

Relaxometry Guided Quantitative Cardiac Magnetic Resonance Image Reconstruction

Zhao, Yidong; Zhang, Yi; Tao, Qian

DOI

[10.1007/978-3-031-52448-6_33](https://doi.org/10.1007/978-3-031-52448-6_33)

Publication date

2024

Document Version

Final published version

Published in

Statistical Atlases and Computational Models of the Heart. Regular and CMRxRecon Challenge Papers - 14th International Workshop, STACOM 2023, Held in Conjunction with MICCAI 2023, Revised Selected Papers

Citation (APA)

Zhao, Y., Zhang, Y., & Tao, Q. (2024). Relaxometry Guided Quantitative Cardiac Magnetic Resonance Image Reconstruction. In O. Camara, E. Puyol-Antón, A. Suinesiaputra, A. Young, M. Sermesant, Q. Tao, & C. Wang (Eds.), *Statistical Atlases and Computational Models of the Heart. Regular and CMRxRecon Challenge Papers - 14th International Workshop, STACOM 2023, Held in Conjunction with MICCAI 2023, Revised Selected Papers* (pp. 349-358). (Lecture Notes in Computer Science (including subseries Lecture Notes in Artificial Intelligence and Lecture Notes in Bioinformatics); Vol. 14507 LNCS). Springer. https://doi.org/10.1007/978-3-031-52448-6_33

Important note

To cite this publication, please use the final published version (if applicable).
Please check the document version above.

Copyright

Other than for strictly personal use, it is not permitted to download, forward or distribute the text or part of it, without the consent of the author(s) and/or copyright holder(s), unless the work is under an open content license such as Creative Commons.

Takedown policy

Please contact us and provide details if you believe this document breaches copyrights.
We will remove access to the work immediately and investigate your claim.

Green Open Access added to TU Delft Institutional Repository

'You share, we take care!' - Taverne project

<https://www.openaccess.nl/en/you-share-we-take-care>

Otherwise as indicated in the copyright section: the publisher is the copyright holder of this work and the author uses the Dutch legislation to make this work public.



Relaxometry Guided Quantitative Cardiac Magnetic Resonance Image Reconstruction

Yidong Zhao^(✉), Yi Zhang, and Qian Tao

Department of Imaging Physics, Delft University of Technology,
Delft, The Netherlands

{y.zhao-8,y.zhang-43,q.tao}@tudelft.nl

Abstract. Deep learning-based methods have achieved prestigious performance for magnetic resonance imaging (MRI) reconstruction, enabling fast imaging for many clinical applications. Previous methods employ convolutional networks to learn the image prior as the regularization term. In quantitative MRI, the physical model of nuclear magnetic resonance relaxometry is known, providing additional prior knowledge for image reconstruction. However, traditional reconstruction networks are limited to learning the spatial domain prior knowledge, ignoring the relaxometry prior. Therefore, we propose a relaxometry-guided quantitative MRI reconstruction framework to learn the spatial prior from data and the relaxometry prior from MRI physics. Additionally, we also evaluated the performance of two popular reconstruction backbones, namely, recurrent variational networks (RVN) and variational networks (VN) with U-Net. Experiments demonstrate that the proposed method achieves highly promising results in quantitative MRI reconstruction.

Keywords: Cardiac MRI · Quantitative mapping · Relaxometry · Image reconstruction

1 Introduction

Quantitative Magnetic Resonance Image (qMRI) has emerged as an indispensable imaging modality in research and clinical applications thanks to its quantitative measurements of tissue properties, such as T_1 and T_2 relaxation times [23]. A common approach for qMRI typically involves a two-step process. Firstly, multiple k-space data of a subject are acquired and reconstructed into a series of weighted images with varying imaging parameters (such as echo times and diffusion weights). Subsequently, the underlying tissue properties are estimated by fitting a signal model to the images [24]. However, acquiring fully sampled k-space data, adhering to the Nyquist criterion, for multiple measurements in qMRI requires a time-consuming endeavor [6]. This extended acquisition process introduces the potential for motion artifacts and leads to patient discomfort due to prolonged scan duration.

Motivated by reducing the acquisition time in qMRI while keeping the reconstruction and estimation quality, *accelarated qMRI* has become one of the central research topics in qMRI. The methods in accelerated qMRI can be divided into two categories according to whether the method includes an intermediate step of image reconstruction from k-space data: If so, the method is categorized as *indirect reconstruction*; Otherwise, the method is categorized as *direct reconstruction* where only the parameter mapping estimations are gained [24]. This paper will focus on indirect reconstruction methods since the goals require both reconstructing images and estimating the mapping parameters. However, our proposed method is flexible yet novel, taking self-supervised quantitative mapping as auxiliary constraints to guide the reconstruction.

In conventional qMRI, the acceleration can be achieved by parallel imaging (PI) [9, 12] or compressed sensing (CS) [2]. With specific undersampling patterns, PI utilizes multiple radio-frequency (RF) coils simultaneously with individual coil sensitivity maps to reduce the aliasing artifacts. The applications of PI include SENSE [20], which is the original implementation, and GRAPPA [5], which allows auto-calibration when sensitivity maps are missing. Meanwhile, CS-based methods achieve acceleration by using partial sampling in k-space, which loosens the requirement of the Nyquist criterion with sparsity assumption in true signals. Though the CS-based methods are well-established, they still struggle with the design of regularizer functions, which varies across different tissues [17].

Accelerated qMRI reconstruction benefits from recent rapid development in deep learning on both computer vision and inference learning since MRI reconstruction can be formulated as an inverse problem: Given a (partial) measurement in k-space, the target is to recover the image signal as close as possible. Deep learning-based methods for MRI reconstruction can be categorized into *iterative* methods [8, 16, 25, 29, 30] and *one-step* methods [11, 31]. One-step methods aim to predict the refined reconstructed images from images reconstructed from corrupted k-space [11] or partially sampled k-space measurements directly [31]. Iterative methods leverage neural networks to learn an incremental refining process, analogous to a conventional optimization process. As a first attempt, Yang et al. [29] proposed ADMM-Net: a neural network to parameterize the alternating direction method of multipliers (ADMM), thus solving the inverse problem of MRI reconstruction in combination with CS. Similarly, Hammernik et al. proposed a variational network [8] in an unrolled gradient descent scheme for generalized CS reconstruction. Inspired by a more general idea in meta-learning, recurrent inference machines (RIM) [21] were designed to find a maximum-a-posteriori (MAP) estimation to solve inverse problems when the corresponding forward model is known. This is then applied in MRI reconstruction [16], where in the RIM framework, the parameters are shared across the iterations with internal hidden states instead of individual units for each iteration in previous works. A more recent work, RecurrentVarNet [30], combines variational networks and RIM with a hybrid domain-learning strategy using convolutional neural networks (CNN) to guide the optimization in k-space.

Unlike naive MRI reconstruction, in qMRI reconstruction, the involvement of physical signal models in methodological design can further improve the quality of both reconstruction and parameter mapping estimation since an anatomical correspondence across images is often assumed when designing the acquisition pipeline [10]. Apart from conventional least-square methods, parameter mappings can be estimated by dictionary matching [7, 19, 24]. Given the reconstructed images or raw k-space measurements, there are several methods utilizing self-supervised learning to estimate parameter mappings [1, 14, 15]. However, few existing works using [3] address the joint tasks of reconstruction and parameter estimation, while none lie in the deep learning paradigm.

1.1 Contributions

In this work, we tackle the quantitative mapping problem in the CMRxRecon challenge and make the following contributions:

- We introduce a novel quantitative mapping network that learns to mimic MR physics in an unsupervised fashion.
- We evaluate two different CNN architectures for image prior learning in the variational reconstruction network.
- We leverage the quantitative mapping network to guide the reconstruction process, ensuring that the output signal conforms to the MR relaxometry.

2 Methods

2.1 Parallel Imaging

Accelerated MRI. In this paper, we focus on the reconstruction of k_t qMRI baseline images $\mathbf{x} \in \mathbb{C}^{k_x \times k_y \times k_t \times 1}$ ¹, given their corresponding k-space measurements $\mathbf{y} \in \mathbb{C}^{n_c \times k_x \times k_y \times k_t}$ of n_c coils, where k_x and k_y are image shapes along the frequency and phase encoding axes, respectively. The c^{th} coil measurement $\mathbf{y}_c$ is formulated as

$$\mathbf{y}_c = U\mathcal{F}S_c\mathbf{x} + \epsilon_c, \quad (1)$$

where $\mathcal{F}$ is the 2D Fourier transform operator, S_c denotes the sensitivity map, U characterises the Cartesian under-sampling pattern and ϵ_c represents the measurement noise. The overall forward operator and its adjoint operator in accelerated parallel imaging can be formulated as $\mathcal{A} = U \circ \mathcal{F} \circ \mathcal{E}$ and $\mathcal{A}^* = \mathcal{R} \circ \mathcal{F} \circ U$ [30], where

$$\mathcal{E}(\mathbf{x}) = [S_1\mathbf{x}, S_2\mathbf{x}, \dots, S_{n_c}\mathbf{x}], \quad (2)$$

$$\mathcal{R}(\mathbf{y}) = \sum_{c=1}^{n_c} S_c^* \mathbf{y}_c. \quad (3)$$

¹ We use $\mathbf{x}$ for both complex and magnitude image for simplicity.

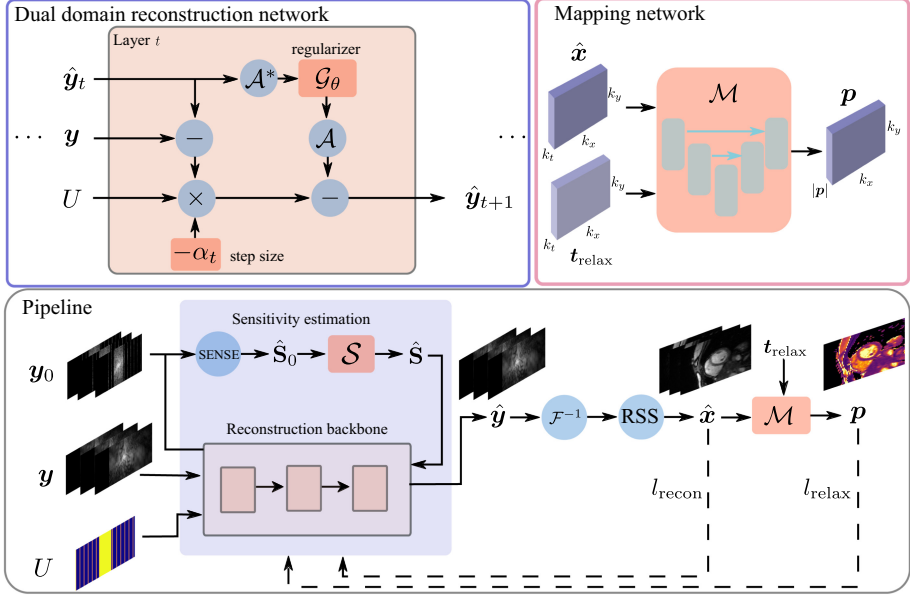


Fig. 1. The reconstruction backbone consists of unrolled gradient descent layers, and the image prior is learned during training by $\mathcal{G}_\theta$. A pre-trained mapping network $\mathcal{M}$ is introduced to predict the quantitative parameters $\mathbf{p}$ and guide the reconstruction with MR relaxometry.

Sensitivity Estimation. The sensitivity maps are estimated using the auto-calibration region U_{AC} in k -space which is always sampled in the low-frequency band. The initial sensitivity maps can be estimated as

$$\hat{S}_c^0 = \frac{\mathcal{F}^{-1} U_{AC} \mathbf{y}_c}{\text{RSS}(\{\mathcal{F}^{-1} U_{AC} \mathbf{y}_l\}_{l=1}^{n_c})}, \quad (4)$$

where RSS denotes the root-sum-of-square operator defined in [13]

$$\text{RSS}(\mathbf{x}_1, \mathbf{x}_2, \dots, \mathbf{x}_{n_c}) = \sqrt{\sum_{c=1}^{n_c} |\mathbf{x}_c|^2}. \quad (5)$$

2.2 Relaxometry Guided Reconstruction

Dual-Domain Reconstruction Network. Reconstruction of an image $\mathbf{x}$ given its under-sampled parallel imaging measurements $\mathbf{y}$ can be formulated as optimizing the Lagrangian $\mathcal{L}(\mathbf{x})$ defined as

$$\arg \min_{\hat{\mathbf{x}}} \mathcal{L}(\mathbf{x}) = \|\mathcal{A}\hat{\mathbf{x}} - \mathbf{y}\|_2^2 + R(\hat{\mathbf{x}}), \quad (6)$$

where the first term constrains the data fidelity via a 2-norm operator and the latter term $R(\cdot)$ is a regularizer to stabilize the solution space. The update rule of gradient-descent for optimization of Eq. 6 is

$$\begin{aligned}\hat{\mathbf{x}}_{t+1} &= \hat{\mathbf{x}}_t - \alpha_t \frac{\partial \mathcal{L}}{\partial \hat{\mathbf{x}}^*} \\ &= \hat{\mathbf{x}}_t - \alpha_t \mathcal{A}^* (\mathcal{A} \hat{\mathbf{x}}_t - \mathbf{y}) - \mathcal{G}_{\theta_t}(\hat{\mathbf{x}}_t),\end{aligned}\quad (7)$$

where $\mathcal{G}_{\theta_t}$ is a convolutional network that learns the scaled Wirtinger derivatives $\frac{\partial R}{\partial \hat{\mathbf{x}}^*}$ of the regularizer *w.r.t.* the current reconstruction, and α_t is a trainable step scalar. Applying Fourier transform on both sides of Eq. 7 yields

$$\hat{\mathbf{y}}_{t+1} = \hat{\mathbf{y}}_t - \alpha_t U(\hat{\mathbf{y}}_t - \mathbf{y}) - \mathcal{F} \circ \mathcal{E} \circ \mathcal{G}_{\theta_t}(\mathcal{R} \circ \mathcal{F}^{-1} \hat{\mathbf{y}}_t), \quad (8)$$

in which the data fidelity part is updated in k-space and the regularization is performed in the spatial domain. In this work, we follow [8, 30] and use the variational network based on the unrolled update rule defined in Eq. 8 for image construction. Specifically, we investigate and compare two different types of CNNs to learn the regularizer $\mathcal{G}_{\theta_t}$: U-Net [22] and convolutional Gated Recurrent Unit (GRU) blocks as in [30]. Additionally, we employ another U-Net $\mathcal{S}$ to refine the initial sensitivity map estimated in Eq. 4 $\hat{S}_c = \mathcal{S}(\hat{S}_c^0)$. The reconstruction network is trained with a combination of L_1 loss and structural similarity index measure (SSIM) [28]:

$$l_{\text{recon}}(\hat{\mathbf{x}}) = \gamma_1 \|\hat{\mathbf{x}} - \mathbf{x}\|_1 + \gamma_2 \text{SSIM}(\hat{\mathbf{x}}, \mathbf{x}), \quad (9)$$

where $\hat{\mathbf{x}}$ and $\mathbf{x}$ denote the predicted image and the fully sampled image, respectively. The architecture of reconstruction backbone layers is shown on the upper left panel in Fig. 1.

Quantitative Mapping Network. Given reconstructed magnitude image $\mathbf{x} \in \mathbb{R}^{k_x \times k_y \times k_t}$, quantitative mapping is usually treated as a parameter fitting problem and solved by least squares or dictionary matching [7, 19, 24]. Voxels in a reconstructed image of good quality should conform to the MR relaxometry and thus have a relatively lower level of fitting error since tissue anatomies are assumed to be spatially aligned. Inspired by this, we propose using the parameter fitting error to guide the reconstruction procedure. However, both least squares and dictionary matching are not differentiable and thus cannot be integrated into the computational graph. To make the mapping procedure differentiable, we propose using a U-Net $\mathcal{M}$ to predict the parameters given the image $\mathbf{x}$ and the inversion time or echo time $\mathbf{t}_{\text{relax}}$. With the MR relaxation physics known, the network $\mathcal{M}$ is trained in a purely unsupervised fashion, and the physics-informed training loss l_{relax} is defined as:

$$l_{\text{relax}}(\mathbf{x}) = \|\mathcal{S}(\mathcal{M}(\mathbf{x}, \mathbf{t}_{\text{relax}}), \mathbf{t}_{\text{relax}}) - \mathbf{x}\|_1, \quad (10)$$

where $s(\cdot)$ can be either s_{T_1} or s_{T_2} which characterises the signal intensity models for T_1 or T_2 relaxation:

$$s_{T_1}(A, B, T_1^*, t_{\text{relax}}) = \left| A - B \exp\left(-\frac{t_{\text{relax}}}{T_1^*}\right) \right|, \quad (11)$$

$$s_{T_2}(A, T_2, t_{\text{relax}}) = A \exp\left(-\frac{t_{\text{relax}}}{T_2}\right). \quad (12)$$

Note that from Eq. 11, the parameter of interest T_1 is derived by $T_1 = (B/A - 1)T_1^*$.

Joint Reconstruction and Quantitative Mapping. The mapping network $\mathcal{M}$ is pre-trained with the fully sampled images only and is frozen during the reconstruction network training process. The physics-informed loss l_{relax} can then be used to enforce a relaxation physics-informed reconstruction. We also penalize the difference between the mapping prediction on fully sampled image $\mathcal{M}(\mathbf{x})$ and on the predicted image $\mathcal{M}(\hat{\mathbf{x}})$, such that the reconstructed image has a consistent parameter map as the fully sampled image. The total training loss is then formulated as:

$$l(\hat{\mathbf{x}}) = l_{\text{recon}}(\hat{\mathbf{x}}) + \gamma_3 l_{\text{relax}}(\hat{\mathbf{x}}) + \gamma_4 \|\mathcal{M}(\hat{\mathbf{x}}) - \mathcal{M}(\mathbf{x})\|_1. \quad (13)$$

An overview of the proposed method is illustrated in Fig. 1.

3 Experiments

3.1 Dataset

We conduct the experiments on the CMRxRecon challenge data² [27], consisting of 120 subjects for training. Imaging was performed on a Siemens 3T MRI scanner (MAGNETOM Vida) and the multi-coil images were compressed to 10 virtual coils, with acceleration factors 4, 8, and 10. More details on the image acquisition protocol are described in [26]. Data of each subject comprise two different qMRI sequences: the modified-look-locker (MOLLI) [18] sequence with 9 baseline images for T_1 -mapping and the T2-prepared (T2prep)-FLASH sequence with 3 baseline images for T_2 -mapping. The validation set contains 59 subjects, and the results are evaluated on the official platform.

3.2 Training Configuration

We first train the mapping network $\mathcal{M}$, a U-Net with 256 base filters and 1 pooling layer, by Adam optimizer with an initial learning rate of $\eta_0 = 10^{-4}$ for 200 epochs. And then we freeze $\mathcal{M}$ for training the reconstruction backbone. We

² <https://cmrxrecon.github.io/>.

studied two different architectures for the reconstruction: the recurrent variational network (RVN) [30] using GRU units for regularizer and the variational network [8] with U-Net as the regularizer (VN-UNet). The number of unrolled layers was set as 10 for both configurations. The loss function weighting was set as $\gamma_1 = 0.2, \gamma_2 = 0.8$. For VN-UNet, we also study the performance with relaxometry guidance, setting $\gamma_3 = 0.01, \gamma_4 = 0.1$ (VN-UNet-relax). During training, the network is trained by the Adam optimizer with an initial learning rate $\eta_0 = 10^{-3}$ for 400 epochs.

Data augmentation was performed on each individual coil image as in [4], with random rotation $[-45^\circ, 45^\circ]$, translation $[-10\%, 10\%]$, shearing $[-20^\circ, 20^\circ]$ and vertical/horizontal flip. Additionally, we contaminate the k-space by additive Gaussian noise with a random signal-to-noise-ratio (SNR) in $[6.67, +\infty]$. The augmentation was performed with a probability of 40%.

4 Results

4.1 Evaluation Results

We evaluate the three aforementioned configurations: RVN, VN-UNet, and VN-UNet-relax on the validation set. For simplicity, we only list the results on T_1 mapping with acceleration factor $R = 4$ in Table 1. The best performance was achieved by VM-UNet-relax with a PSNR of 42.59 dB. The evaluation results on all acceleration factors of both T_1 and T_2 mapping are shown in Table 2.

Table 1. Ablation of model architecture settings on the T_1 mapping validation set ($R = 4$). The relaxometry guided variational network with U-Nets achieved the best performance.

Model Settings	PSNR (dB) $\uparrow$	NMSE ($\times 10^{-3}$) $\downarrow$	SSIM (%) $\uparrow$
RVN	41.55	3.4	97.53
VN-UNet	42.50	2.6	97.91
VN-UNet-relax	42.59	2.6	97.93

4.2 Qualitative Results

We show a few exemplar reconstructed images and their corresponding quantitative maps in Fig. 2. The baseline images with the shortest and longest inversion or echo times are listed.

5 Discussion and Conclusion

Table 1 shows that the VN-UNet achieved better performance than RVN, and the performance is slightly improved by introducing relaxometry guidance. However, only image-based metrics like SSIM are provided during the validation

Table 2. Evaluation results of VN-UNet-relax on the validation set. A higher acceleration factor causes a performance drop.

Dataset	PSNR (dB)	NMSE ($\times 10^{-3}$)	SSIM (%)
T_1 - $R = 4$	43.1	2	98.1
T_1 - $R = 8$	37.8	7	95.6
T_1 - $R = 10$	36.6	9	95.1
T_2 - $R = 4$	38.9	3	97.1
T_2 - $R = 8$	35.2	6	94.9
T_2 - $R = 10$	34.4	7	94.6

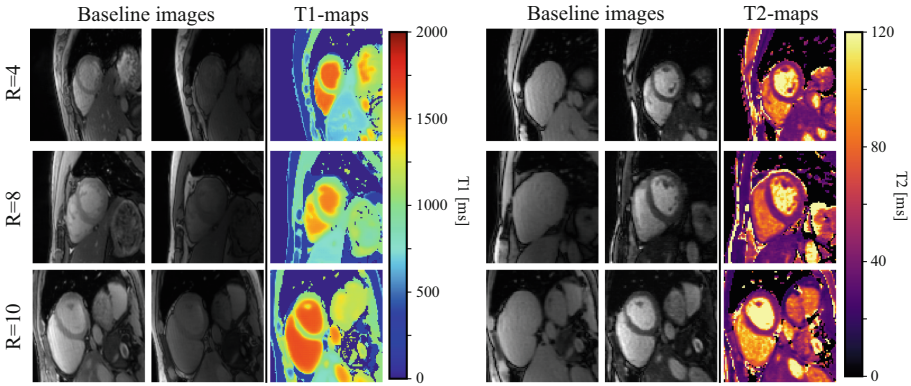


Fig. 2. Qualitative results for T1 and T2 mapping sequences. The baseline images with the shortest and longest inversion/echo times are shown. The proposed method can generate both images and quantitative maps simultaneously. Perceptually, the reconstructed images of all acceleration factors are of good quality.

phase. Therefore, the effect of introducing MR relaxometry-related terms in the loss function needs further investigation because the discrepancy between the predicted and the ground truth quantitative maps can not be evaluated. From Table 2, we observe a performance drop with the increase of acceleration factor. However, it can barely be perceived from the qualitative results shown in Fig. 2.

In conclusion, we proposed a learning-based framework for qMRI reconstruction with a variational network as the reconstruction backbone and introduced an additional mapping network. The proposed framework can output both the baseline images and the mapping result simultaneously. Choosing U-Net as the regularizer achieved better performance, which is further improved by introducing MR relaxometry.

References

1. Barbieri, M., et al.: A deep learning approach for magnetic resonance fingerprinting: scaling capabilities and good training practices investigated by simulations. *Phys. Med.* **89**, 80–92 (2021)
2. Donoho, D.L.: Compressed sensing. *IEEE Trans. Inf. Theory* **52**(4), 1289–1306 (2006)
3. Eliasi, P.A., Feng, L., Otazo, R., Rangan, S.: Fast magnetic resonance parametric imaging via structured low-rank matrix reconstruction. In: 2014 48th Asilomar Conference on Signals, Systems and Computers, pp. 423–428. IEEE (2014)
4. Fabian, Z., Heckel, R., Soltanolkotabi, M.: Data augmentation for deep learning based accelerated MRI reconstruction with limited data. In: International Conference on Machine Learning, pp. 3057–3067. PMLR (2021)
5. Griswold, M.A., et al.: Generalized autocalibrating partially parallel acquisitions (GRAPPA). *Magn. Reson. Med.: Off. J. Int. Soc. Magn. Reson. Med.* **47**(6), 1202–1210 (2002)
6. Haacke, E.M.: Magnetic resonance imaging: physical principles and sequence design (1999)
7. Haaf, P., Garg, P., Messroghli, D.R., Broadbent, D.A., Greenwood, J.P., Plein, S.: Cardiac T1 mapping and extracellular volume (ECV) in clinical practice: a comprehensive review. *J. Cardiovasc. Magn. Reson.* **18**(1), 1–12 (2017)
8. Hammernik, K., et al.: Learning a variational network for reconstruction of accelerated MRI data. *Magn. Reson. Med.* **79**(6), 3055–3071 (2018)
9. Heidemann, R.M., et al.: A brief review of parallel magnetic resonance imaging. *Eur. Radiol.* **13**, 2323–2337 (2003)
10. Huizinga, W., et al.: PCA-based groupwise image registration for quantitative MRI. *Med. Image Anal.* **29**, 65–78 (2016)
11. Hyun, C.M., Kim, H.P., Lee, S.M., Lee, S., Seo, J.K.: Deep learning for undersampled mri reconstruction. *Phys. Med. Biol.* **63**(13), 135007 (2018)
12. Larkman, D.J., Nunes, R.G.: Parallel magnetic resonance imaging. *Phys. Med. Biol.* **52**(7), R15 (2007)
13. Larsson, E.G., Erdogmus, D., Yan, R., Principe, J.C., Fitzsimmons, J.R.: Snr-optimality of sum-of-squares reconstruction for phased-array magnetic resonance imaging. *J. Magn. Reson.* **163**(1), 121–123 (2003)
14. Liu, F., Feng, L., Kijowski, R.: MANTIS: model-augmented neural network with incoherent k-space sampling for efficient MR parameter mapping. *Magn. Reson. Med.* **82**(1), 174–188 (2019)
15. Liu, F., Kijowski, R., El Fakhri, G., Feng, L.: Magnetic resonance parameter mapping using model-guided self-supervised deep learning. *Magn. Reson. Med.* **85**(6), 3211–3226 (2021)
16. Lønning, K., Putzky, P., Sonke, J.J., Reneman, L., Caan, M.W., Welling, M.: Recurrent inference machines for reconstructing heterogeneous MRI data. *Med. Image Anal.* **53**, 64–78 (2019)
17. Lustig, M., Donoho, D., Pauly, J.M.: Sparse MRI: the application of compressed sensing for rapid MR imaging. *Magn. Reson. Med.: Off. J. Int. Soc. Magn. Reson. Med.* **58**(6), 1182–1195 (2007)
18. Messroghli, D.R., Radjenovic, A., Kozerke, S., Higgins, D.M., Sivananthan, M.U., Ridgway, J.P.: Modified look-locker inversion recovery (MOLLI) for high-resolution t1 mapping of the heart. *Magn. Reson. Med.: Off. J. Int. Soc. Magn. Reson. Med.* **52**(1), 141–146 (2004)

19. O'Brien, A.T., Gil, K.E., Varghese, J., Simonetti, O.P., Zareba, K.M.: T2 mapping in myocardial disease: a comprehensive review. *J. Cardiovasc. Magn. Reson.* **24**(1), 1–25 (2022)
20. Pruessmann, K.P., Weiger, M., Börnert, P., Boesiger, P.: Advances in sensitivity encoding with arbitrary k-space trajectories. *Magn. Reson. Med.: Off. J. Int. Soc. Magn. Reson. Med.* **46**(4), 638–651 (2001)
21. Putzky, P., Welling, M.: Recurrent inference machines for solving inverse problems. arXiv preprint [arXiv:1706.04008](https://arxiv.org/abs/1706.04008) (2017)
22. Ronneberger, O., Fischer, P., Brox, T.: U-net: convolutional networks for biomedical image segmentation. In: Navab, N., Hornegger, J., Wells, W.M., Frangi, A.F. (eds.) *MICCAI 2015, Part III*. LNCS, vol. 9351, pp. 234–241. Springer, Cham (2015). https://doi.org/10.1007/978-3-319-24574-4_28
23. Seraphim, A., Knott, K.D., Augusto, J., Bhuva, A.N., Manisty, C., Moon, J.C.: Quantitative cardiac MRI. *J. Magn. Reson. Imaging* **51**(3), 693–711 (2020)
24. Shafieizargar, B., Byanju, R., Sijbers, J., Klein, S., den Dekker, A.J., Poot, D.H.: Systematic review of reconstruction techniques for accelerated quantitative MRI. *Magn. Reson. Med.* (2023)
25. Sriram, A., et al.: End-to-end variational networks for accelerated MRI reconstruction. In: Martel, A.L., et al. (eds.) *MICCAI 2020, Part II*. LNCS, vol. 12262, pp. 64–73. Springer, Cham (2020). https://doi.org/10.1007/978-3-030-59713-9_7
26. Wang, C., et al.: Recommendation for cardiac magnetic resonance imaging-based phenotypic study: imaging part. *Phenomics* **1**, 151–170 (2021)
27. Wang, C., et al.: CMRxRecon: an open cardiac MRI dataset for the competition of accelerated image reconstruction. arXiv preprint [arXiv:2309.10836](https://arxiv.org/abs/2309.10836) (2023)
28. Wang, Z., Bovik, A.C., Sheikh, H.R., Simoncelli, E.P.: Image quality assessment: from error visibility to structural similarity. *IEEE Trans. Image Process.* **13**(4), 600–612 (2004)
29. Yang, Y., Sun, J., Li, H., Xu, Z.: ADMM-CSNet: a deep learning approach for image compressive sensing. *IEEE Trans. Pattern Anal. Mach. Intell.* **42**(3), 521–538 (2018)
30. Yiasemis, G., Sonke, J.J., Sánchez, C., Teuwen, J.: Recurrent variational network: a deep learning inverse problem solver applied to the task of accelerated MRI reconstruction. In: *Proceedings of the IEEE/CVF Conference on Computer Vision and Pattern Recognition*, pp. 732–741 (2022)
31. Zhu, B., Liu, J.Z., Cauley, S.F., Rosen, B.R., Rosen, M.S.: Image reconstruction by domain-transform manifold learning. *Nature* **555**(7697), 487–492 (2018)