

BIOMEDICAL IMAGING & ANALYTICS

Seminar 4. Image reconstruction and MRI

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What is MRI?

- Non-invasive medical imaging technique with **excellent soft-tissue contrast**
- Does **not involve X-rays** or the use ionizing **radiation**, which makes it harmless (just keep magnetizable objects aside)
- Fun fact: MRI was initially called NMRI (nuclear magnetic resonance imaging) but “nuclear” was dropped to avoid negative associations

How to get an MRI image?



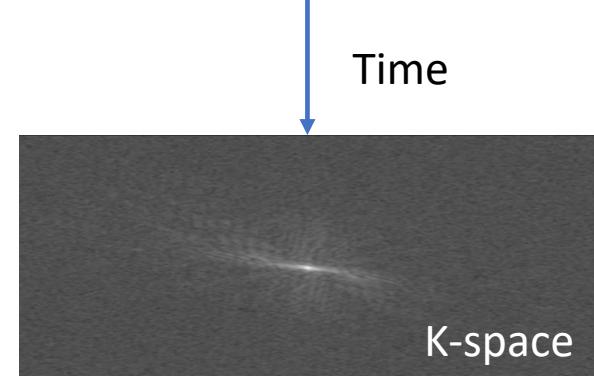
In the **simplest** case we need to:

1. Place a patient in the scanner

How to get an MRI image?

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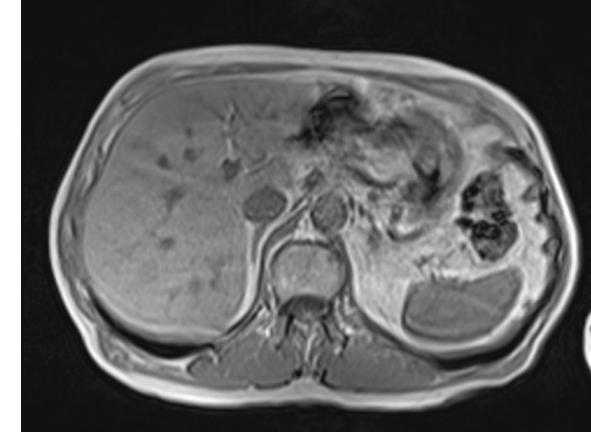
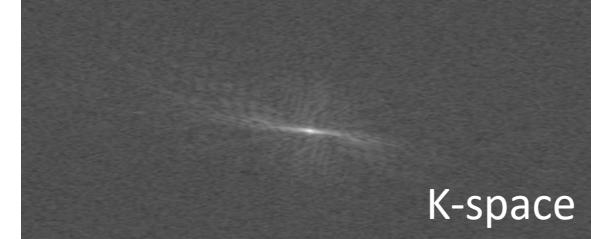
1. Place a patient in the scanner
2. Acquire raw data – **k-space** (wait and do not move)



How to get an MRI image?

In the **simplest** case we need to:

1. Place a patient in the scanner
2. Acquire raw data – **k-space** (wait and do not move)
3. Reconstruct an image using inverse **Fast Fourier Transform (iFFT)**



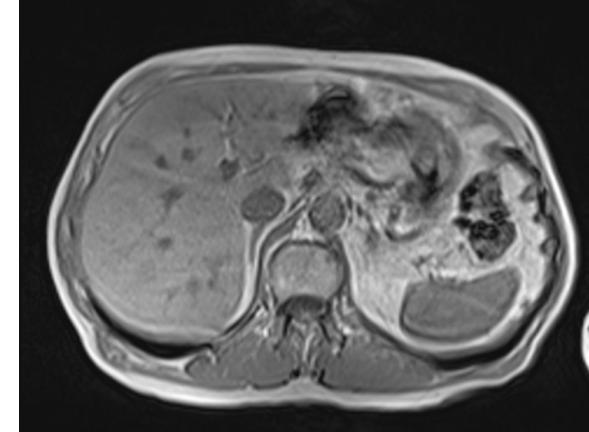
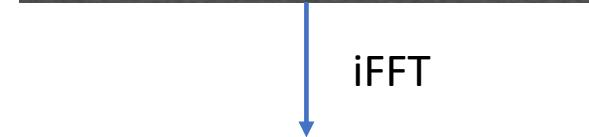
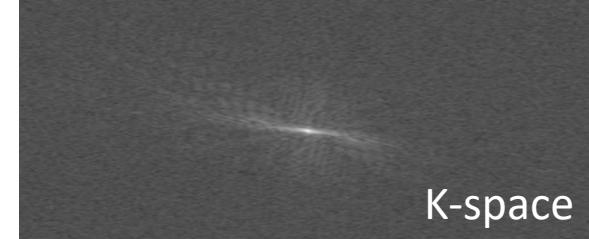
How to get an MRI image?

In the **simplest** case we need to:

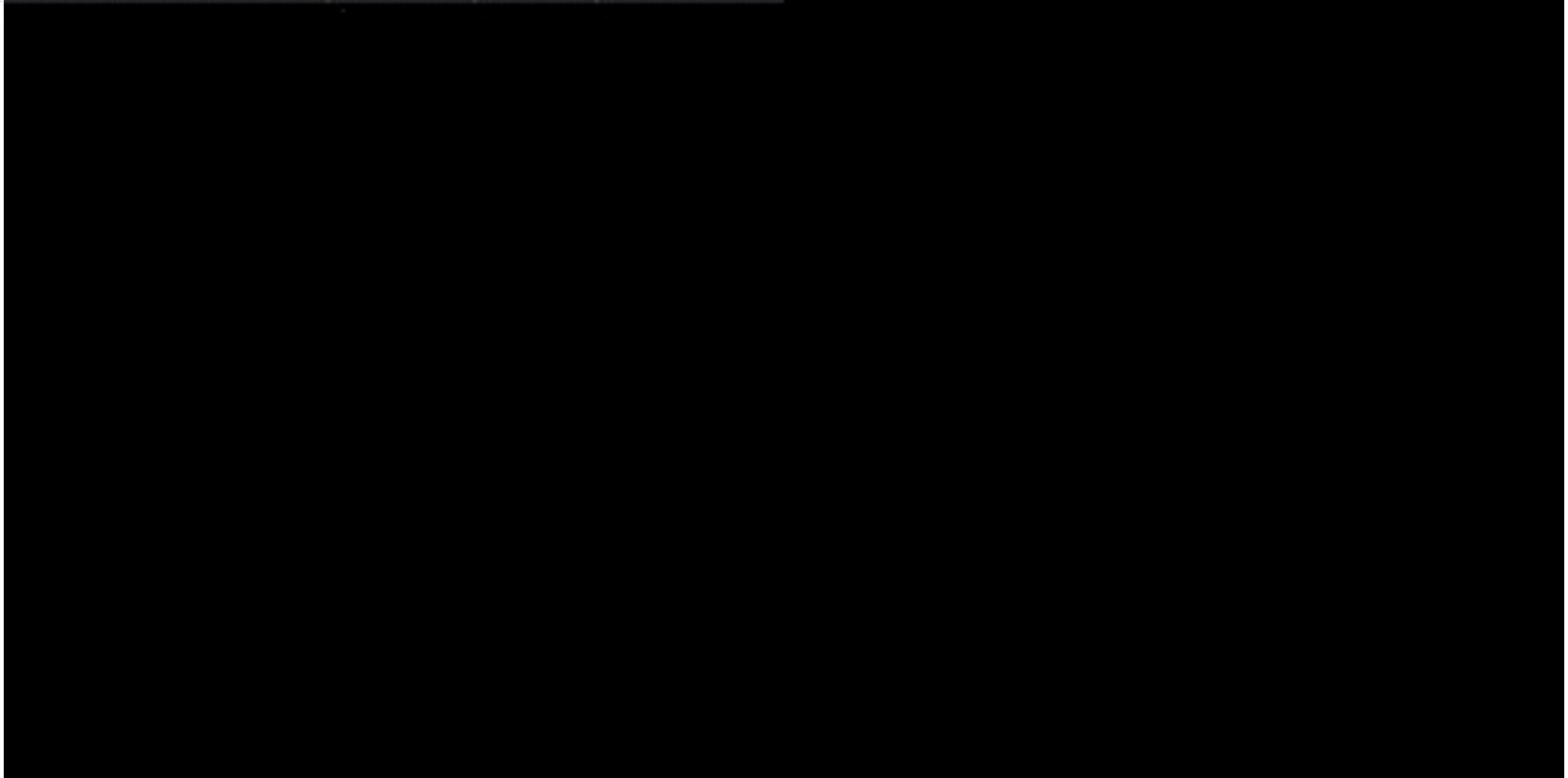
1. Place a patient in the scanner
2. Acquire raw data – **k-space** (wait and do not move)
3. Reconstruct an image using inverse **Fast Fourier Transform (iFFT)**

Problem

Data acquisition takes **a lot of time**: typically, from 20 to 60 min

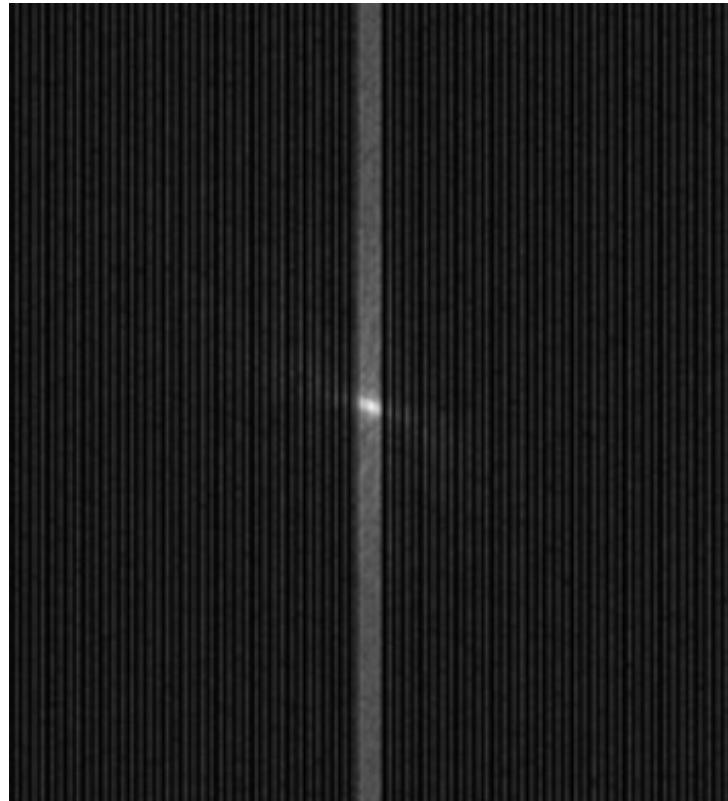


Why so long?



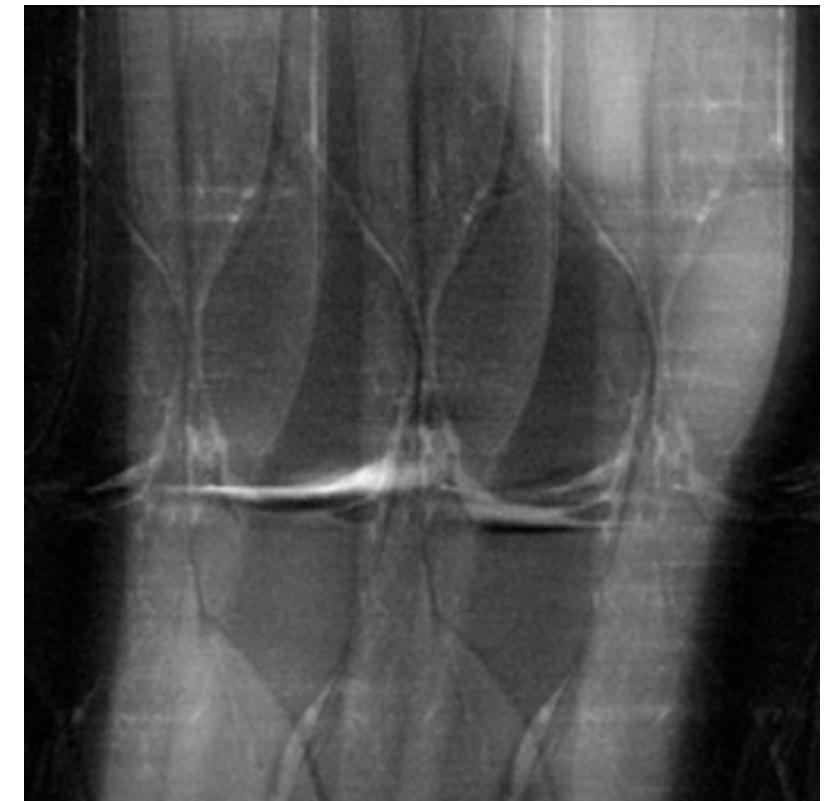
Standard sequential Cartesian filling of k-space (Courtesy of Brian Hargreaves)

Problem statement



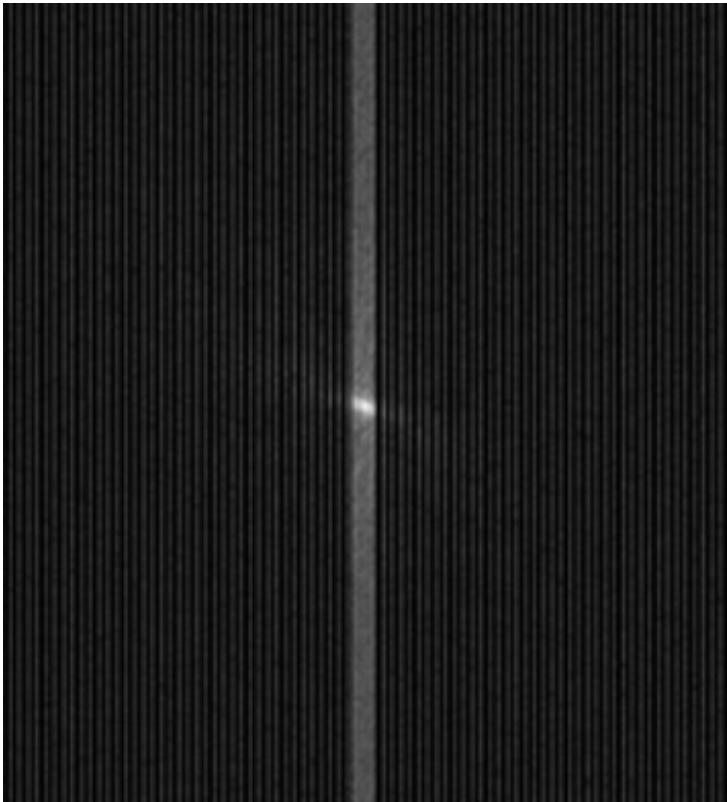
k-space

Standard
reconstruction



Reconstructed image

Problem statement



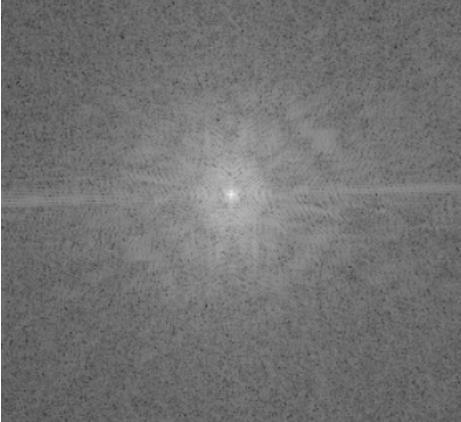
k-space



Reconstructed image

Accelerating Magnetic Resonance Imaging

k-space



iFFT 



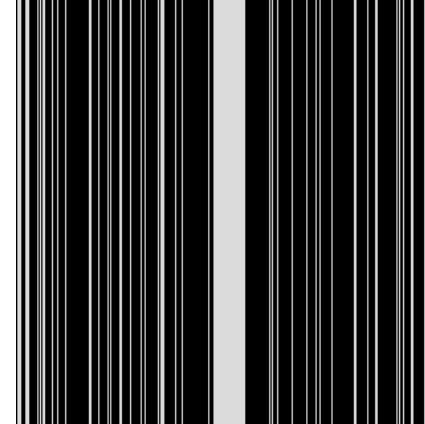
Fully-Sampled recon

Accelerating Magnetic Resonance Imaging

k-space



Sampling mask



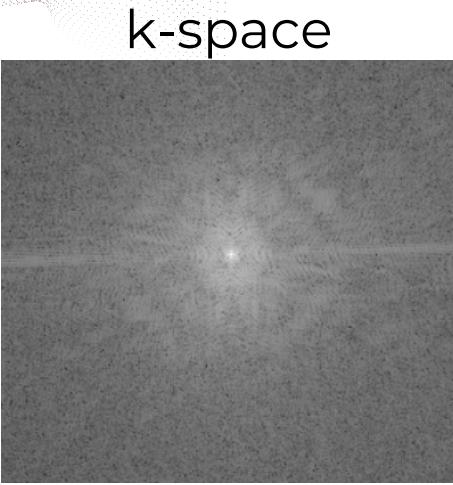
X

iFFT



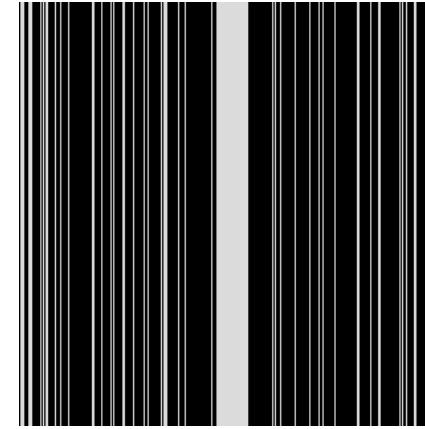
Fully-Sampled recon

Accelerating Magnetic Resonance Imaging



Sampling mask

X



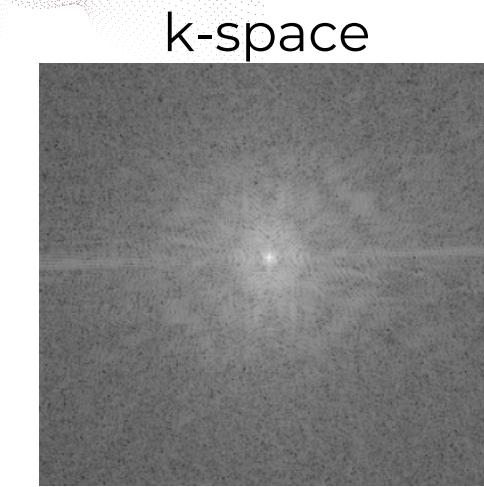
iFFT
→

Zero-filled recon



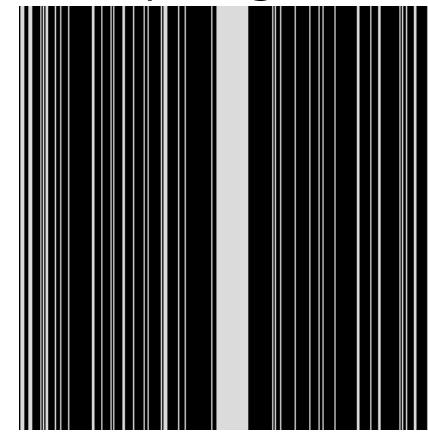
Fully-Sampled recon

Accelerating Magnetic Resonance Imaging



Sampling mask

X



iFFT
→

Zero-filled recon



iFFT
↓



Fully-Sampled recon



Our recon

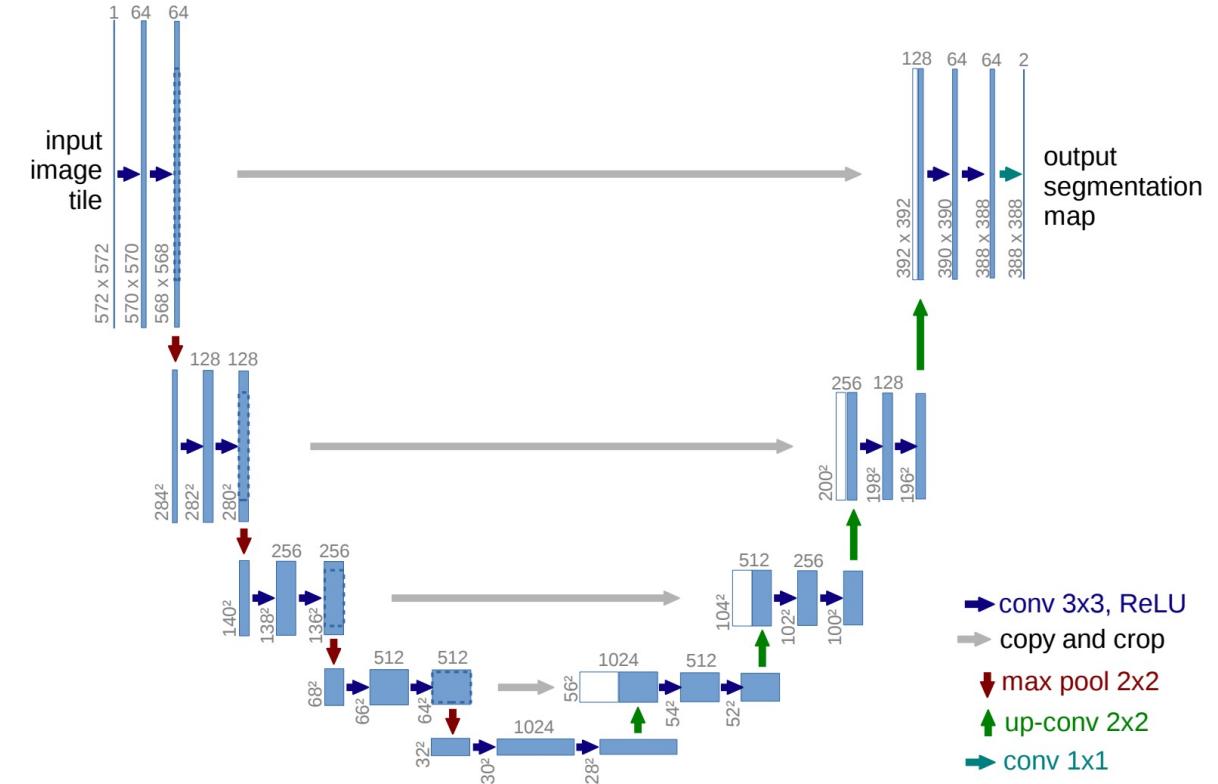
←

Our algorithm



What algorithm to choose? A baseline approach

- U-Net [1] architecture is an easy first candidate for virtually any image-to-image task
- Copy-pasted between codebases since 2015
- No one remembers why U-Net, but still use it
- Initially proposed for medical image segmentation but can be adapted for virtually any task by changing a couple of first layers



Demo 1 – Baseline approaches

Find the code in the [S4_Image_reconstruction_and_MRI_basic.ipynb](#) file
in the BIA course repository

Everyone uses U-Net, it should be enough, right?

Images from Demo 1



Zero-filled R=4

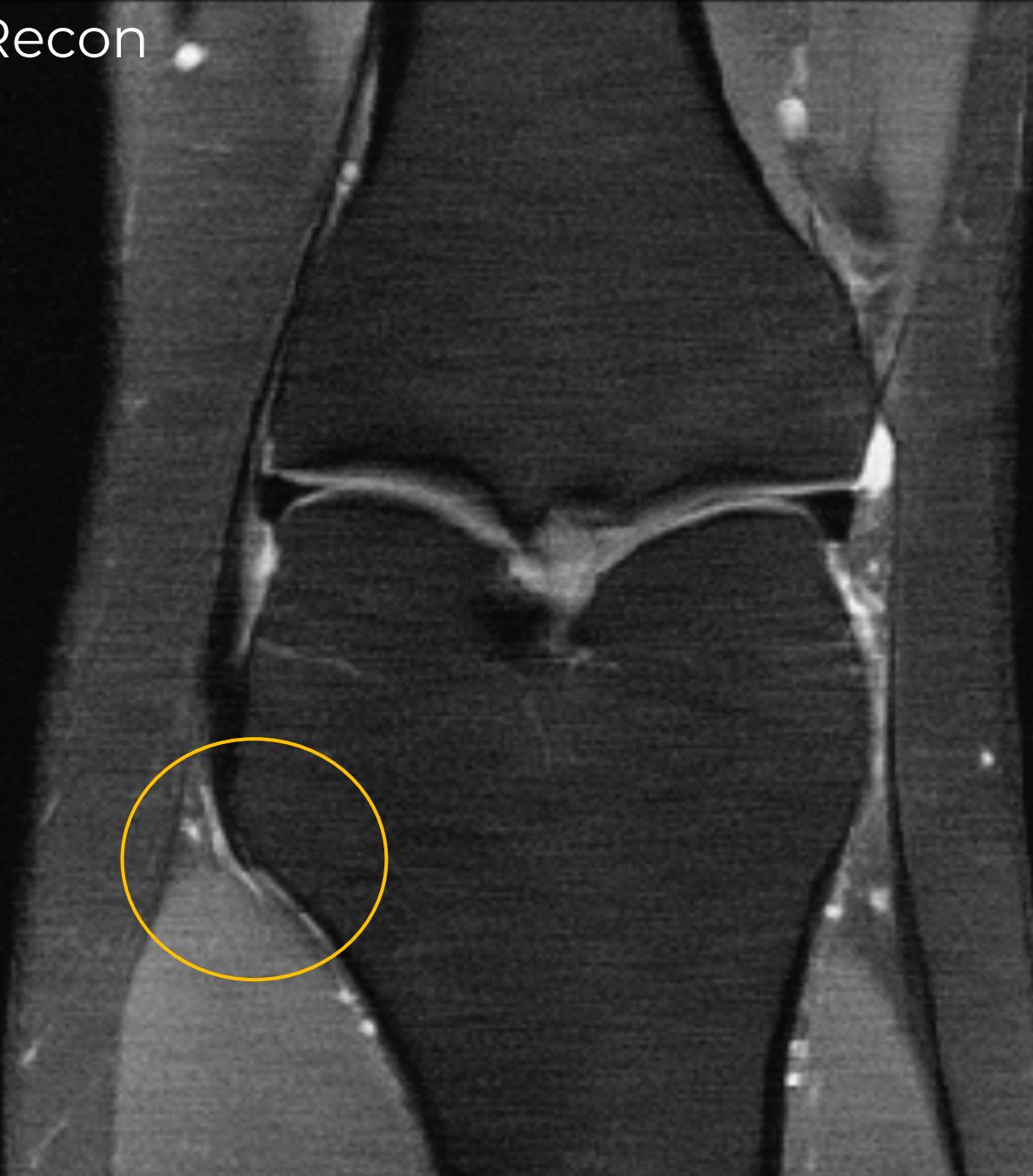


Recon

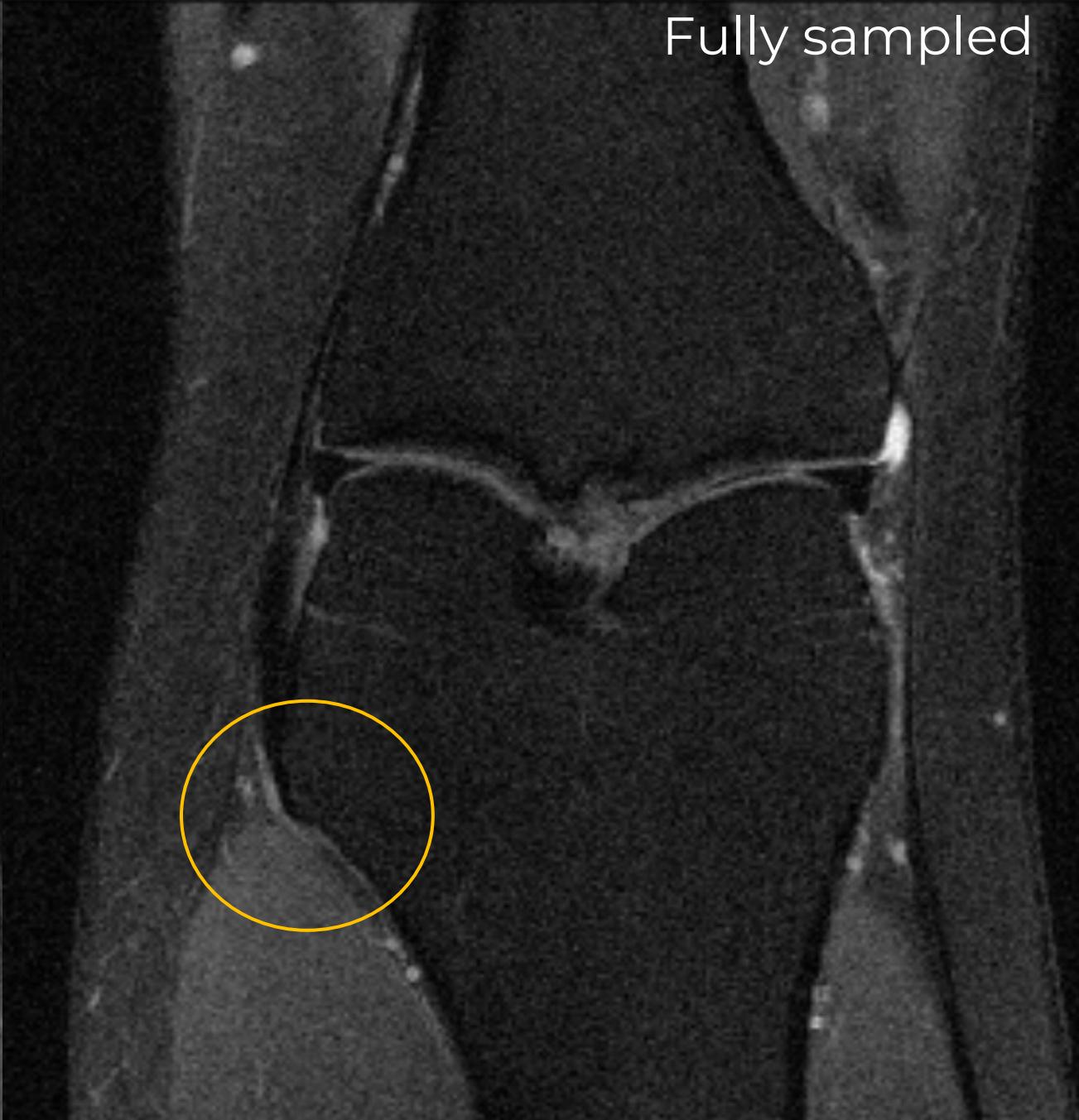


Fully sampled

Recon



Fully sampled



OK, let us use GANs then!

The original paper [2] (2015) on Generative Adversarial Networks (GANs) introduced the following objective

$$\min_G \max_D V(D, G) = \mathbb{E}_{\mathbf{x} \in p_{\text{data}}(\mathbf{x})} [\log D(\mathbf{x})] + \mathbb{E}_{\mathbf{z} \in p_{\mathbf{z}}(\mathbf{z})} [\log(1 - D(G(\mathbf{z})))] \quad (1)$$

Equation 1 can be simplified as follows:

$$\begin{aligned} V(D, G) &= \int_{\mathbf{x}} p_{\text{data}}(\mathbf{x}) \log(D(\mathbf{x})) d\mathbf{x} + \int_{\mathbf{z}} p_{\mathbf{z}}(\mathbf{z}) \log(1 - D(G(\mathbf{z}))) d\mathbf{z} \\ &= \int_{\mathbf{x}} p_{\text{data}}(\mathbf{x}) \log(D(\mathbf{x})) + p_g(\mathbf{x}) \log(1 - D(\mathbf{x})) d\mathbf{x} \end{aligned} \quad (2)$$

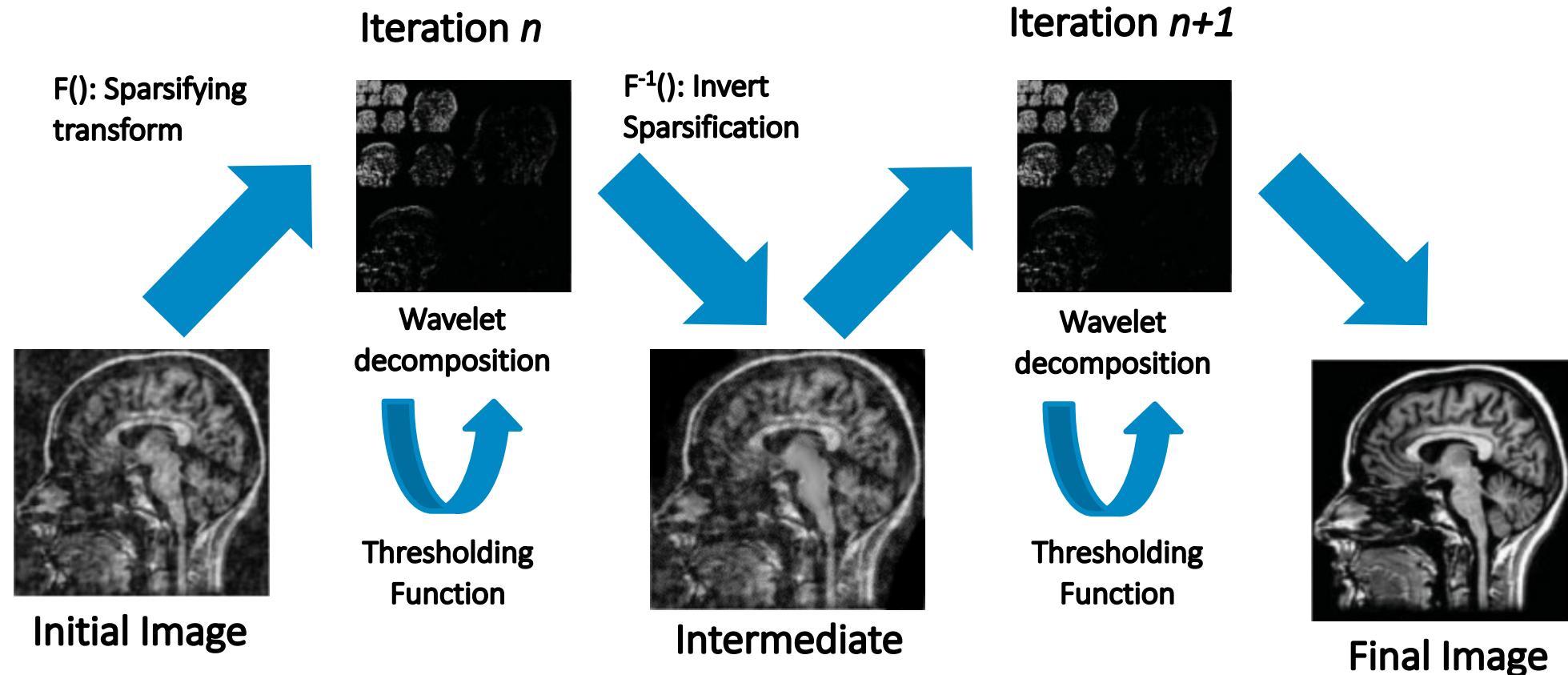
Hence, the optimal discriminator is:

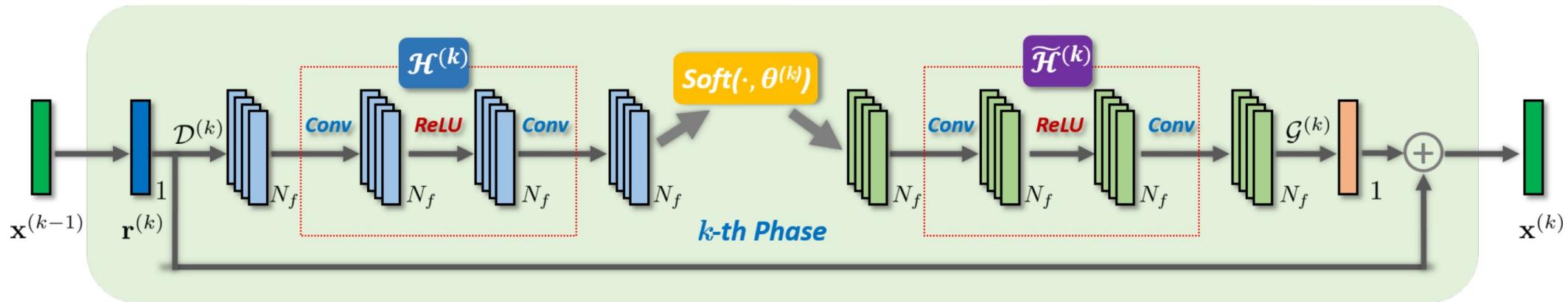
$$D_G^*(\mathbf{x}) = \frac{p_{\text{data}}(\mathbf{x})}{p_{\text{data}}(\mathbf{x}) + p_g(\mathbf{x})} \quad (3)$$

But that is **not** what we want.

We want to achieve the **best reconstruction quality** but **not make it similar** to images from train set!

Compressed Sensing for MR reconstruction (commercial SOTA)

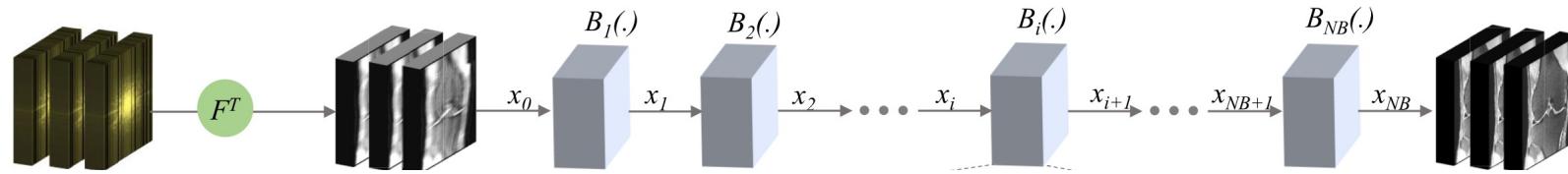




- **ISTA-Net** [3] introduced in 2017 but did not gain a lot of popularity
- **ISTA** stands for **Iterative Shrinkage-Thresholding Algorithm**
- Natural extension of iterative approaches, using **learned** blocks (convolutional layers) instead of **deterministic** (e.g., wavelet) functions
- Applies soft thresholding – threshold values are learnable parameters
- There is an extension called ISTA-Net⁺ (same authors, same paper) , where residual computations are adopted

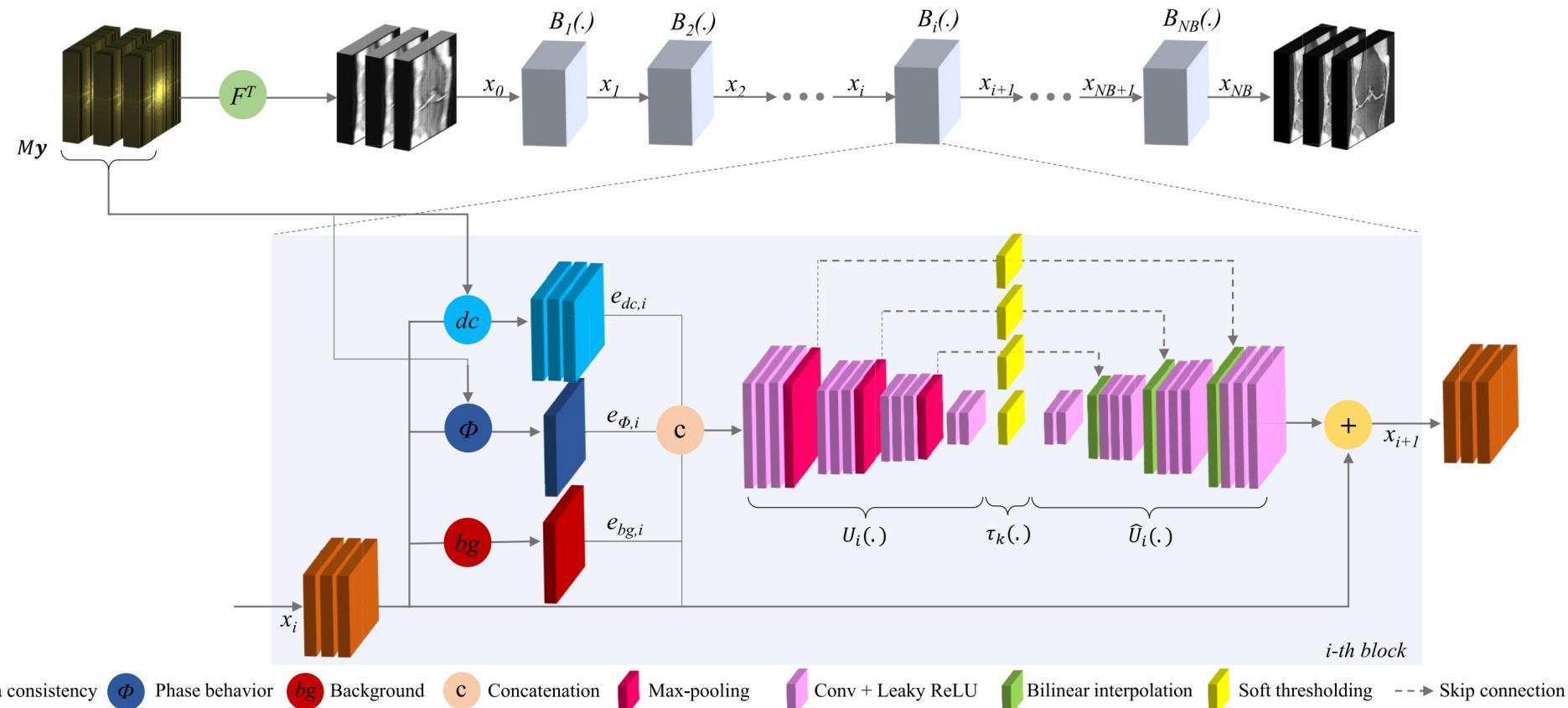
Adaptive CS-Net [4] Architecture

Winner of the 2019 fastMRI challenge [5] in 2 out of 3 most clinically-relevant tracks



Adaptive CS-Net [4] Architecture

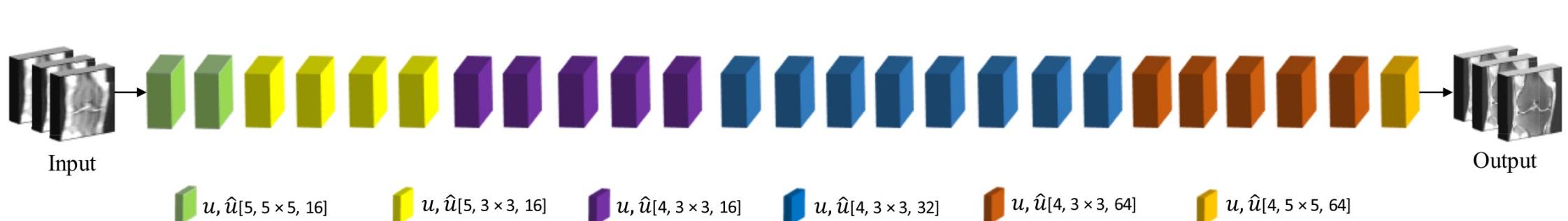
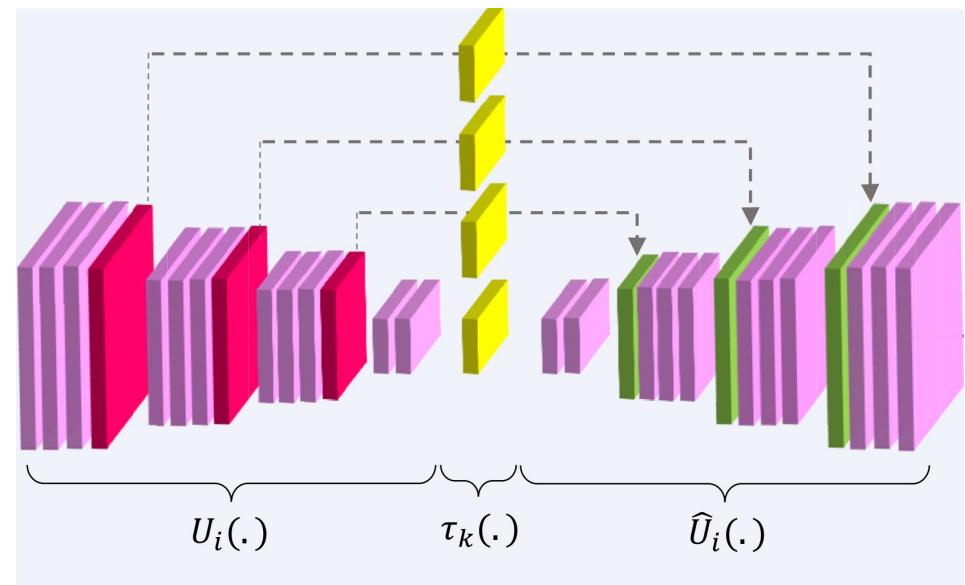
Winner of the 2019 fastMRI challenge [5] in 2 out of 3 most clinically-relevant tracks



Adaptive CS-Net [4] Building Blocks

U-Net – like architecture of fully convolutional blocks:

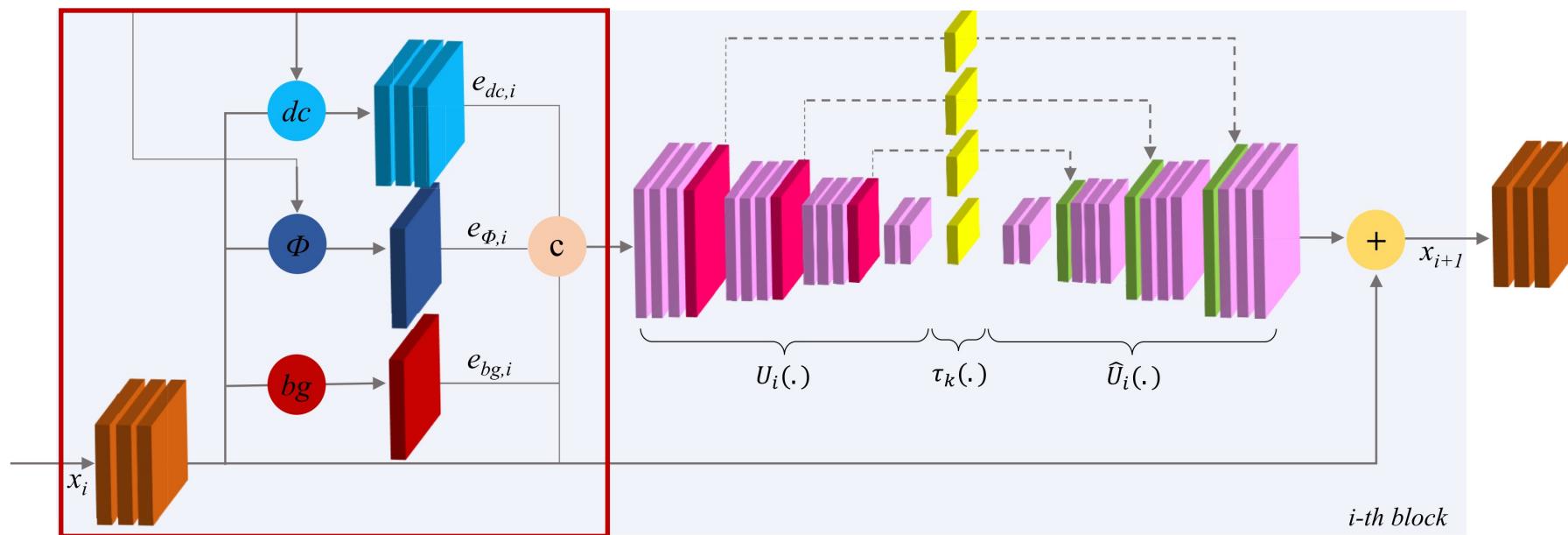
- Each feature map is filtered
- Only basic Deep Learning operations to retain control
 - 2D convolutions without bias term
 - Leaky ReLu
 - Max Pooling
 - Interpolation 2D
- 2.5D convolutions – loss applied to the central slice only



MR-specific knowledge

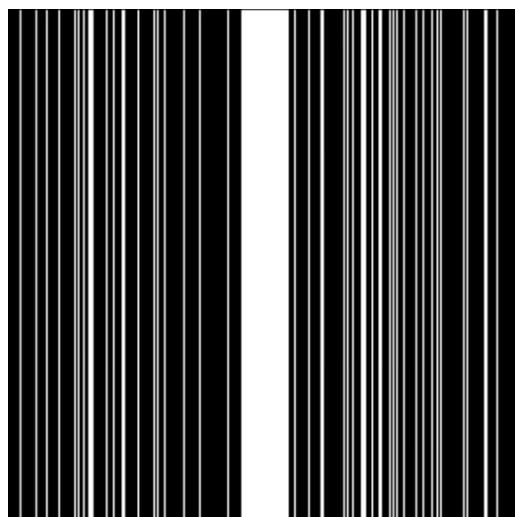
The following soft priors capture some properties on an MR image that cannot be easily learned due to the limited size of the receptive field:

- Soft data consistency
- Phase behavior
- Background information



Soft data consistency $e_{dc,i}$

- Our algorithms **do not know** what they should fix and what not
- Data consistency is a way to “undo” **unnecessary** corrections of our algorithm

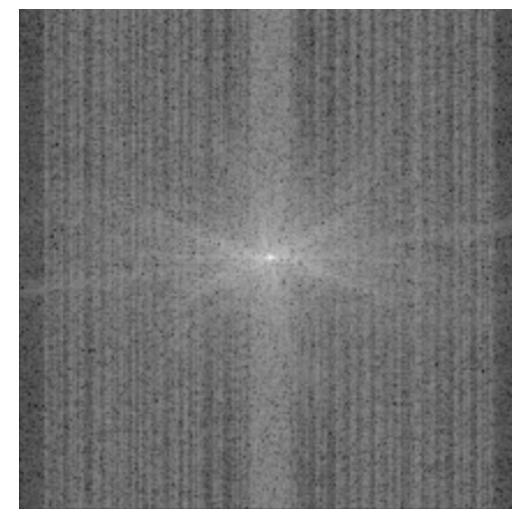


Sampling mask

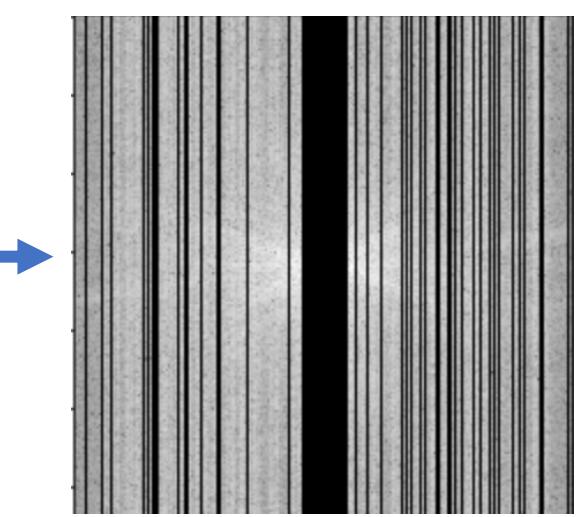


Inverted
Sampling mask

X



Recon k-space



Unnecessary
corrections

Soft data consistency $e_{dc,i}$

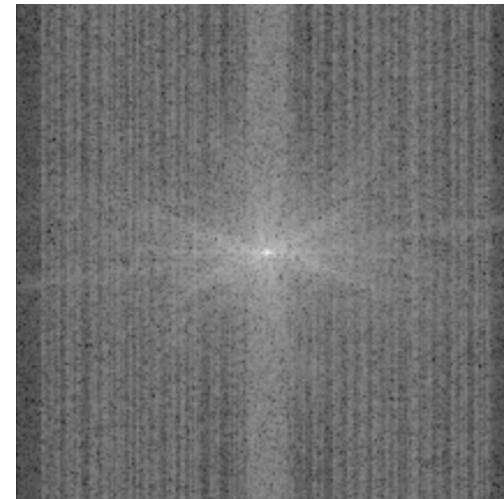
- **Data consistency**

Subtract unnecessary corrections from the recon k-space to get implement data consistency

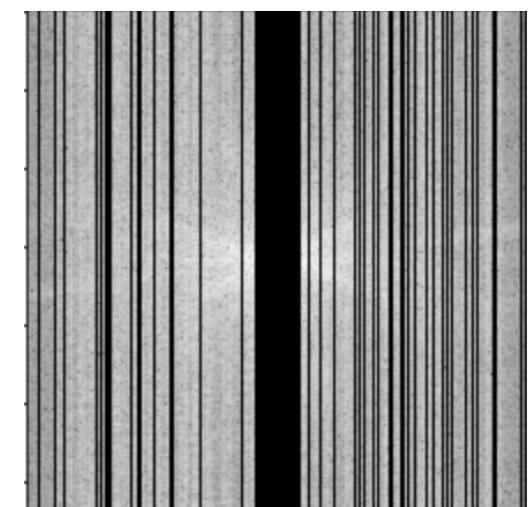
- **Soft data consistency**

Provide unnecessary corrections as an additional input for reconstruction module to let the network learn to evaluate reliability of the acquired data and potentially compensate

Data consistency



Recon k-space



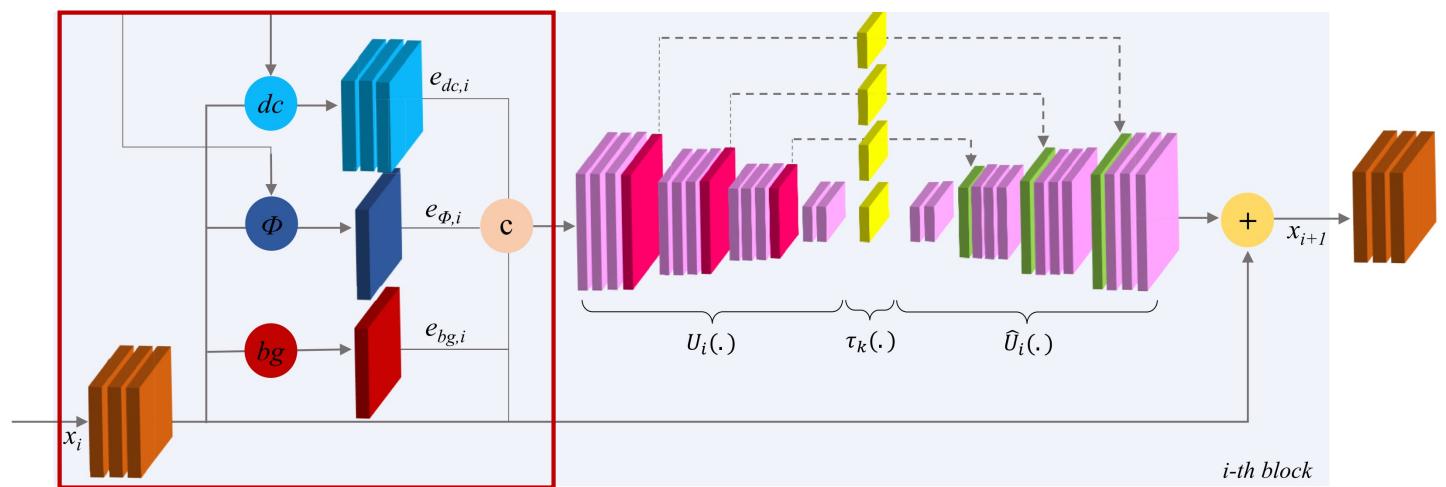
Unnecessary
corrections

Phase behavior $e_{\Phi,i}$

- Theoretically, spin-echo sequences have zero phase everywhere in the image
- In practice, slow varying phase will occur only in low frequencies due to hardware and acquisition imperfections
- It means that final recon image should be real-valued after removing this component

$$\mathbf{e}_{\phi,i} = \left\{ \mathbf{x}_i \cdot \frac{\mathbf{x}_{i,\text{lpf}}^*}{\|\mathbf{x}_{i,\text{lpf}}\|_2} \right\}_{\text{imag}},$$

where lpf refers to low pass filtering.

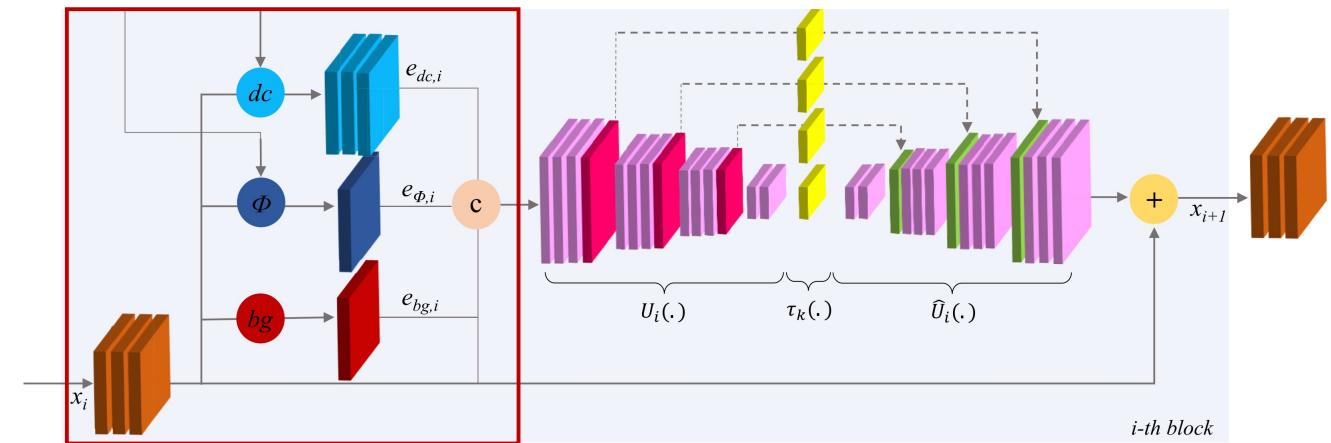


Background information $e_{bg,i}$

- Estimate the location in the \mathbf{x}_i where the background is found

$$\mathbf{e}_{bg,i} = \frac{\mathbf{x}_i}{\|\mathbf{x}_{i,\text{lpf}}\|_2}.$$

- This prior penalize estimated signal content where $\|\mathbf{x}_{i,\text{lpf}}\|$ is low, i.e., within the background
- Note that last two priors are commonly used techniques applied in parallel imaging. However, in our demo we used artificial single-coil data for simplicity, which has different properties.

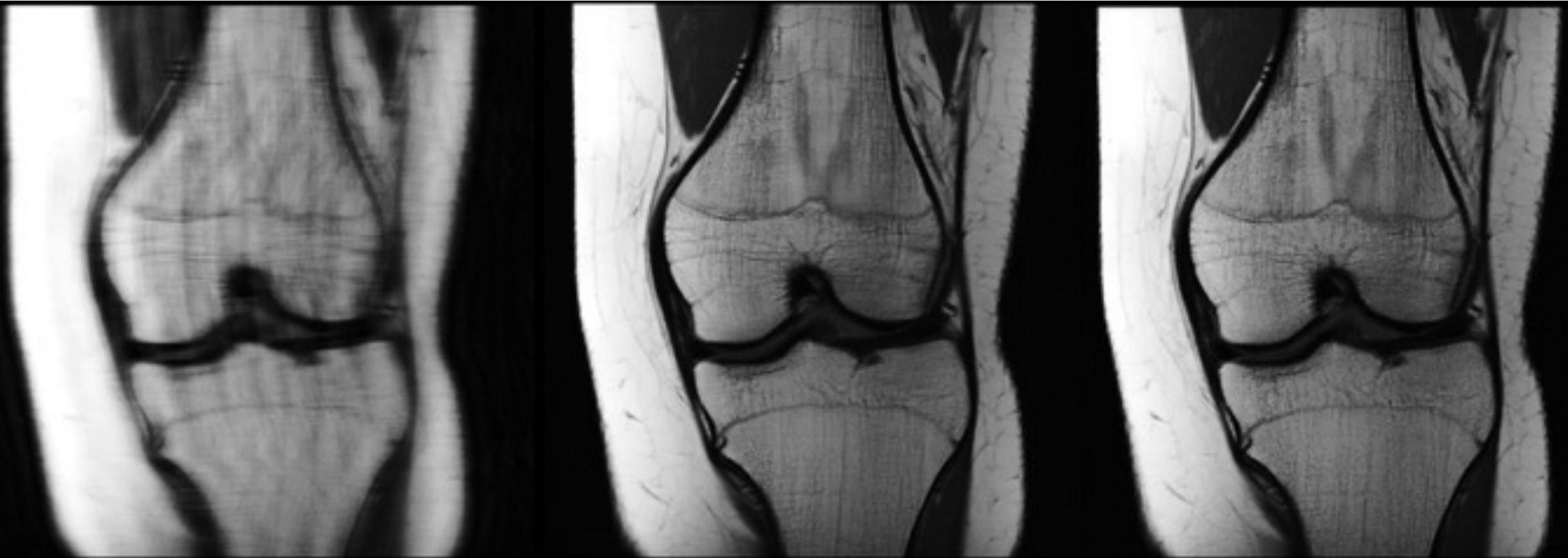


Demo 2 – Data consistency

Find the code in the [S4_Image_reconstruction_and_MRI_advanced.ipynb](#) file
in the BIA course repository.

Is it worth it? – Results from the 2019 fastMRI challenge

Effective acceleration: 3.64x – SSIM: 0.9885



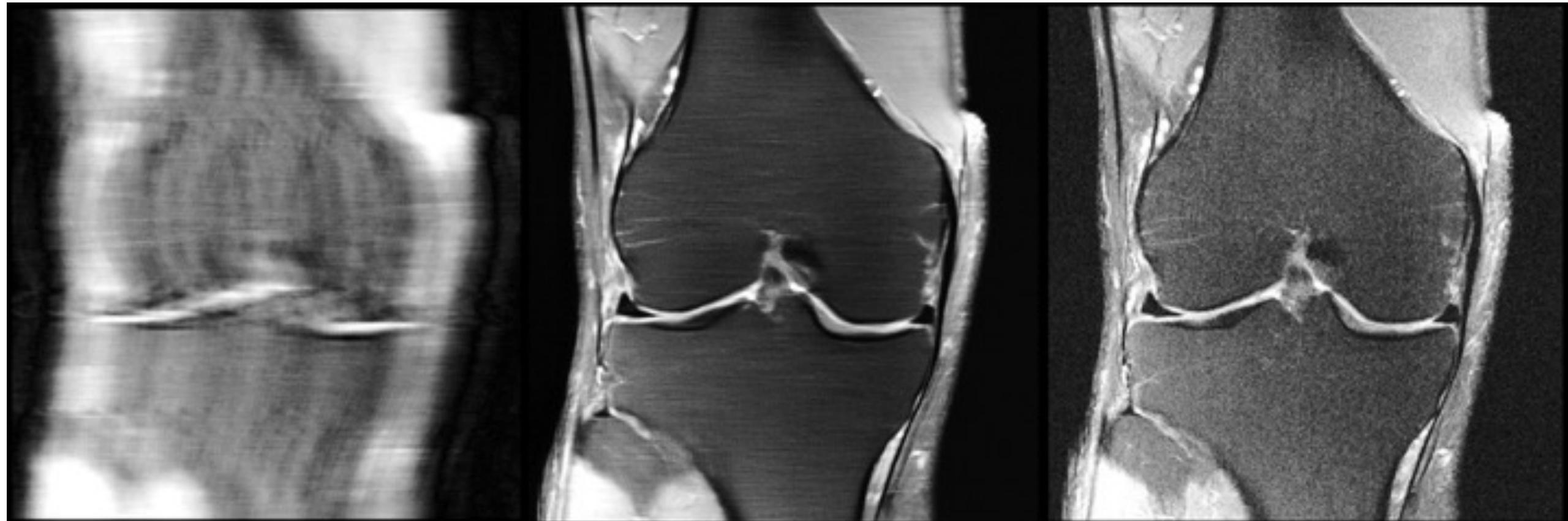
Zero-filled

Recon

Fully sampled

Is it worth it? - Results from the 2019 fastMRI challenge

Effective acceleration: 9.68x – SSIM: 0.8254



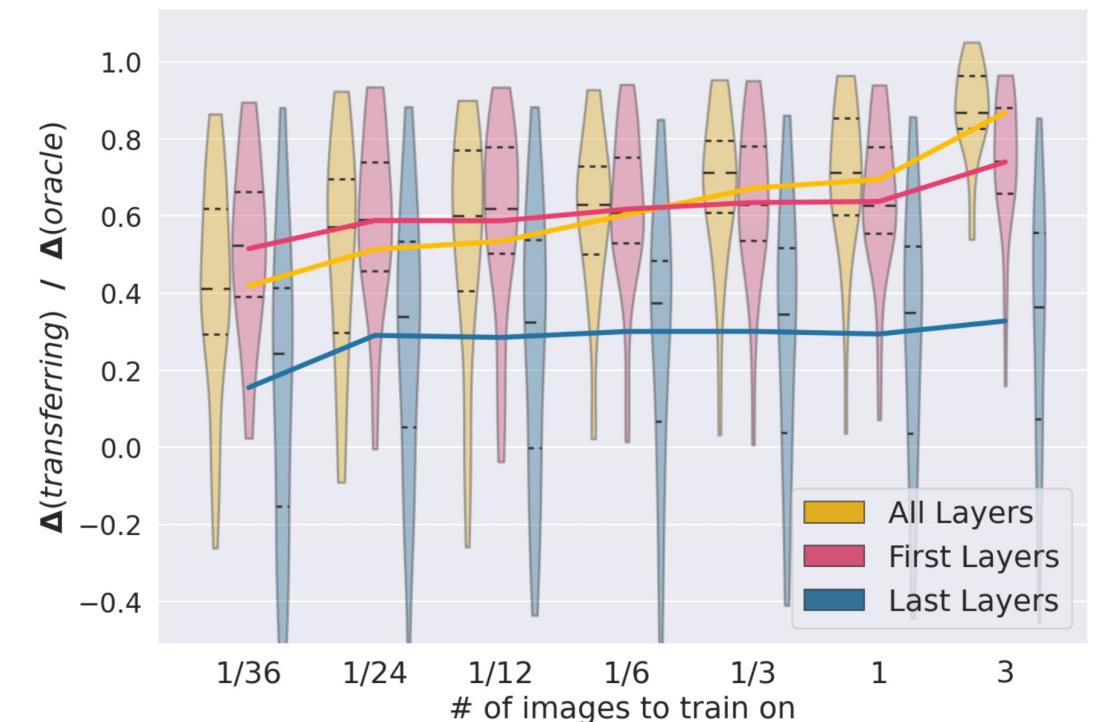
Zero-filled

Recon

Fully sampled

Ongoing work - vendor/domain adaptation

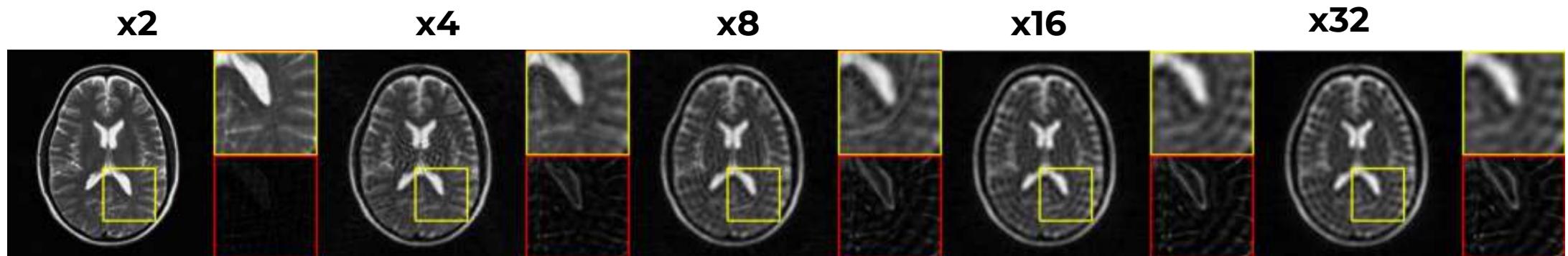
- Major vendors of MRI systems (Philips, GE, Siemens) have **proprietary implementations**, which causes **domain shift** between their data
- Domain adaptation between vendor data remains **ongoing work** with **high commercial interest** for all involved parties (patients, vendors, clinics, third parties)
- FastMRI challenge 2020 had a special track for cross-vendor domain adaptation
- The problem reveals new **theoretical insights**.
For instance, recent work from Shirokikh, Zakazov et. al. [6] (2020) shows that **first** U-Net layers contain more domain-specific information than the **last ones**



Dependence of the relative Surface Dice improvement (y-axis) on the target domain data availability

Ongoing work – generative models for scan acceleration

- Recent work from Belov, Stadelmann, Kastrulyin and Dylov [7] (2021) experiments with generative models for reconstruction from **extremely undersampled** k-spaces (acceleration up to x64)
- Experiments show that:
 - GANs **cannot** produce reliable diagnostic quality (confirm previous discussions)
 - Generated images can be applied for **different MRI applications** such as computer-assisted surgery or radiation planning



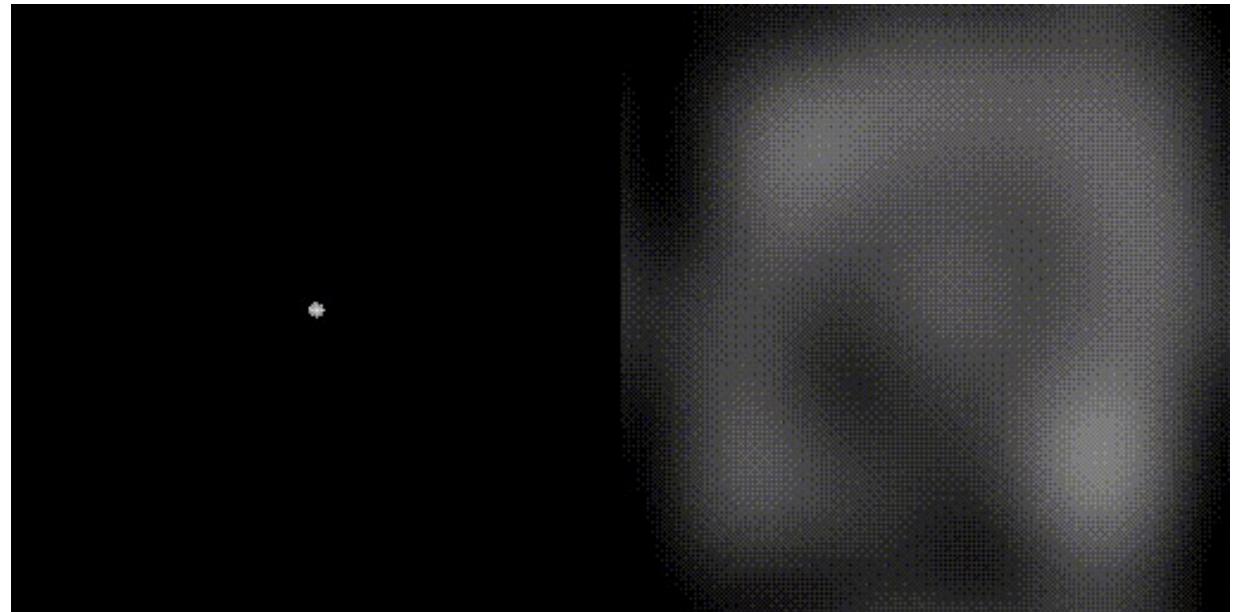
Examples of images obtained using different acceleration factors.

The red crop shows the absolute difference between the ground truth and the undersampled images.

Ongoing work – sampling patterns

- As we saw, **sampling** pattern play a **huge** role in accelerated image formation
- **Cartesian** sampling (the one that we discussed on the seminar) is the **most widespread** technique for sampling of **2D** MR data
- It is possible to find **optimal parameters** of Cartesian sampling w.r.t. **subsequent reconstruction** model
- **Other** sampling techniques exist, and their verification and application is a **hot research topic**

Outward radial ordering of k-space



Courtesy of Brian Hargreaves



Find better sampling pattern

- As we discussed, sampling heavily influences the quality of resulting images
- In Demo 1 we achieved SSIM scores close to 0.61 for $R = 4$ and 0.53 for $R = 8$ (in the single-coil track)
- Find sampling pattern that would beat the baseline SSIM score
 - Target $R = 4$: SSIM > 0.7
 - Target $R = 8$: SSIM > 0.6

<Solution>

- **Code....**

© Credits

- **Slide |7, 34|** <http://mri-q.com/k-space-trajectories.html> animations are courtesy of Brian Hargreaves



Resources

1. Olaf Ronneberger et. al, U-Net: Convolutional Networks for Biomedical Image Segmentation, 2015, [arXiv:1505.04597](https://arxiv.org/abs/1505.04597)
2. Ian Goodfellow et. al., Generative Adversarial Networks, 2014, [arXiv:1406.2661](https://arxiv.org/abs/1406.2661)
3. Jian Zhang and Bernard Ghanem, ISTA-Net: Interpretable Optimization-Inspired Deep Network for Image Compressive Sensing, 2017, [arXiv:1706.07929](https://arxiv.org/abs/1706.07929)
4. Pezzotti, ..., **Kastruylin** et. al, An Adaptive Intelligence Algorithm for Undersampled Knee MRI Reconstruction, 2021, [doi: 10.1109/ACCESS.2020.3034287](https://doi.org/10.1109/ACCESS.2020.3034287)
5. Jure Zbontar et. al., FastMRI: An Open Dataset and Benchmarks for Accelerated MRI, 2018, [arXiv:1811.08839](https://arxiv.org/abs/1811.08839)
6. Boris Shirokikh, Ivan Zakazov et. al., First U-Net Layers Contain More Domain Specific Information Than The Last Ones, 2020, [arXiv:2008.07357](https://arxiv.org/abs/2008.07357)
7. Belov, Stadelmann, **Kastruylin, Dylov**, Towards Ultrafast MRI via Extreme k-Space Undersampling and Superresolution, 2021, [arXiv:2103.02940](https://arxiv.org/abs/2103.02940)