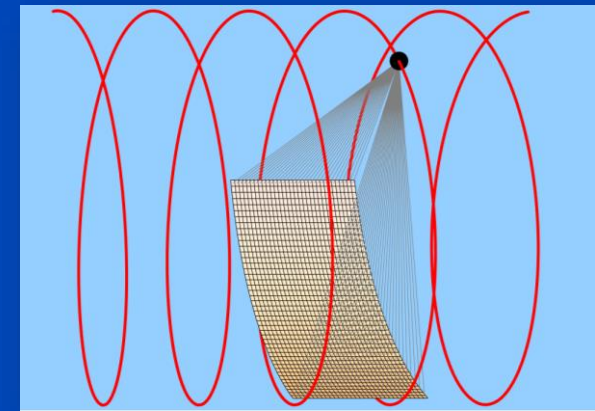
²Siemens Healthineers, Forchheim, Germany

³Heidelberg University, Heidelberg, Germany

www.dkfz.de/ct

Windmill Artifacts in Spiral CT

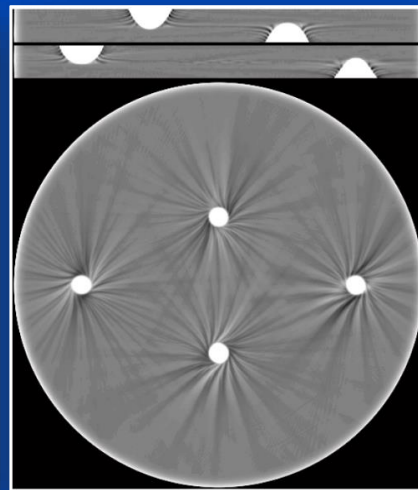
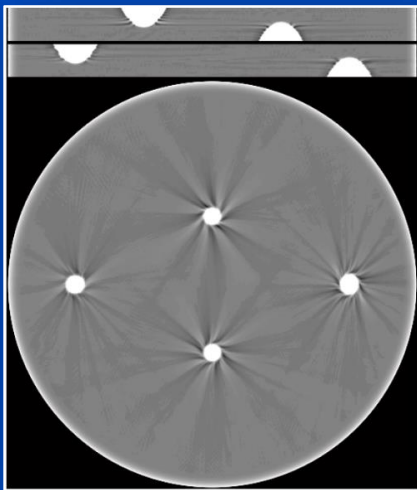
- Interpolation between adjacent detector rows is performed during backprojection in multislice spiral CT.
- Not satisfying the Nyquist criterion due to longitudinal undersampling leads to the so-called windmill artifacts.
- They are characterized by streaks diverging from a focal high-density structure.



$C = 0$ HU, $W = 200$ HU
(32×0.6 mm)

Pitch: 0.5

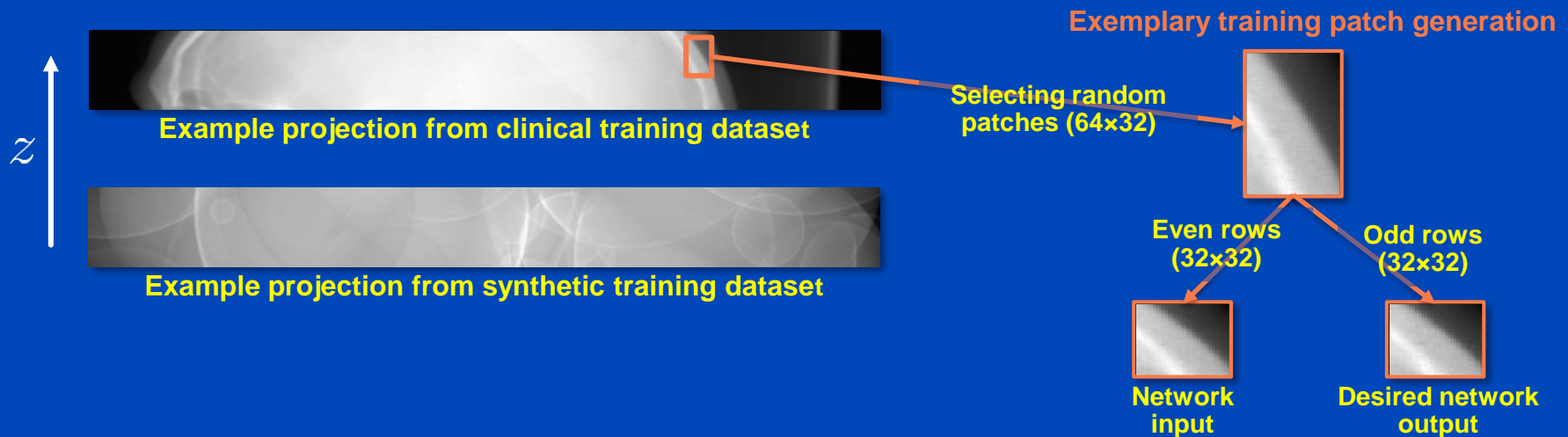
Pitch: 1.0



Reconstructions of a phantom scanned with a Siemens Somatom Sensation 16 CT system (16×0.75 mm)

Row Interpolation with Deep Learning (RIDL)

- CNN trained to predict upsampled rows for a given input.
- Previously presented network architecture¹ was further simplified.
- An experimental synthetic dataset was provided².
- Training two separate RIDL-CNN networks:
 - **Clinical** dataset consisting of projection data from patient scans acquired with zFFS.
 - **Synthetic** dataset consisting of simulated noise-free projection data of spherical shells.
- Value range of synthetic data linearly scaled to range of clinical data.



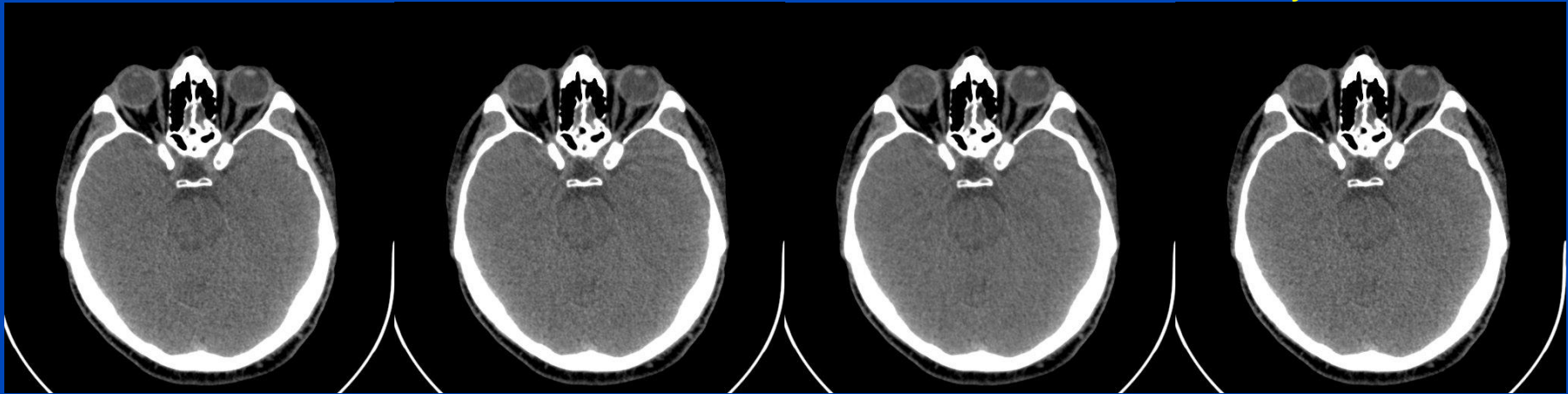
Comparison of Windmill Artifact Reduction

Ground truth
with zFFS

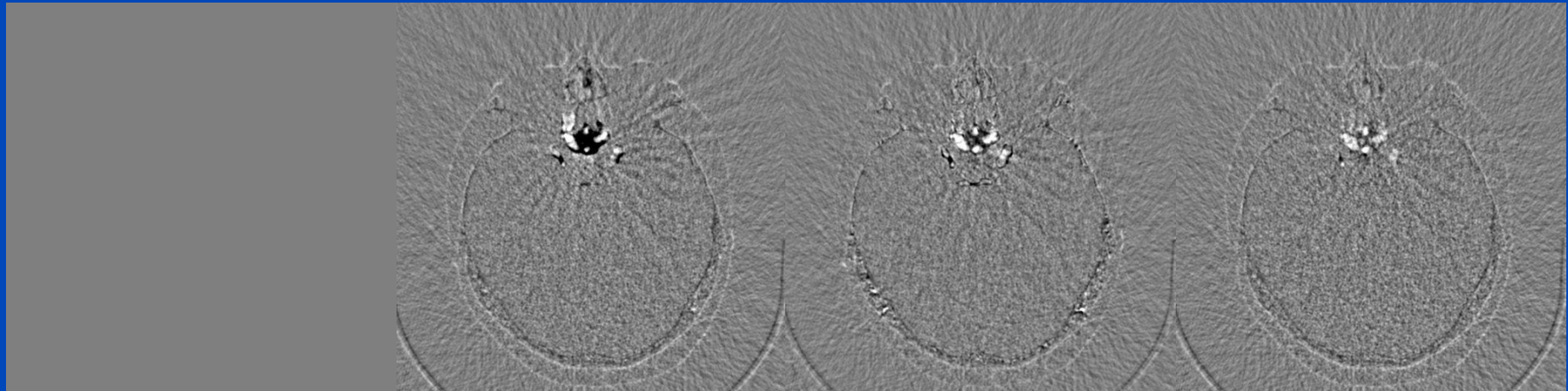
Without zFFS

RIDL-CNN trained with
clinical dataset

RIDL-CNN trained with
synthetic dataset



C = 60 HU, W = 260 HU; collimation: 64×0.6 mm;
reconstructed slice width 0.6 mm



C = 0 HU, W = 100 HU

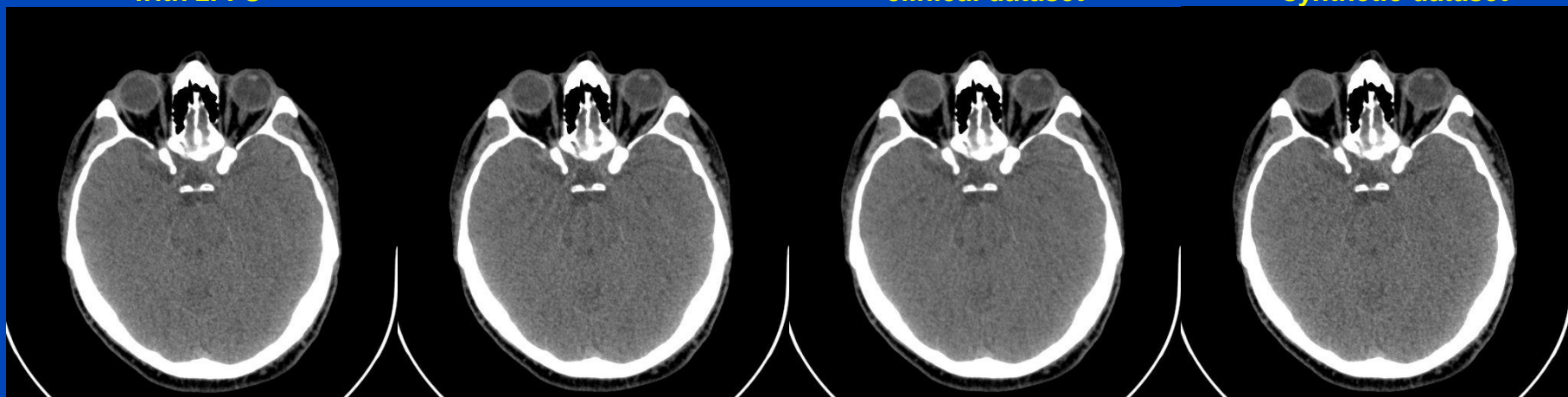
Comparison of Windmill Artifact Reduction

Ground truth
with zFFS

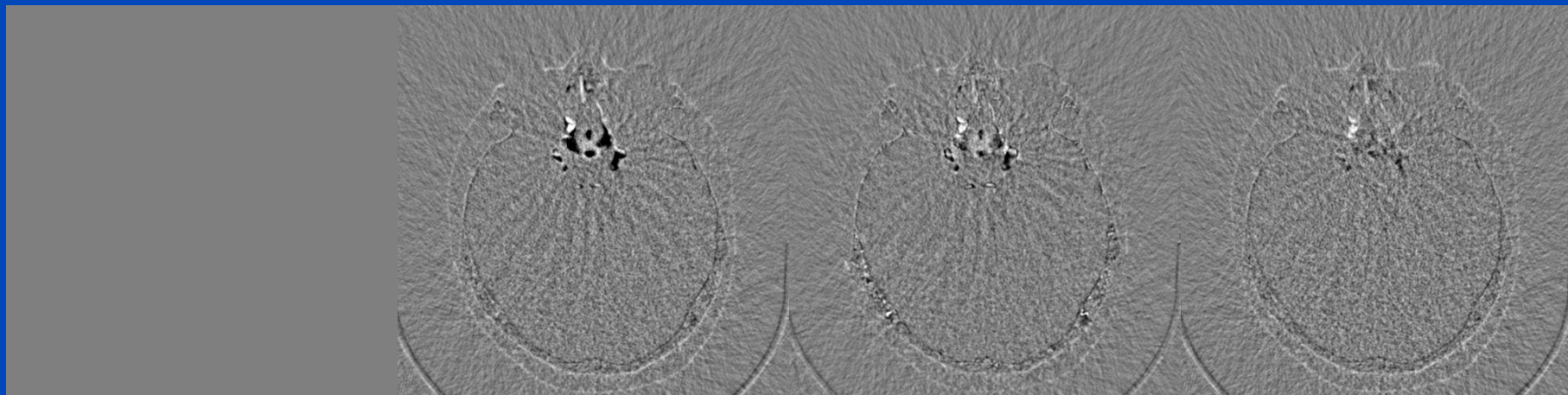
Without zFFS

RIDL-CNN trained with
clinical dataset

RIDL-CNN trained with
synthetic dataset



$C = 60 \text{ HU}$, $W = 260 \text{ HU}$; collimation: $64 \times 0.6 \text{ mm}$;
reconstructed slice width 0.6 mm



$C = 0 \text{ HU}$, $W = 100 \text{ HU}$

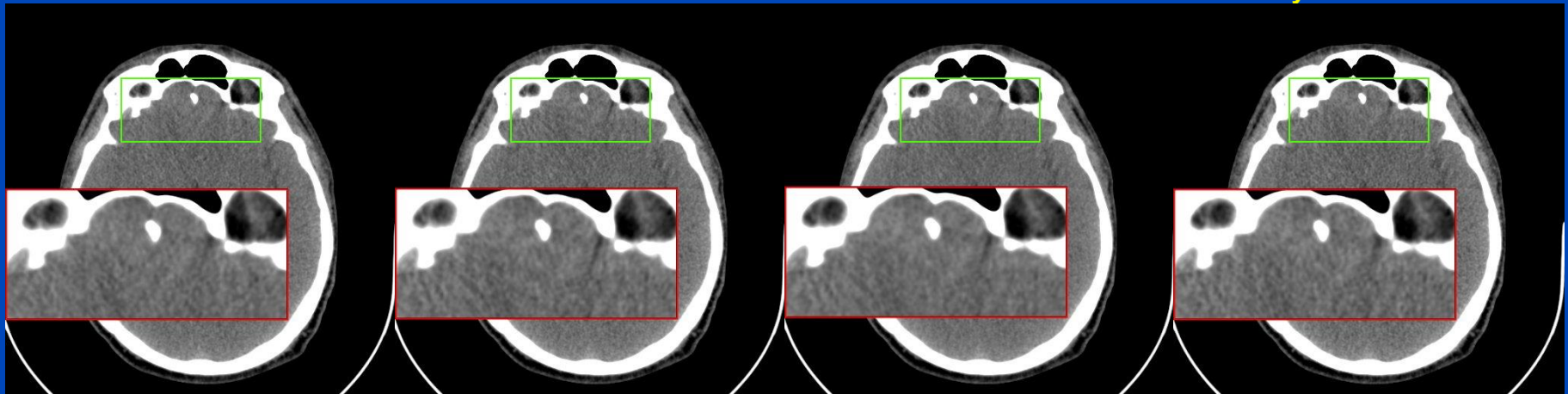
Comparison of Windmill Artifact Reduction

Ground truth
with zFFS

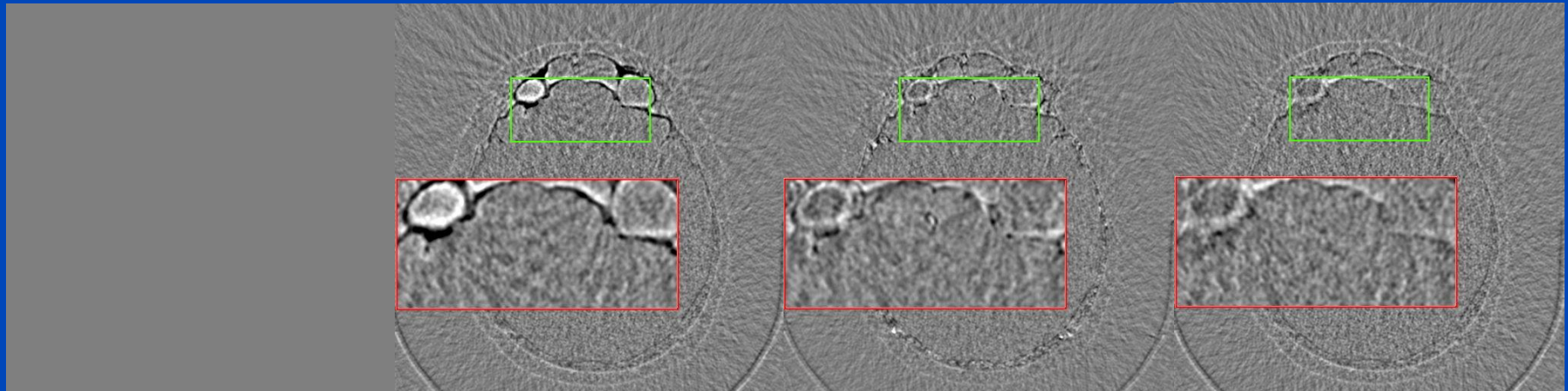
Without zFFS

RIDL-CNN trained with
clinical dataset

RIDL-CNN trained with
synthetic dataset



$C = 60 \text{ HU}$, $W = 260 \text{ HU}$; collimation: $64 \times 0.6 \text{ mm}$;
reconstructed slice width 0.6 mm



$C = 0 \text{ HU}$, $W = 100 \text{ HU}$

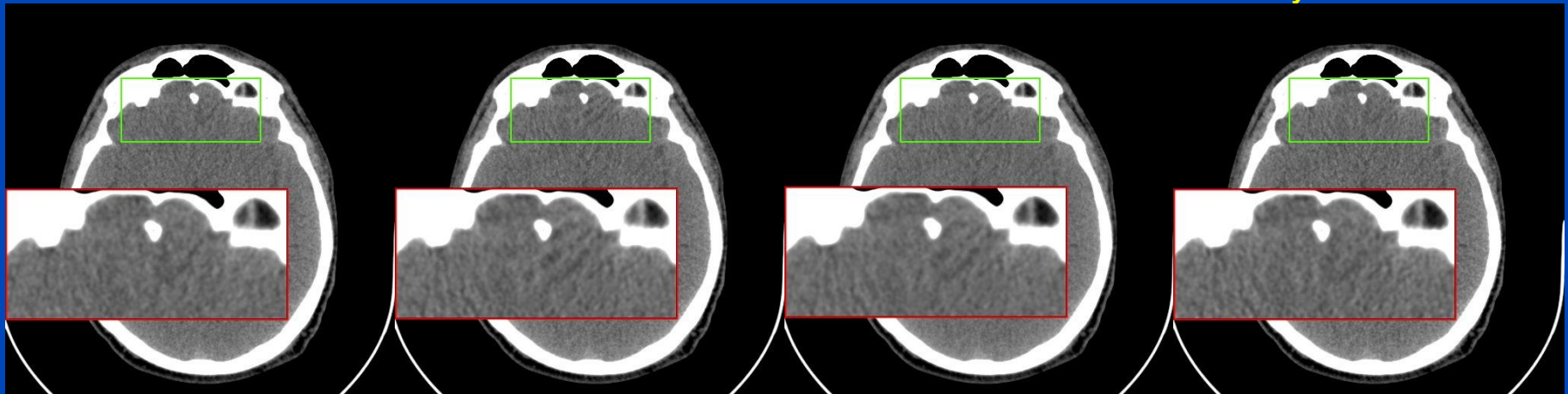
Comparison of Windmill Artifact Reduction

Ground truth
with zFFS

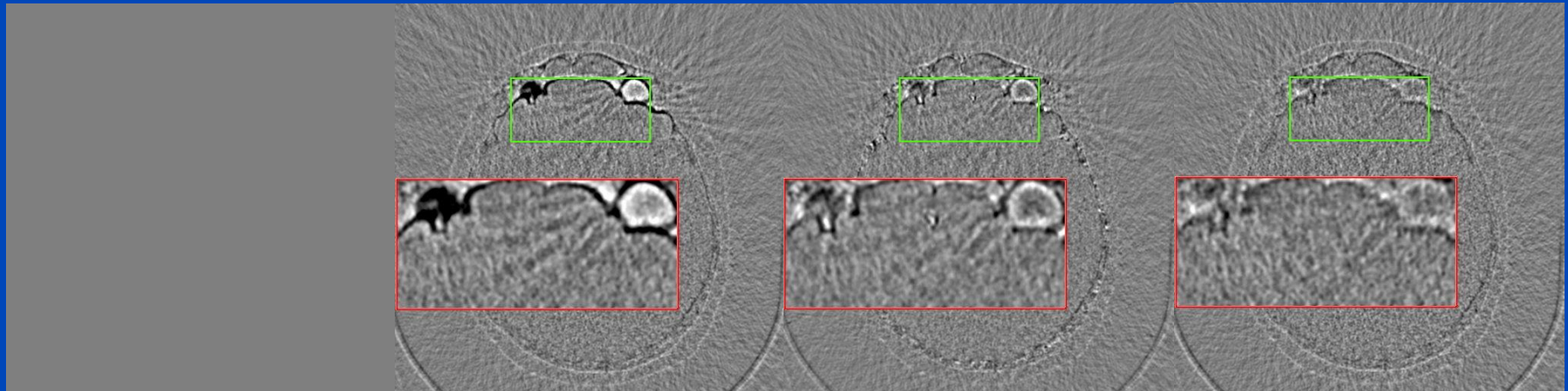
Without zFFS

RIDL-CNN trained with
clinical dataset

RIDL-CNN trained with
synthetic dataset



$C = 60 \text{ HU}$, $W = 260 \text{ HU}$; collimation: $64 \times 0.6 \text{ mm}$;
reconstructed slice width 0.6 mm



$C = 0 \text{ HU}$, $W = 100 \text{ HU}$

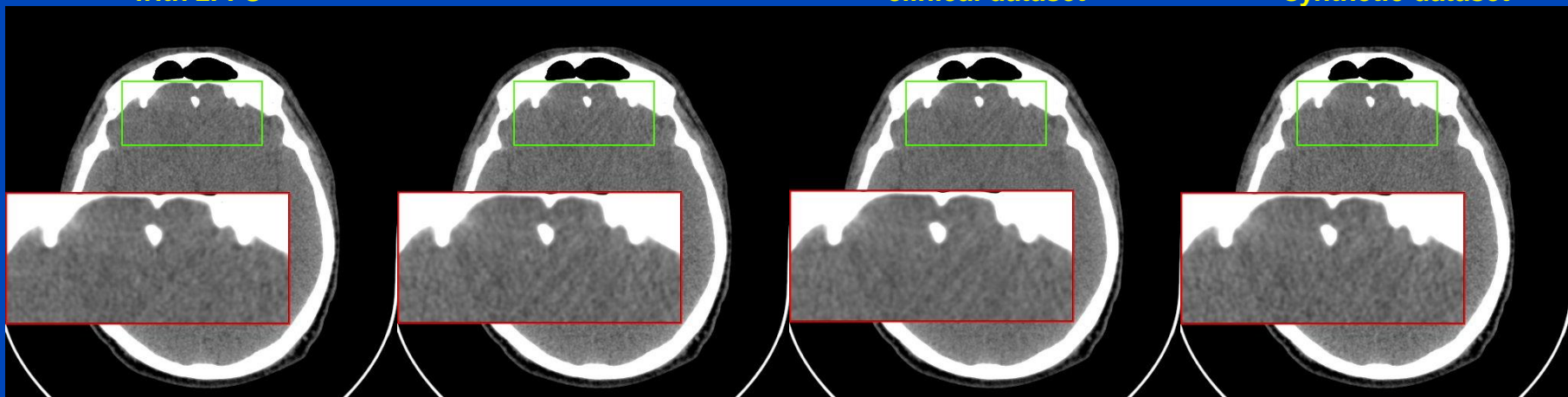
Comparison of Windmill Artifact Reduction

Ground truth
with zFFS

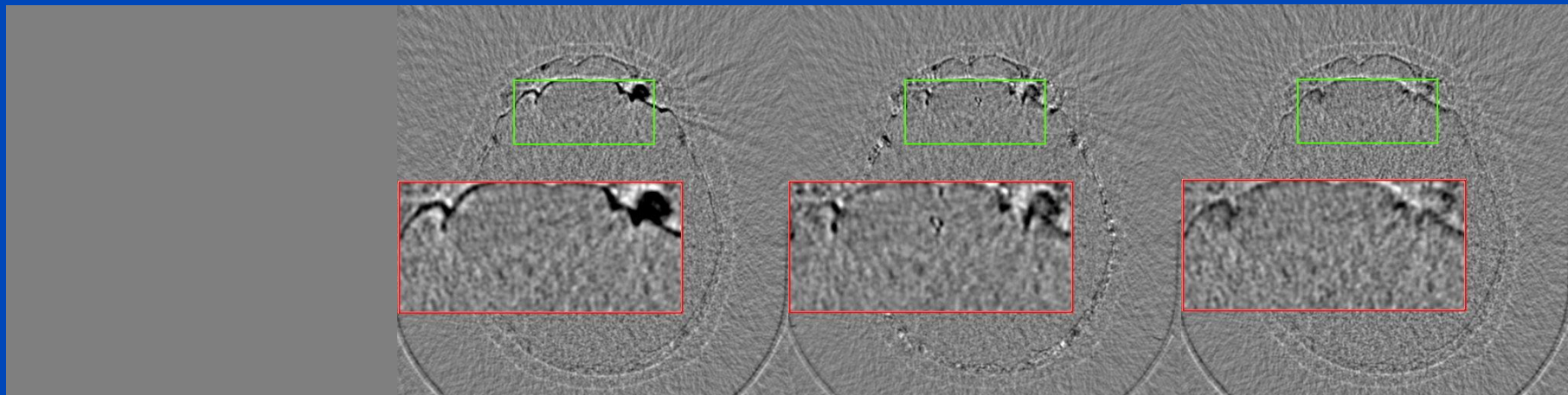
Without zFFS

RIDL-CNN trained with
clinical dataset

RIDL-CNN trained with
synthetic dataset



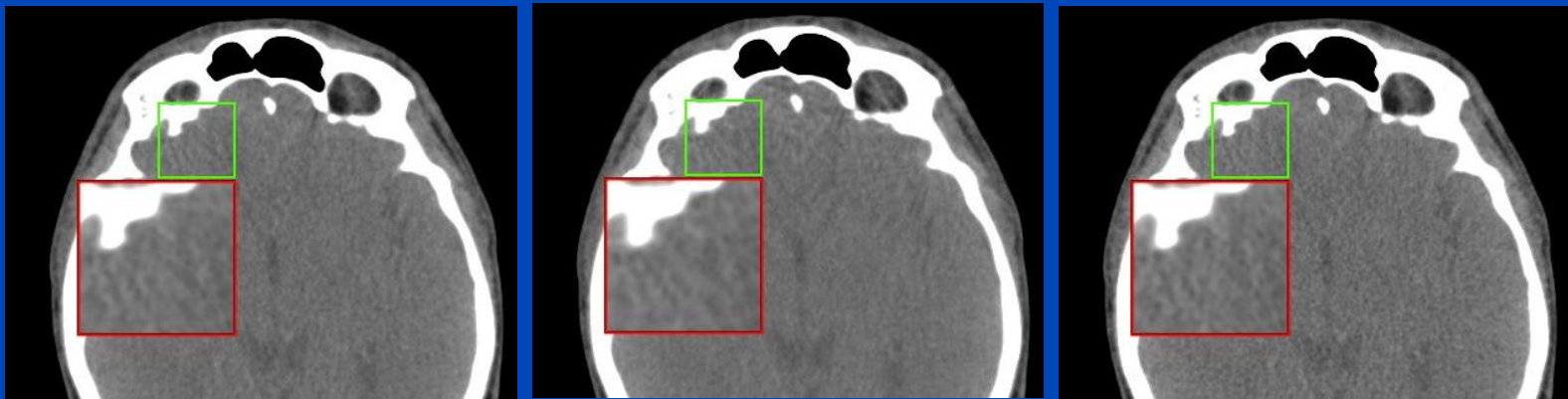
$C = 60$ HU, $W = 260$ HU; collimation: 64×0.6 mm;
reconstructed slice width 0.6 mm



$C = 0$ HU, $W = 100$ HU

Conclusions

- RIDL significantly reduces windmill artifacts.
- However, it cannot outperform zFFS.
- But it does not require special hardware.
- Training with noise-free synthetic data leads to superior performance!
 - This can, probably, be attributed to noise in the clinical data.
 - Further evaluation to be performed with more patient scans.



RIDL

trained on:

noisy clinical data

noisy synthetic data

noise-free synthetic data

Thank You!

This presentation will soon be available at www.dkfz.de/ct

Job opportunities through DKFZ's international PhD or Postdoctoral Fellowship programs (www.dkfz.de), or directly through Prof. Dr. Marc Kachelrieß (marc.kachelriess@dkfz.de).