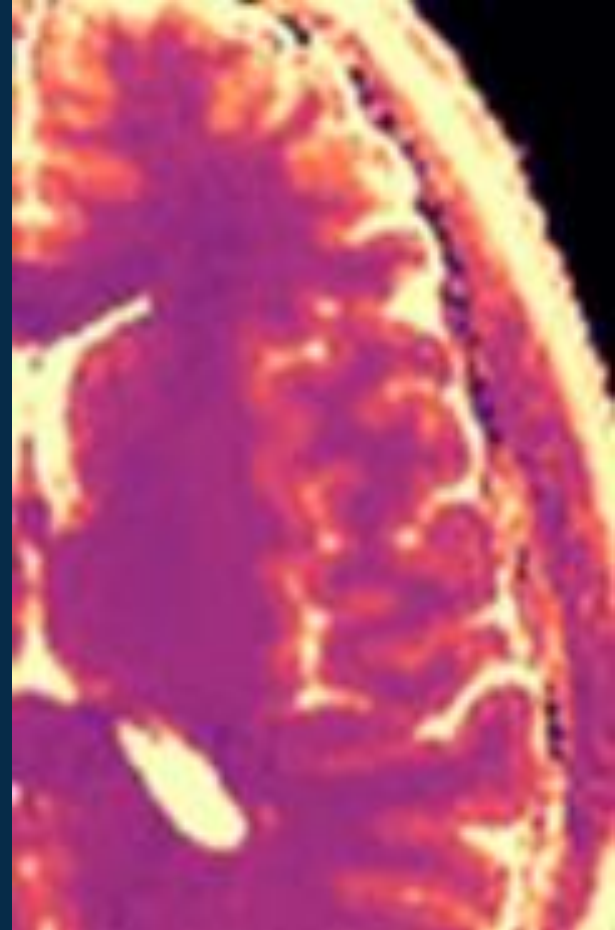


Deep inverse problems – a novel approach to MRI image reconstruction



Matteo Cencini – INFN



IMAGO7

| Overview

1. Background: a short introduction on MRI
2. AI-based MR image reconstruction
3. DeepMR framework

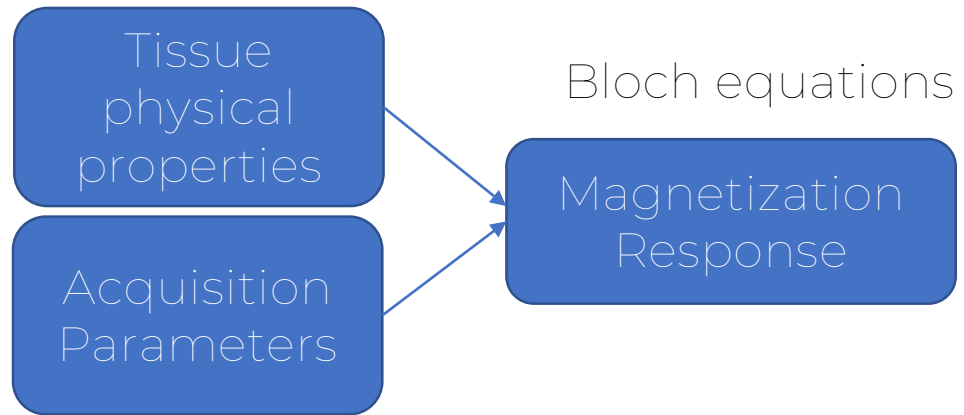
Magnetic Resonance Imaging

| Magnetic Resonance Imaging

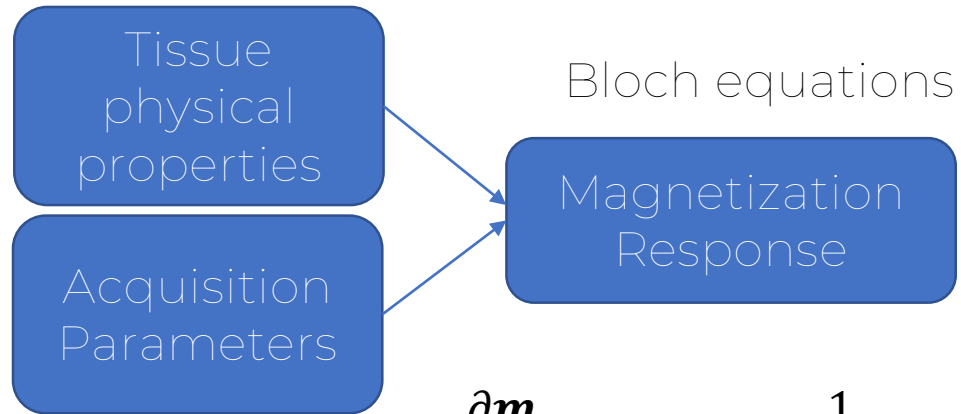
Versatile imaging technique based on the interaction of nuclear spins within the human body with magnetic fields

- ✓ Excellent soft tissue contrast
- ✓ It is non invasive! (non-ionizing radiation)
- Relatively slow due to the characteristics of MR signal encoding

Magnetic Resonance Imaging



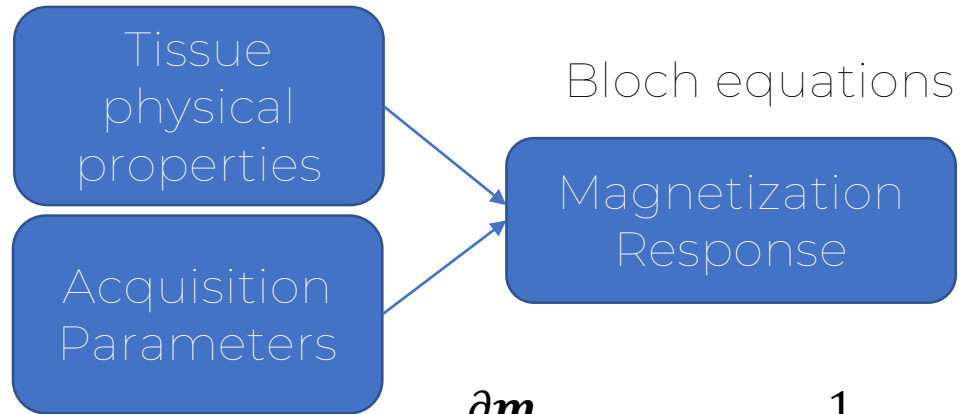
Magnetic Resonance Imaging



$$\frac{\partial \mathbf{m}}{\partial t} = \gamma \mathbf{m} \times \mathbf{B} + \frac{1}{T_1} (m_0 - m_z) \hat{z} + \frac{1}{T_2} \mathbf{m}_\perp$$

Contrast encoding

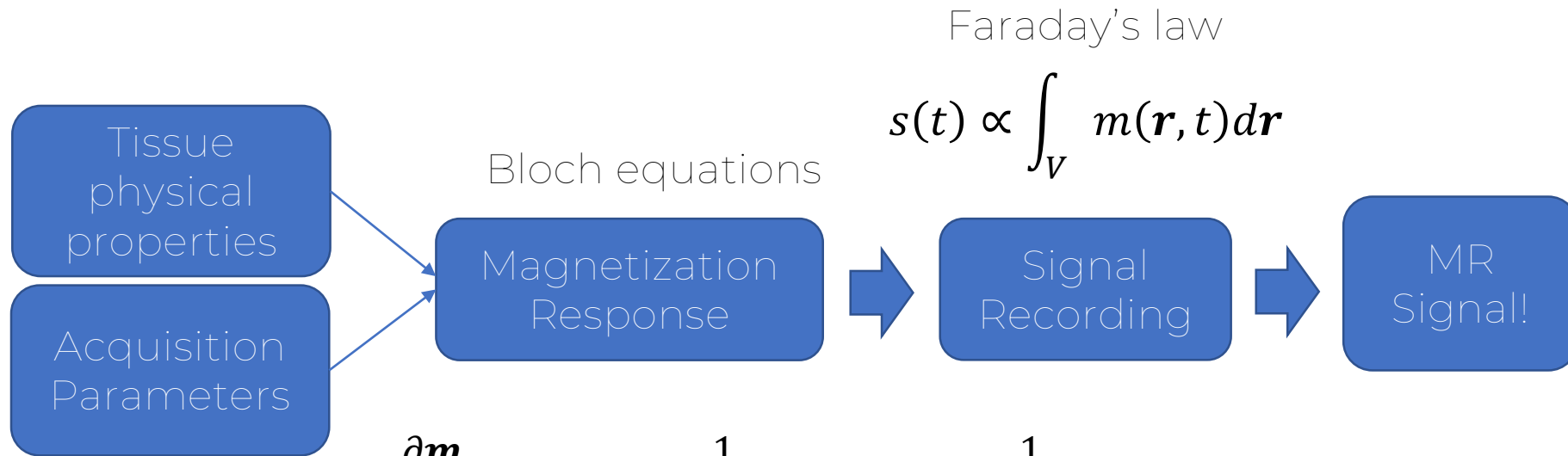
Magnetic Resonance Imaging



$$\frac{\partial \mathbf{m}}{\partial t} = \gamma \mathbf{m} \times \mathbf{B} + \frac{1}{T_1} (m_0 - m_z) \hat{z} + \frac{1}{T_2} \mathbf{m}_\perp \quad \text{Contrast encoding}$$

$$\mathbf{B}(\mathbf{r}) = B_0 \hat{z} + \mathbf{G}(t) \cdot \mathbf{r} \quad \text{Spatial encoding}$$

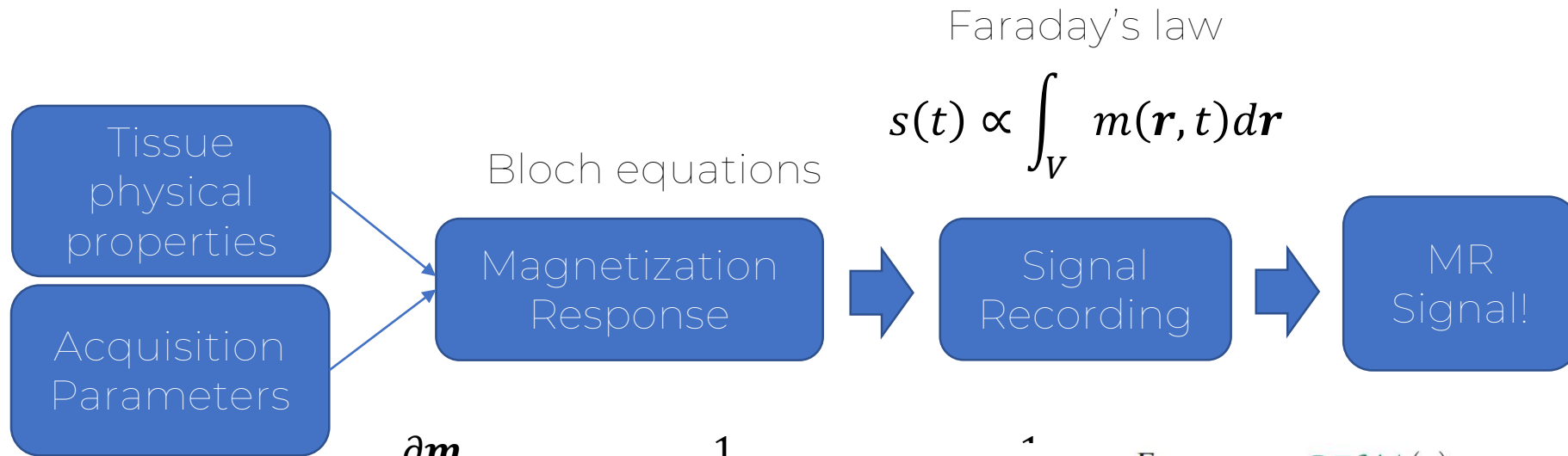
Magnetic Resonance Imaging



$$\frac{\partial \mathbf{m}}{\partial t} = \gamma \mathbf{m} \times \mathbf{B} + \frac{1}{T_1} (m_0 - m_z) \hat{z} + \frac{1}{T_2} \mathbf{m}_\perp \quad \text{Contrast encoding}$$

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Magnetic Resonance Imaging

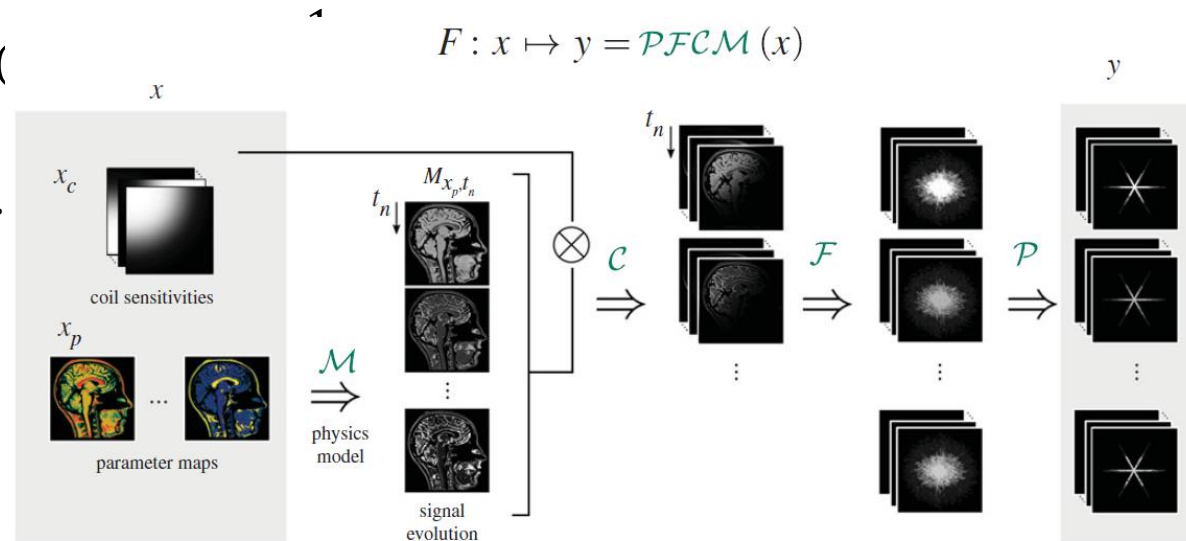


$$\frac{\partial \mathbf{m}}{\partial t} = \gamma \mathbf{m} \times \mathbf{B} + \frac{1}{T_1} (\dots)$$

$$\mathbf{B}(\mathbf{r}) = B_0 \hat{z} + \mathbf{G}(t) \cdot \mathbf{r}$$

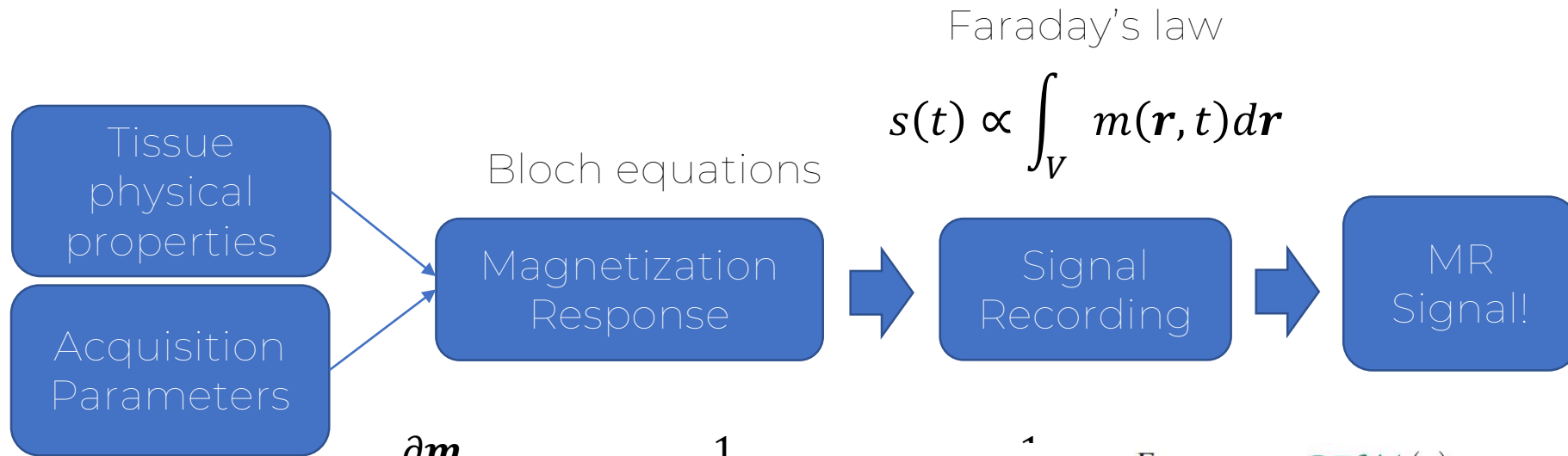
Signal Encoding

$$\mathbf{y} = \mathcal{PFCM}(\mathbf{x}) + n$$



$n = \text{noise}$

Magnetic Resonance Imaging



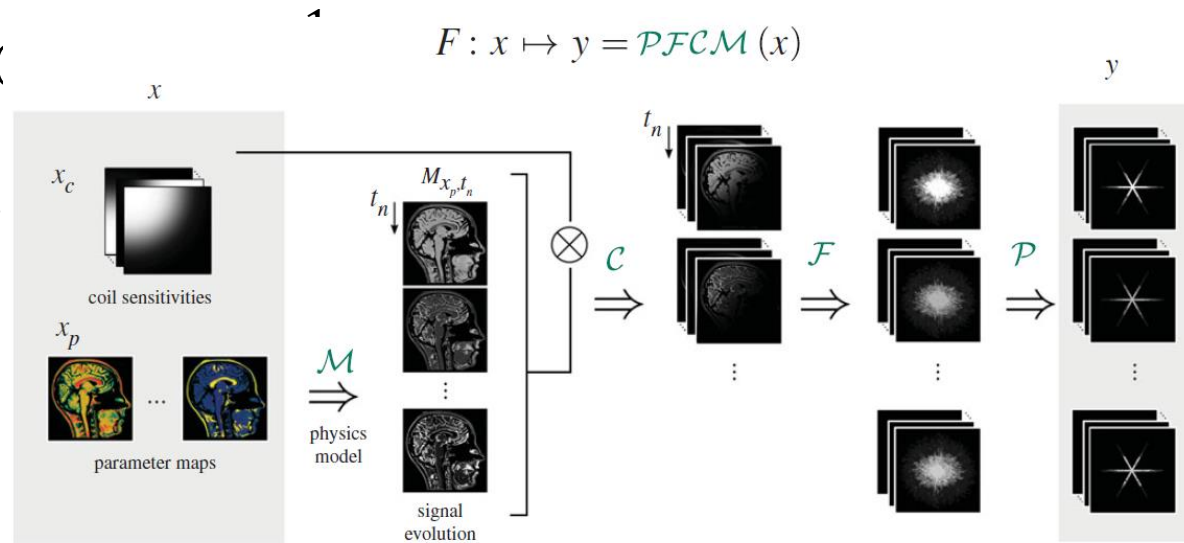
$$\frac{\partial \mathbf{m}}{\partial t} = \gamma \mathbf{m} \times \mathbf{B} + \frac{1}{T_1} (\dots)$$

$$\mathbf{B}(\mathbf{r}) = B_0 \hat{z} + \mathbf{G}(t) \cdot \mathbf{r}$$

Signal Encoding

$$\mathbf{y} = \mathcal{PFCM}(\mathbf{x}) + \mathbf{n}$$

$$\mathbf{y} = \mathcal{PFC}\mathbf{x}' + \mathbf{n}$$



$\mathbf{n} = \text{noise}$

| Problem formulation

Direct inversion

$$\hat{x} = F^H(y)$$

Reconstructed image Acquired data

Encoding operator

| Problem formulation

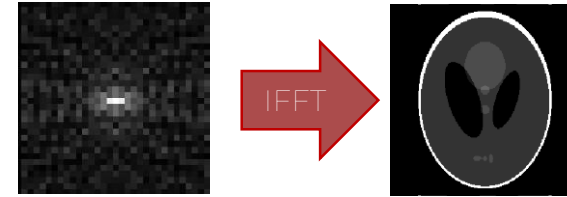
Direct inversion

$$\hat{x} = F^H(y)$$

Reconstructed image

Acquired data

Encoding operator



| Problem formulation

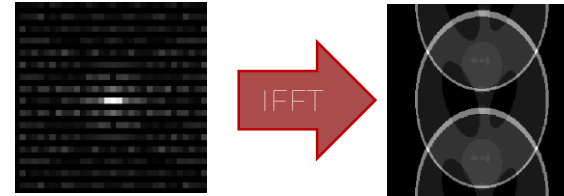
Direct inversion

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Encoding operator



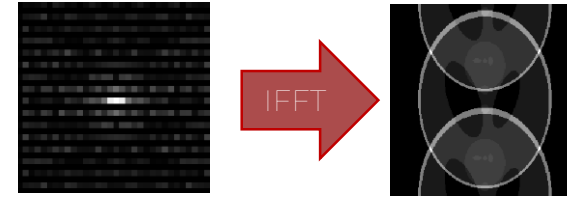
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Direct inversion

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Encoding operator



Imaging Equation

$$\hat{x} = \overbrace{\operatorname{argmin}_x \|F(x) - y\|_2^2}^{\text{Data consistency term}} + \underbrace{\sum_i \lambda_i R_i(x)}_{\text{Regularization term}}$$

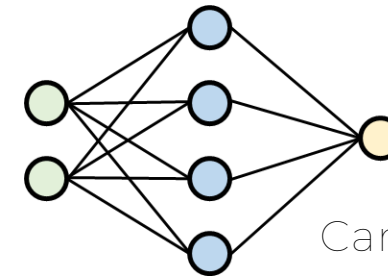
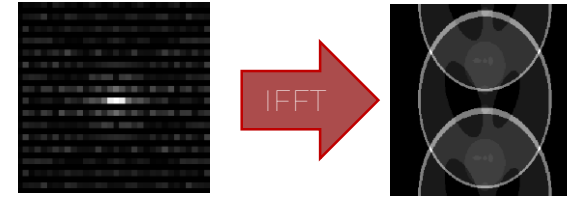
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Reconstructed image Acquired data



Can AI help here?

Imaging Equation

$$\hat{x} = \overbrace{\operatorname{argmin}_x \|F(x) - y\|_2^2}^{\text{Data consistency term}} + \underbrace{\sum_i \lambda_i R_i(x)}_{\text{Regularization term}}$$

AI-based MR reconstruction

| AI-recon classification

Supervised



Both ground-truth and undersampled data are available

| AI-recon classification

Supervised



Both ground-truth and undersampled data are available

Self-Supervised



Undersampled data only are available

| AI-recon classification

Supervised



Both ground-truth and undersampled data are available

Self-Supervised



Undersampled data only are available

Unsupervised



Neither are available (single subject)

Supervised DL models

Ground truth image is available → can compute loss both in measurement and image space

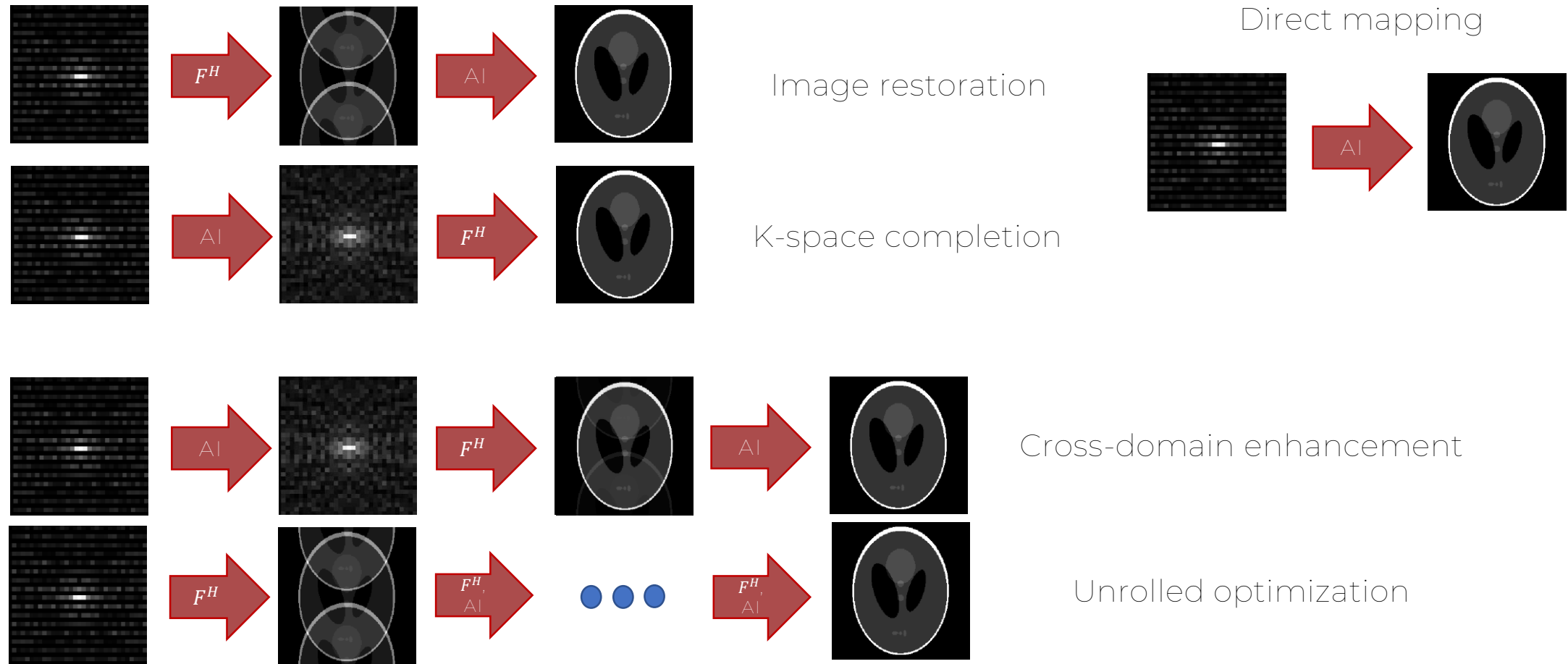
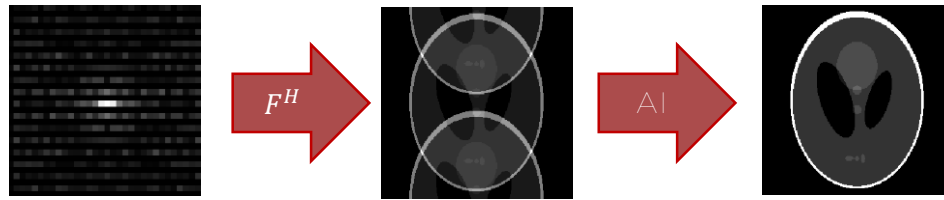
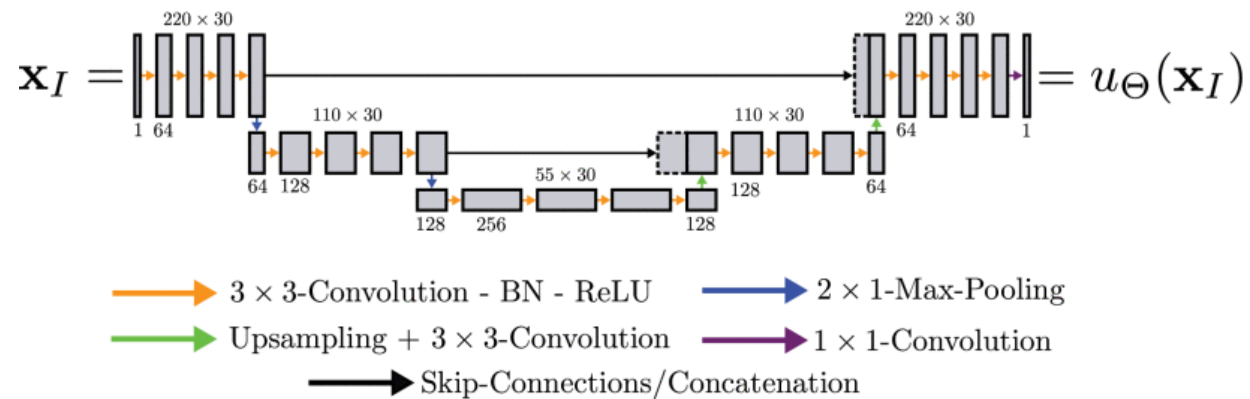


Image restoration

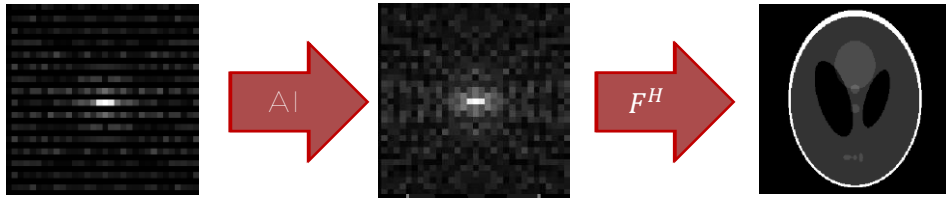


Can rely on rich body of literature on CNN-based denoising (also from non-medical context) – e.g., U-Net



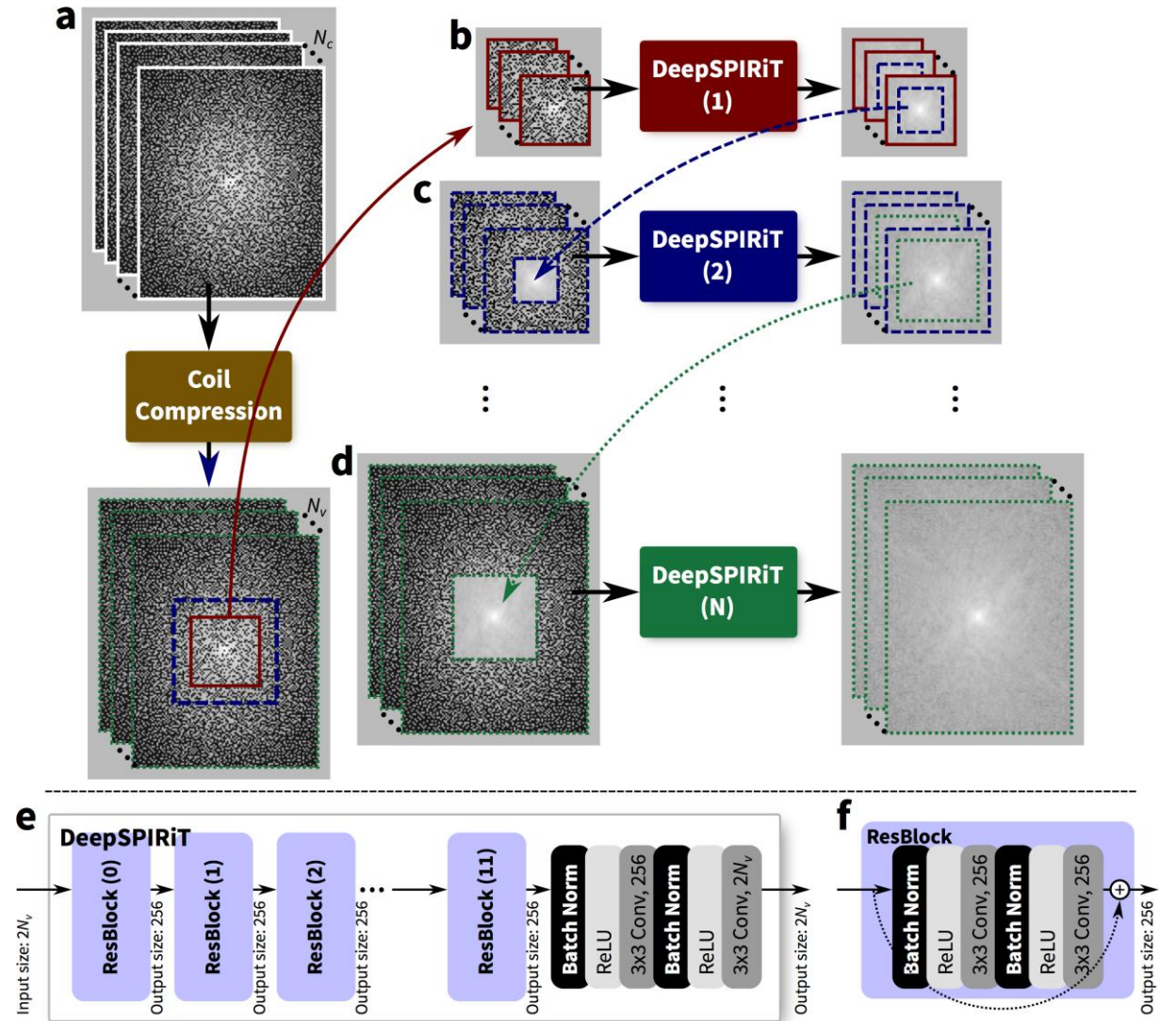
Aliasing should have noise-like pattern \rightarrow does not apply well to common undersampling artifact

K-space completion

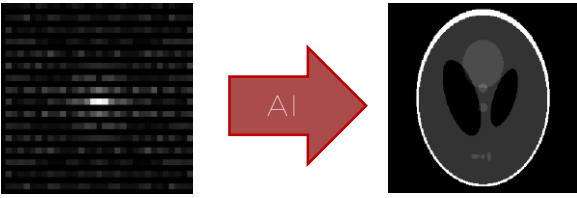


More closely related to the natural representation of MR data (k-space)

More complicated data pre-processing and overall architecture

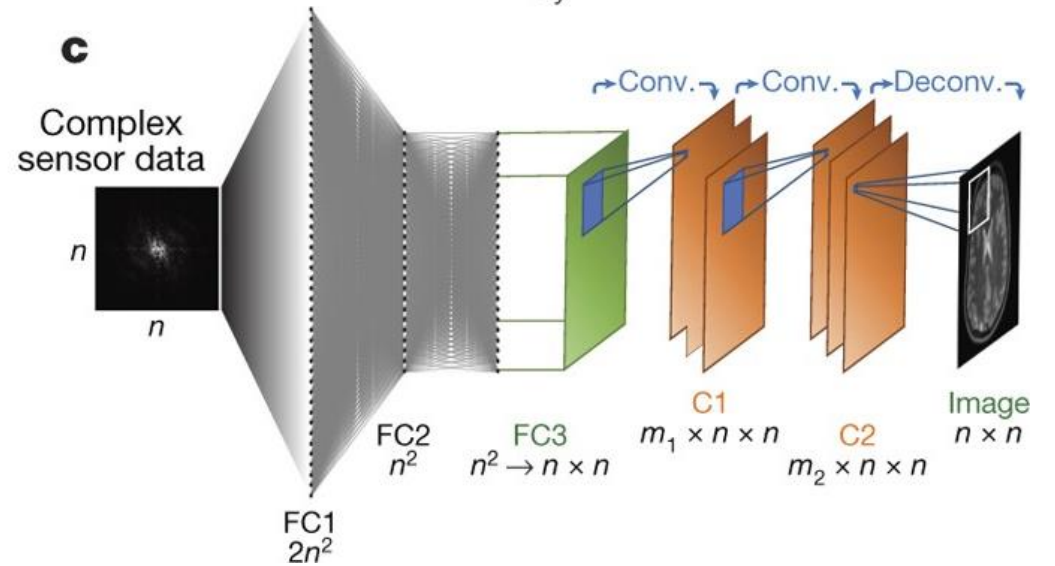
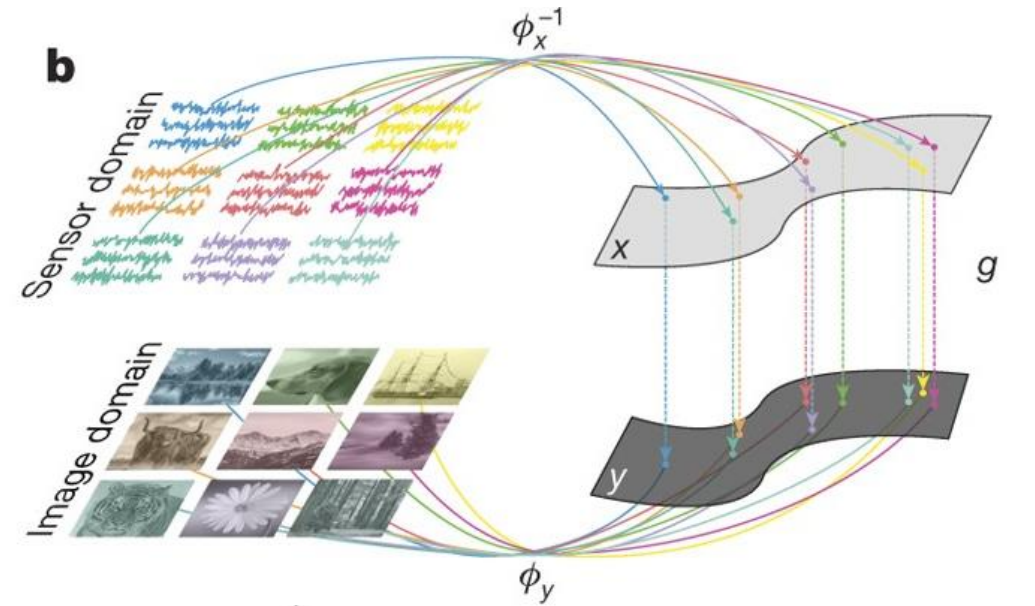


Direct mapping

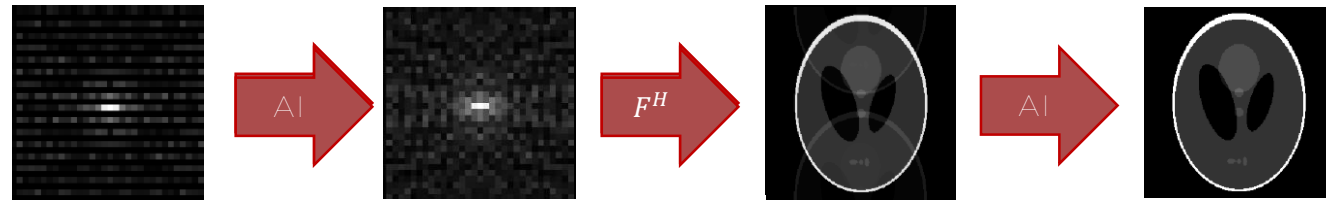


Can capture (and correct) artifacts related to system imperfections

High memory footprint – limited to low dimensional data



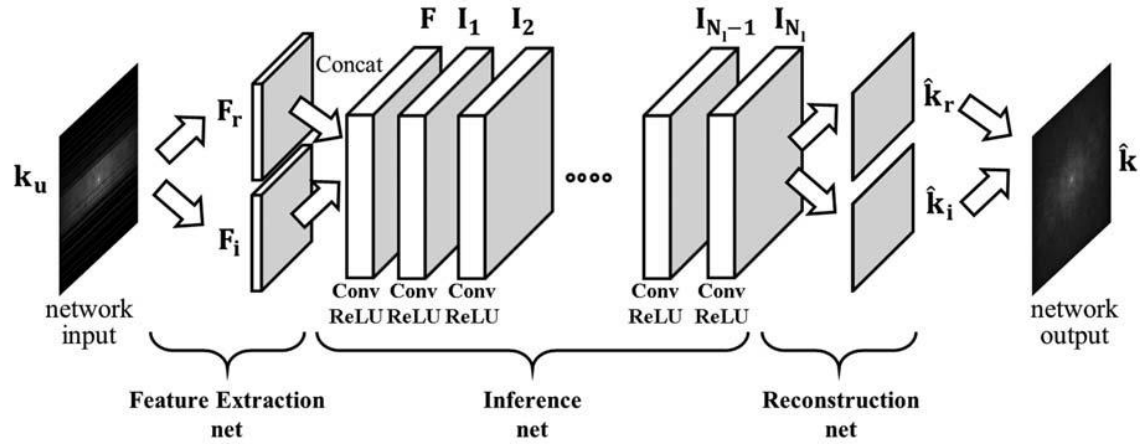
Cross-domain enhancement



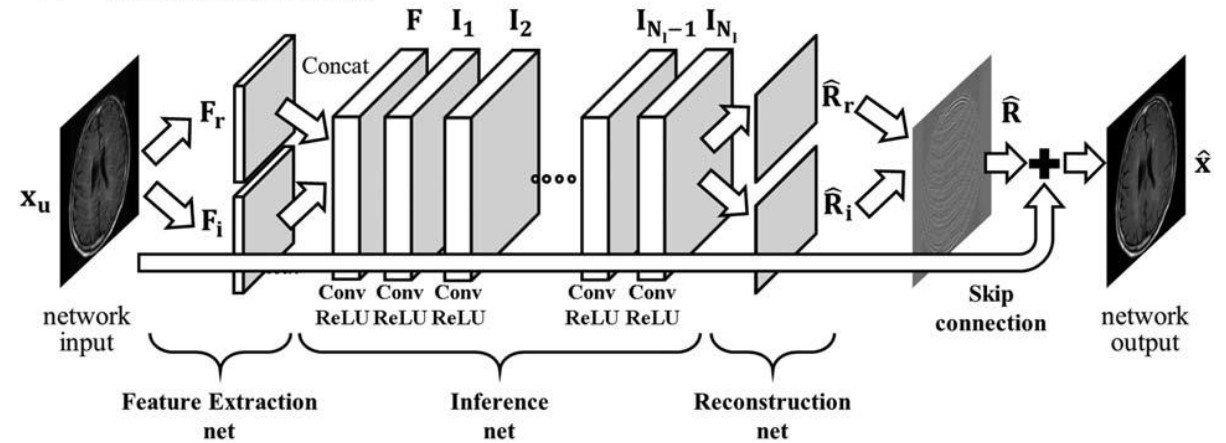
Improved performances by exploiting highly coupled information between the image and k-space domains

Increased complexity

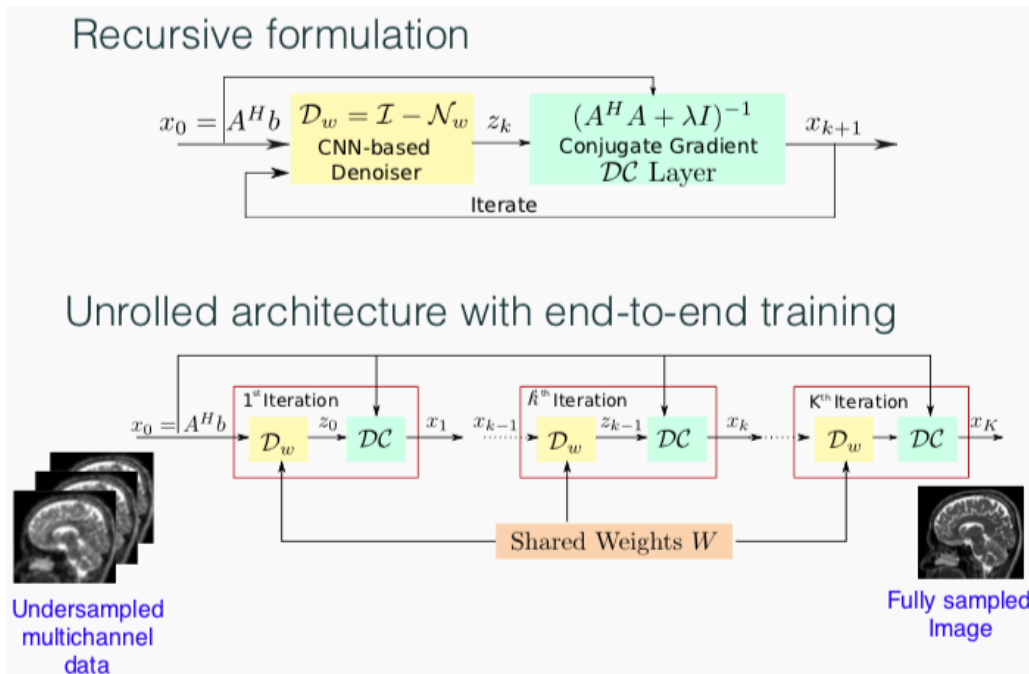
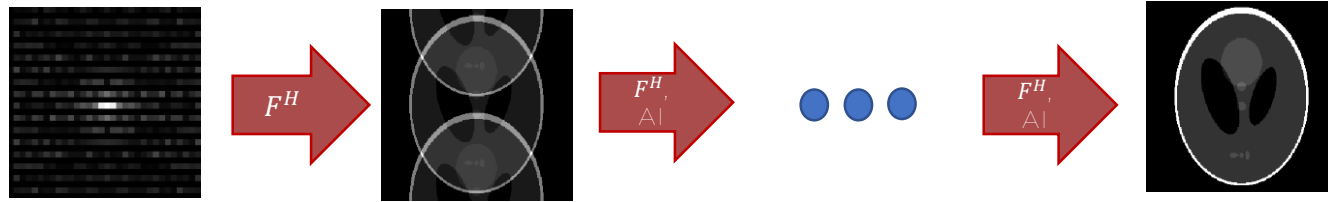
A Architecture of KCNN



B Architecture of ICNN



Unrolled optimization



Require less data for training

Involve potentially inefficient conventional image encoding operators

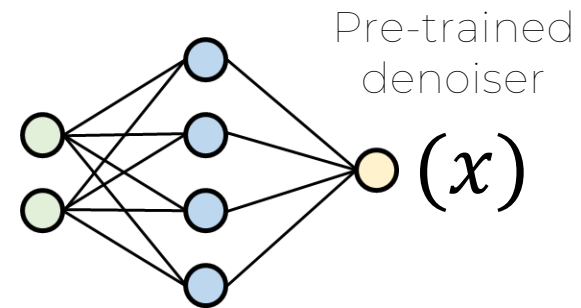
| Self-/un-supervised DL models

Ground truth image not available → can compute loss both in measurement space only

In the limit case of $N_{train} = 1$, training is performed during reconstruction itself

- Example: PnP reconstruction

$$\hat{x} = \operatorname{argmin}_x \|F(x) - y\|_2^2 +$$



DeepMR framework

PREDATOR project

Aim 1:

- Development of an open-source framework for development of advanced qMR reconstruction
- Acquisition of high-quality data for Deep Learning based qMR reconstruction



Aim 2:

- Application to real-world MR acquisition (healthy + patients), either retrospectively on previously acquired data (MR Fingerprinting) and prospectively (QSM)

Aim 3:

- Implement on-line reconstruction on MR scanner
- Measure scalability of the algorithms on low-end hardware (consumer-level PCs, single-board computers) for low-income countries



DeepMR framework

I/O

- Raw data loading
- Dicom Export

Linear Operators

- FFT
- NUFFT
- Wavelet Transform
- ...

Regularization Operators

- L1 Wavelet
- Total Variation
- Local Low Rank
- PnP denoisers
- ...

Solvers

- Gradient Method
- Conjugate Gradient
- ADMM
- ...

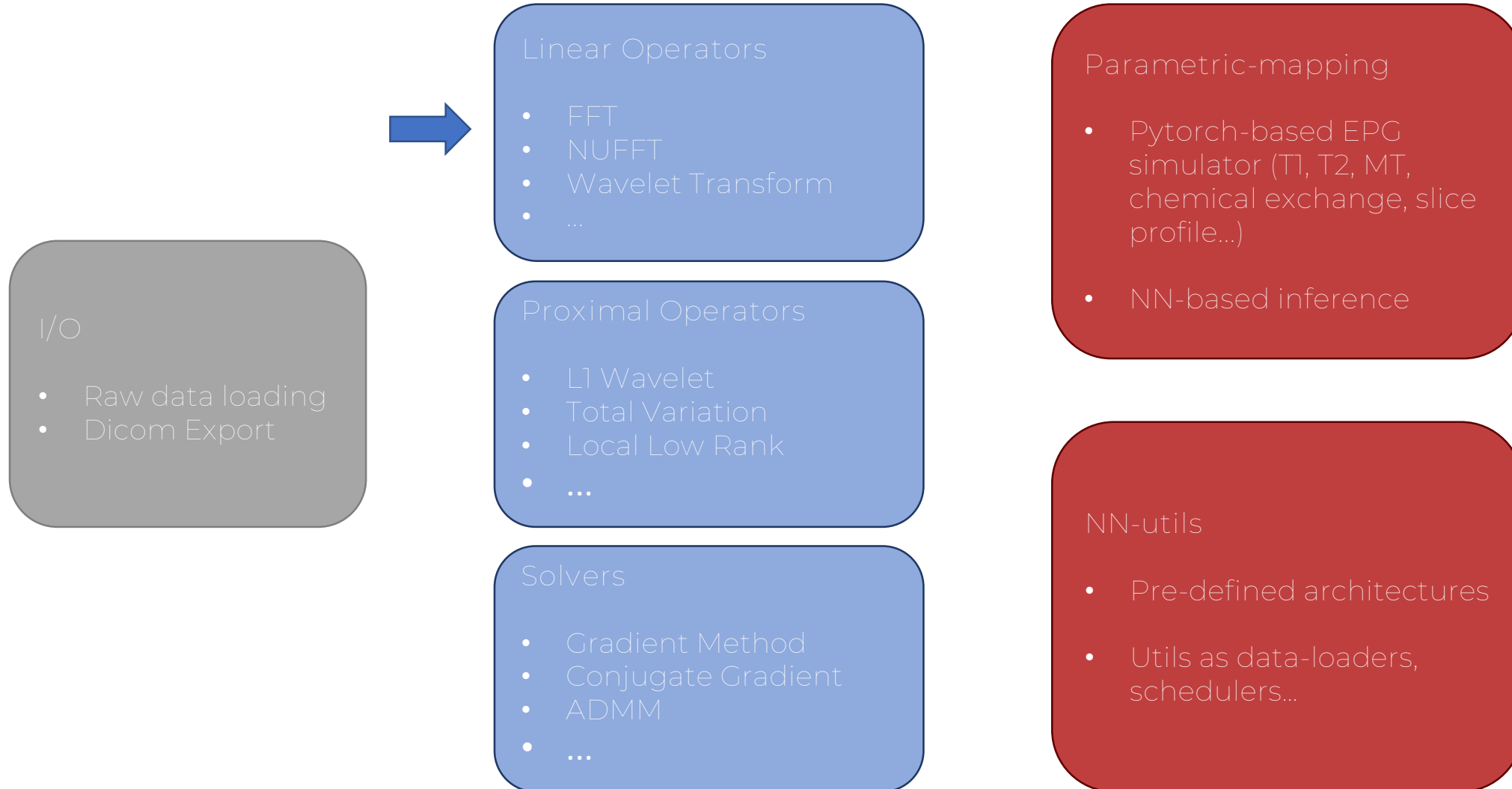
Parametric-mapping

- Pytorch-based EPG simulator (T1, T2, MT, chemical exchange, slice profile...)
- NN-based inference

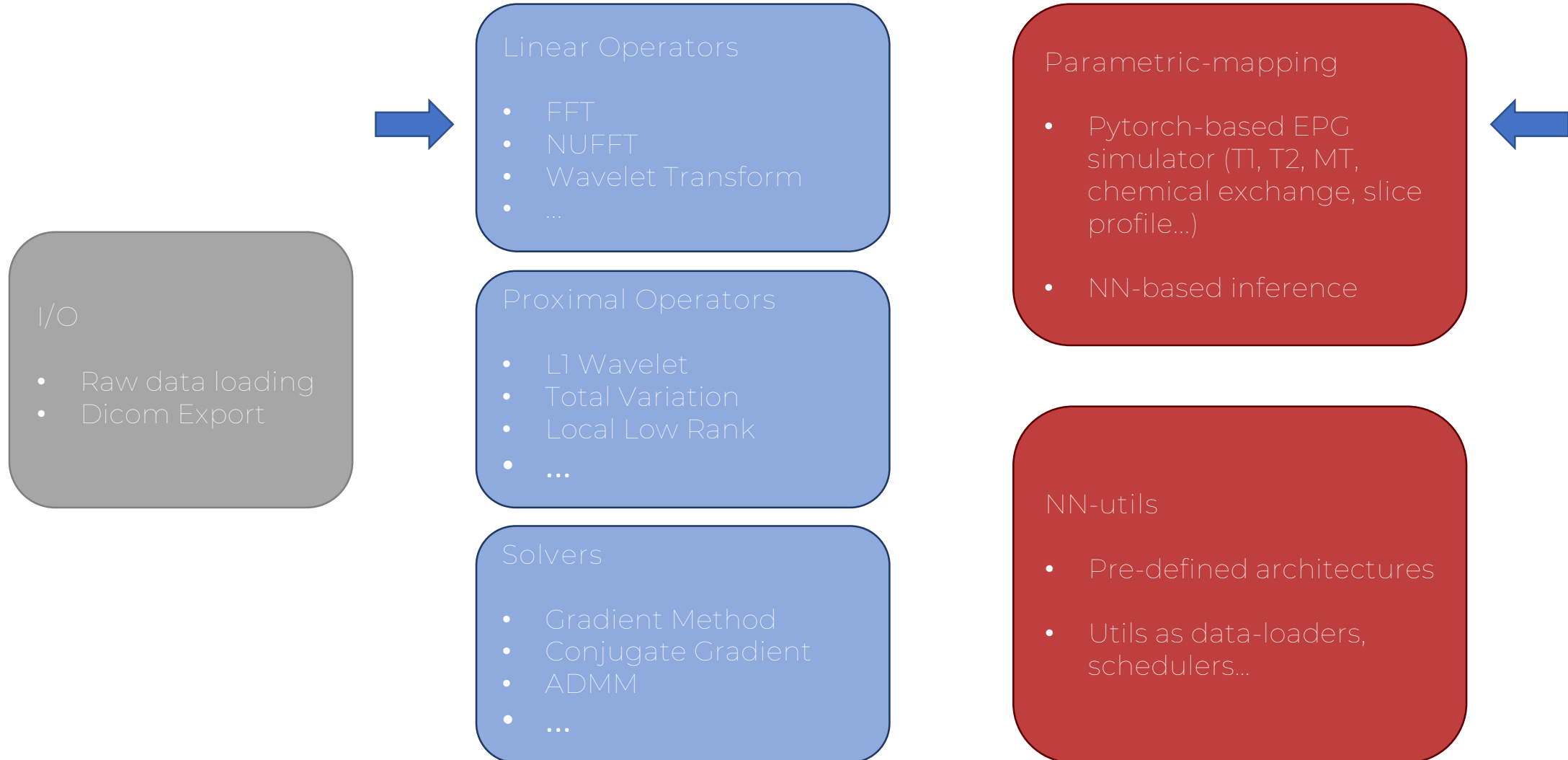
NN-utils

- Pre-defined architectures
- Utils as data-loaders, schedulers...

DeepMR framework



DeepMR framework



Efficient Bloch simulator

| Efficient Bloch simulator

Pytorch implementation based on Extended Phase Graphs formalism

- Effect of RF pulse and spin relaxation is represented by matrix-matrix product operations
- Effect of gradient is represented in the Fourier space as a shift between different (discrete) states representing different dephasing orders

| Efficient Bloch simulator

Pytorch implementation based on Extended Phase Graphs formalism

- Effect of RF pulse and spin relaxation is represented by matrix-matrix product operations
- Effect of gradient is represented in the fourier space as a shift between different (discrete) states representing different dephasing orders

Features:

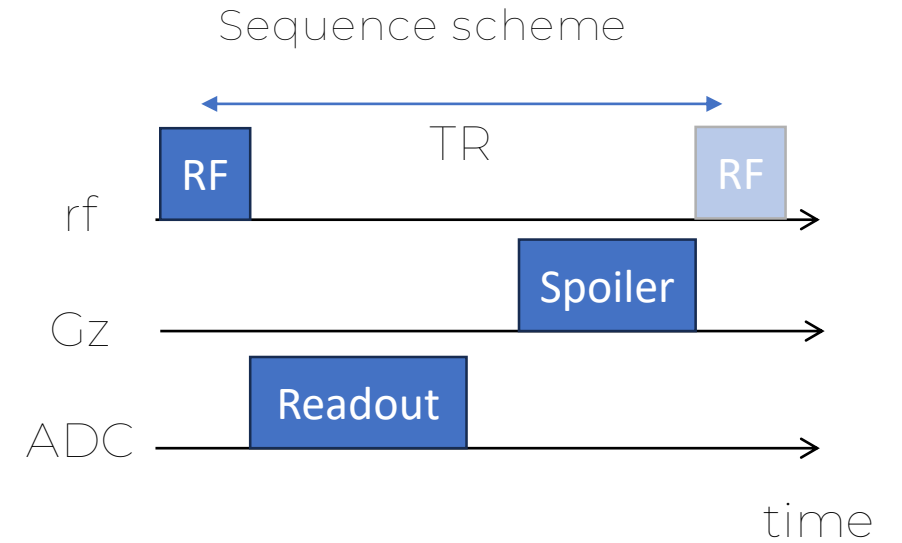
- Include all main MR related parameters (relaxation, diffusion, chemical exchange, magnetization transfer, system imperfections)
- Highly parallelized (both on CPU and GPU)
- Supports gradient calculation via forward automatic differentiation (enable backpropagation in NN training, efficient parameter fitting and optimization of acquisition parameters)

Efficient Bloch simulator

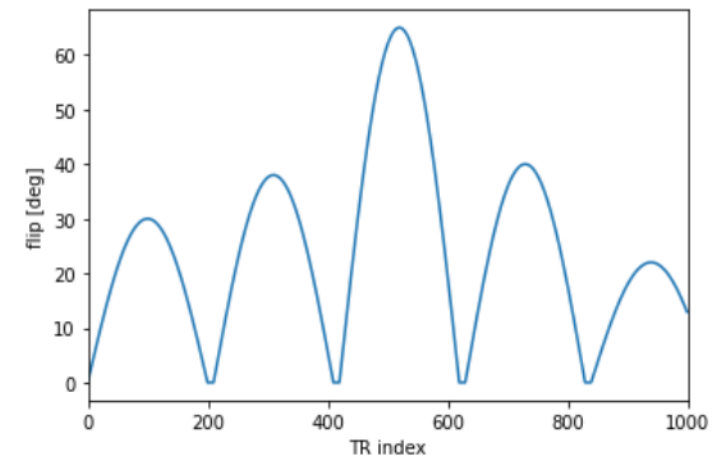
Validation:

- Acquisition: Variable Flip Angle unbalanced SSFP with 1000 echoes (TR=8.5ms)
- Virtual Object: Three representative tissues:
 - White Matter (T1=500ms; T2=70ms)
 - Gray Matter (T1=833ms; T2=83ms)
 - CSF (T1=2569ms; T2=329ms)

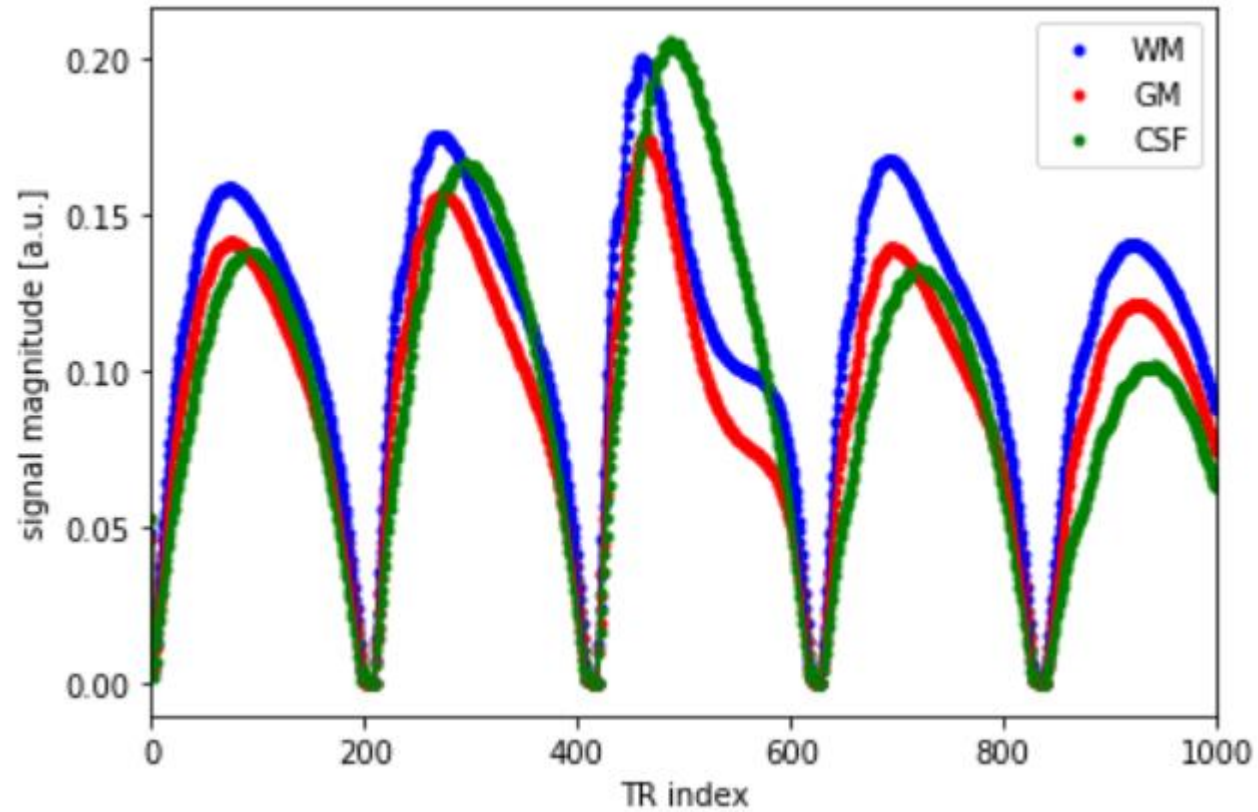
Benchmark: comparison of computations time against *sycomore* (state-of-the-art C++ implementation) on a batch of tissue species of size 65536



Flip angle variation



Efficient Bloch simulator

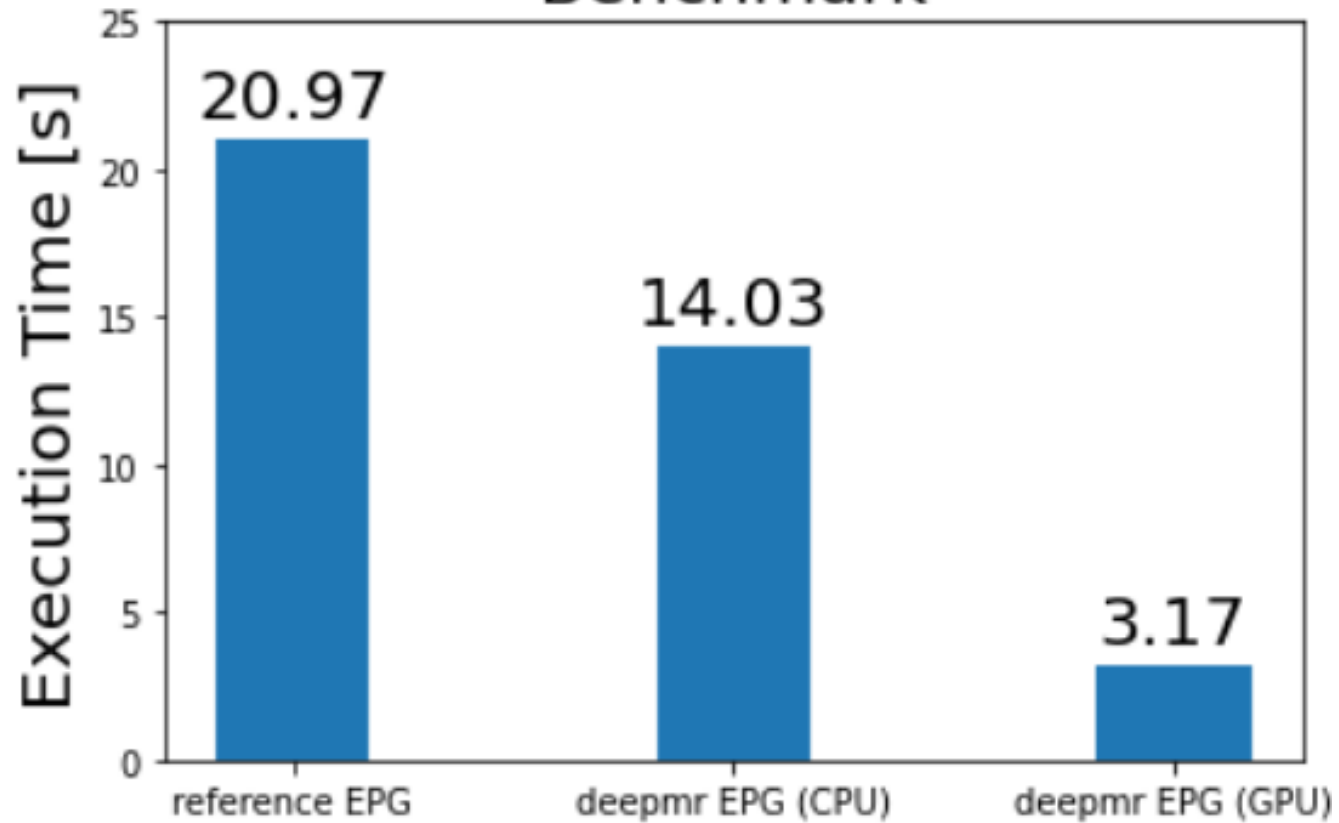


No differences compared to the reference implementation

- : reference implementation
- : deepmr implementation

Efficient Bloch simulator

Benchmark



Up to one-order of magnitude of speed-up
(on GPU)

Efficient encoding operator

| Efficient encoding operator

$$y = PFM(x) + n$$

P: sampling operator

F: Fourier Transform

M: Bloch response

PF: Non-Uniform Sparse FFT

| Efficient encoding operator

$$y = \mathbf{PFM}(x) + n$$

P: sampling operator

F: Fourier Transform

M: Bloch response

PF: Non-Uniform Sparse FFT

| Efficient encoding operator

$$y = PFM(x) + n$$

P: sampling operator

F: Fourier Transform

M: Bloch response

PF: Non-Uniform Sparse FFT

M is often replaced with a projection on a linear subspace for speedup

→ jointly perform P, F and M!

| Efficient encoding operator

$$y = PFM(x) + n$$

M is often replaced with a projection on a linear subspace for speedup

→ jointly perform P, F and M!

Implementation: Pytorch + Numba

- Can operate both on CPU and GPU
- Efficient backpropagation noticing that, for linear operators, backward pass is equivalent to adjoint operator

Efficient encoding operator

Validation:

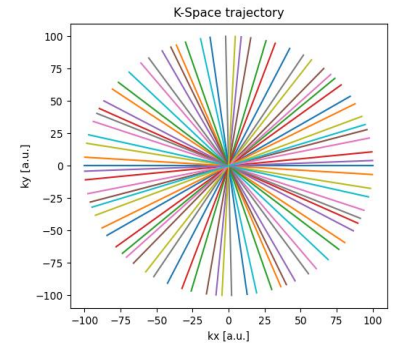
- Acquisition: Fast Spin-Echo with 1000 echoes and golden-angle incremented 2D radial sampling (100 spokes / echo)
- Virtual Object: Shepp-Logan phantom with three regions:
 - White Matter (T2=70ms)
 - Gray Matter (T2=83ms)
 - CSF (T2=329ms)
- Basis determination: SVD of an ensemble of 300 signal evolution (T2:1 – 329ms)
- K-Space data: calculated via forward NUFFT from *torchkbnufft*

Benchmark: comparison of both forward and adjoint NUFFT computations time against *torchkbnufft* implementation

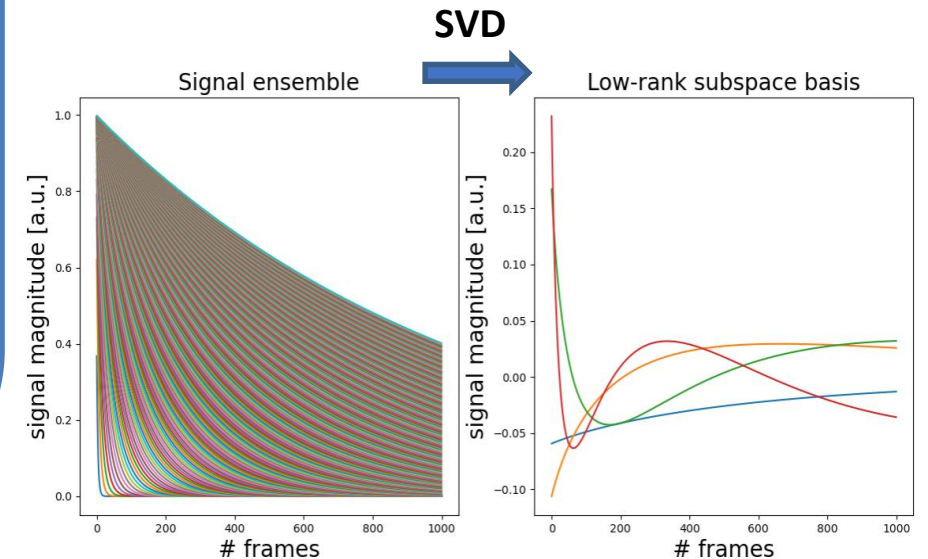
Signal equation

$$S(TE) = \exp\left(-\frac{TE}{T2}\right)$$

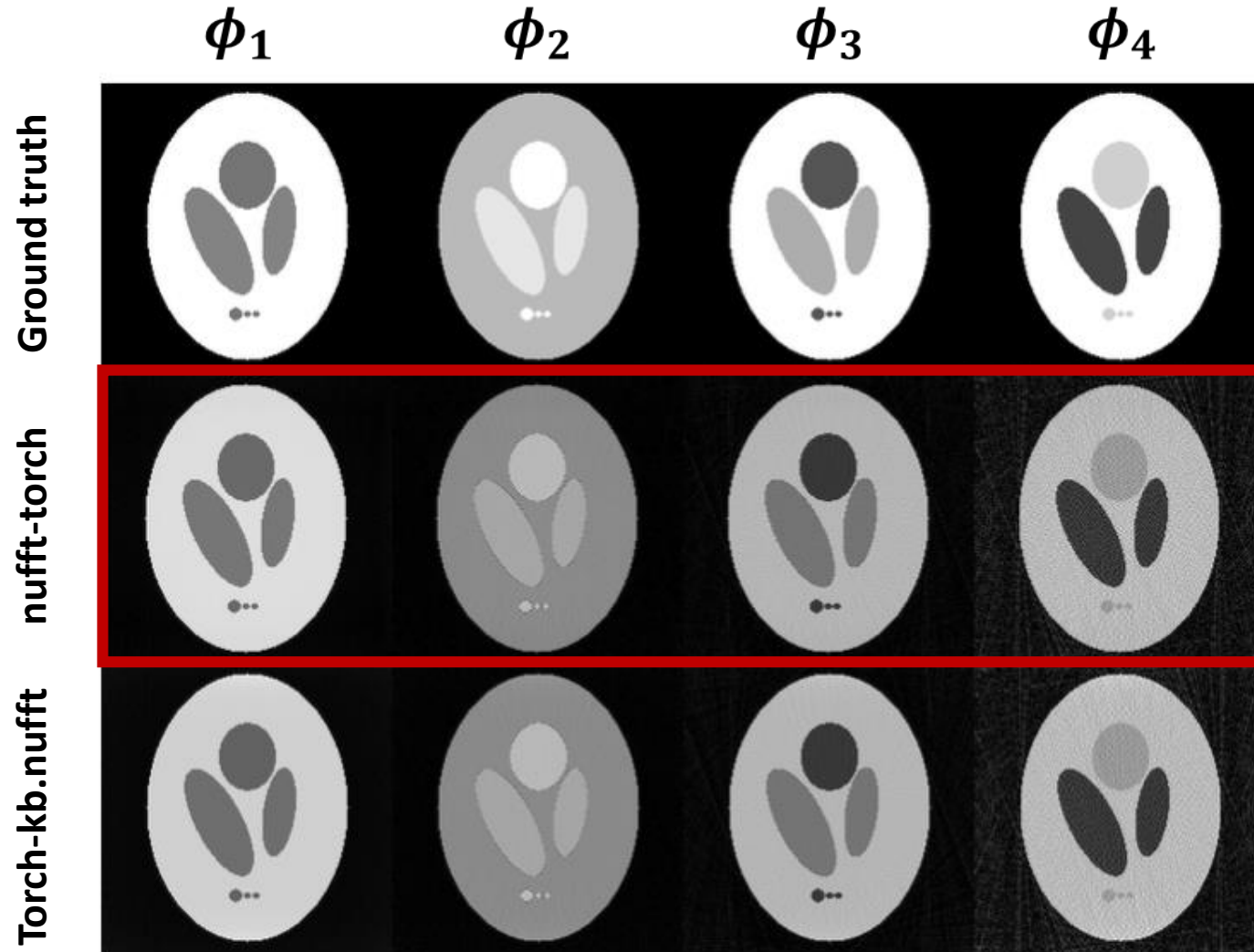
Trajectory



Basis determination

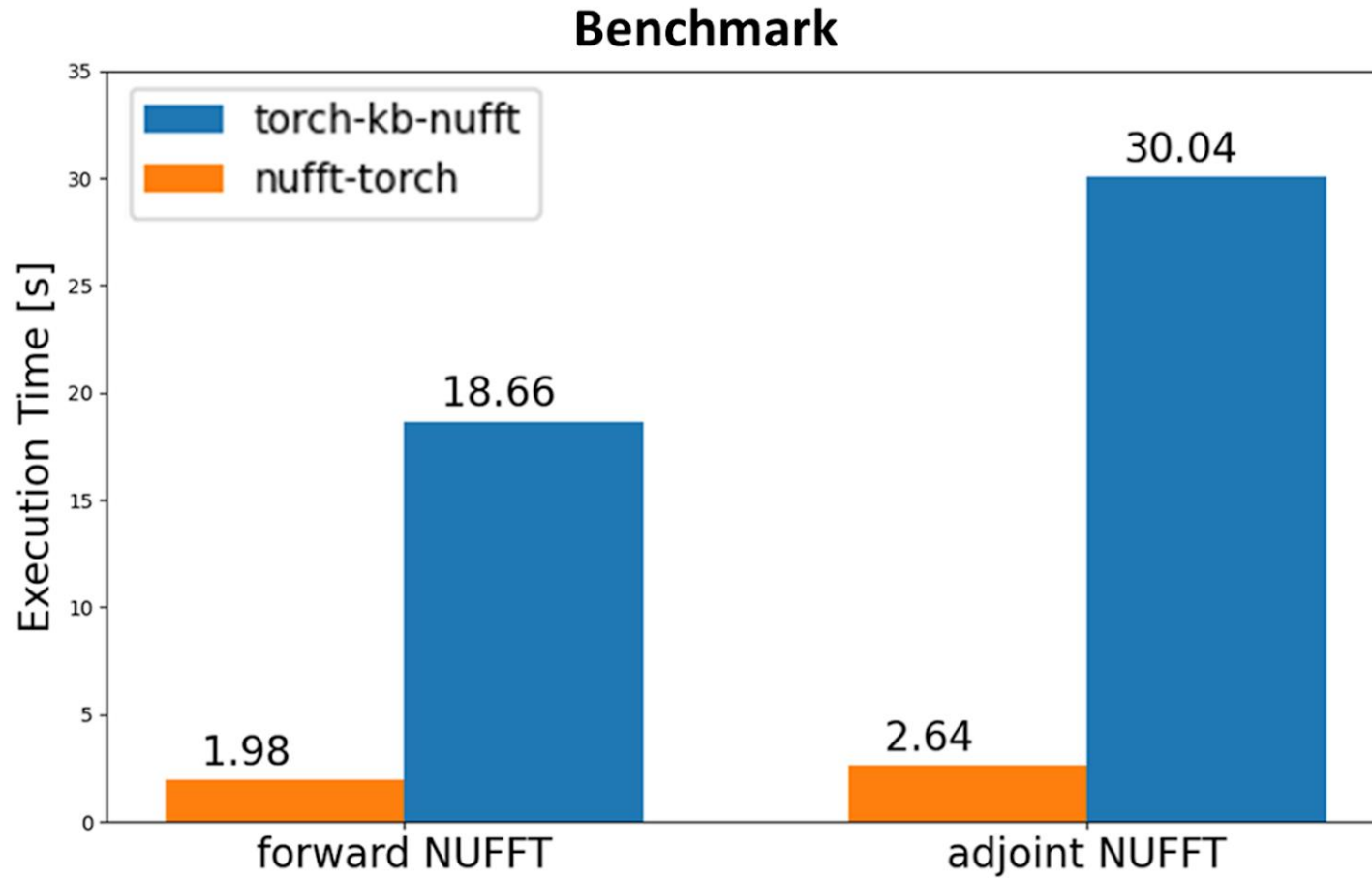


Efficient encoding operator



No differences compared to the reference implementation

Efficient encoding operator



One order of magnitude of speed-up compared to the reference implementation

qMR database creation

qMR database creation

M2. Creation of training qMRI database of healthy volunteers (31 Dec 2023): Generation of sub-millimetric T1, T2, T2*/susceptibility maps of at least 10 subjects.

Protocol

Sequence Type	Parameter	FA [deg]	RF phase inc [deg]	TE [ms]	TR [ms]	FoV [mm]	Resolution [mm]	Acceleration factor	Scan Time [min:sec]
SPGR	T1	3, 11, 28	quadratic	3.3	8.5	224x224x160	0.8 iso	4	9:06
bSSFP	T2	10, 20, 30, 35	0, 180	2	4	224x224x160	0.8 iso	4	6:00
ME-GRE	T2*, B0, QSM	15	quadratic	3.9, 8.5, 13.1, 17.7	30	224x224x160	0.8 iso	4	5:06

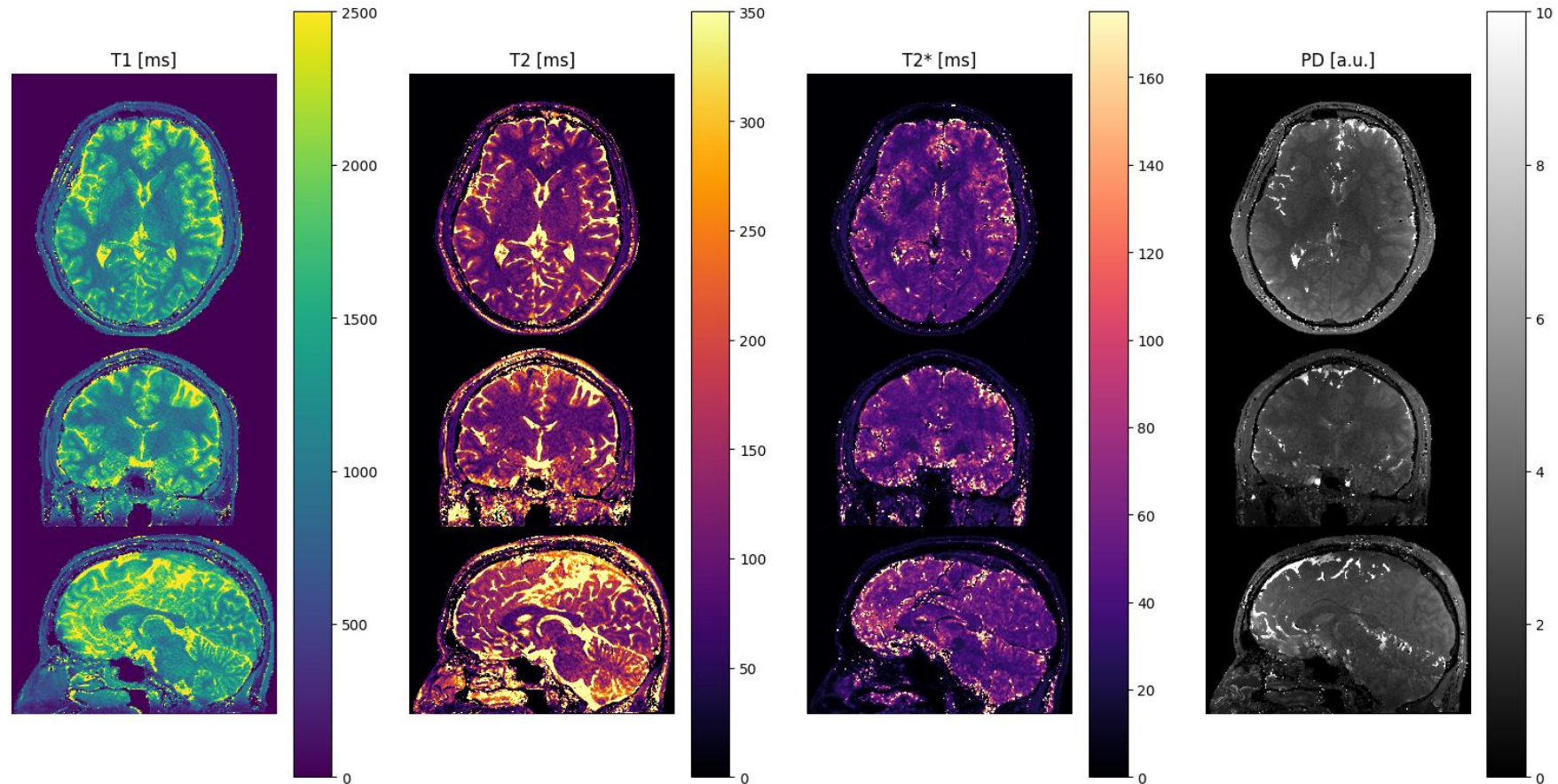
Protocol also include a B1+ calibration scan (Bloch-Siegert) (whole-brain, 4x4x5 mm³ resolution)

Dataset

- 1 healthy volunteer (Male, 34 years old)

qMR database creation

M2. Creation of training qMRI database of healthy volunteers (31 Dec 2023): Generation of sub-millimetric T1, T2, T2*/susceptibility maps of at least 10 subjects.



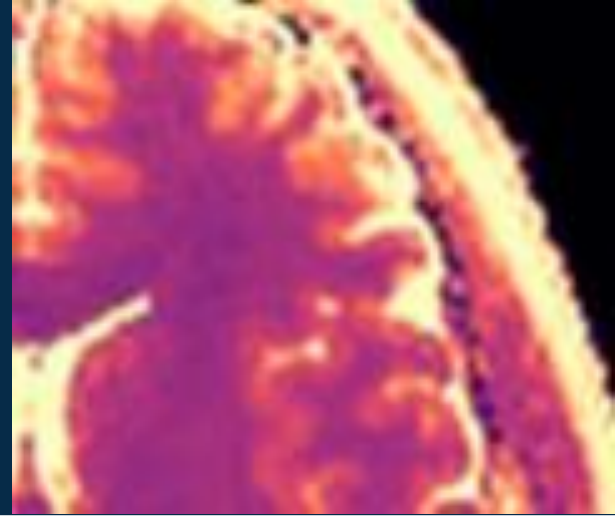
Good geometrical quality and reasonable quantification.

To-Do: improve accuracy and precision of fitting

| Conclusions

- AI-based approaches have great potential in accelerating MR acquisition and reconstruction
- The DeepMR framework will contribute to the field providing easy-to-use, extensible and efficient tools to develop novel reconstruction approaches

Pisa
October 16 2023



Thank you for your attention!



IRCCS FONDAZIONE
STELLA MARIS

IMAGO7