

Global k-Space Interpolation for Dynamic MRI

Reconstruction using Masked Image Modeling

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Project Page



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Conventional MR Reconstruction with Image-domain Denoising

Typical Works:
 DcCNN [Schlemper TMI2018]
 VN-Net [Hammernik MRM18]
 CRNN [Qin TMI2018]
 MoDL [Aggarwal TMI2018]
 ...

Major Limitations:

1. **Lack of generalizability** due to artifacts-specific priors/regularizers
2. Artifacts **distort** and/or **obscure** tissues of interest before denoising kick-in
3. Work on the **corrupted data**

MR Reconstruction via k-Space Interpolation (work on reliable data without any corruption)

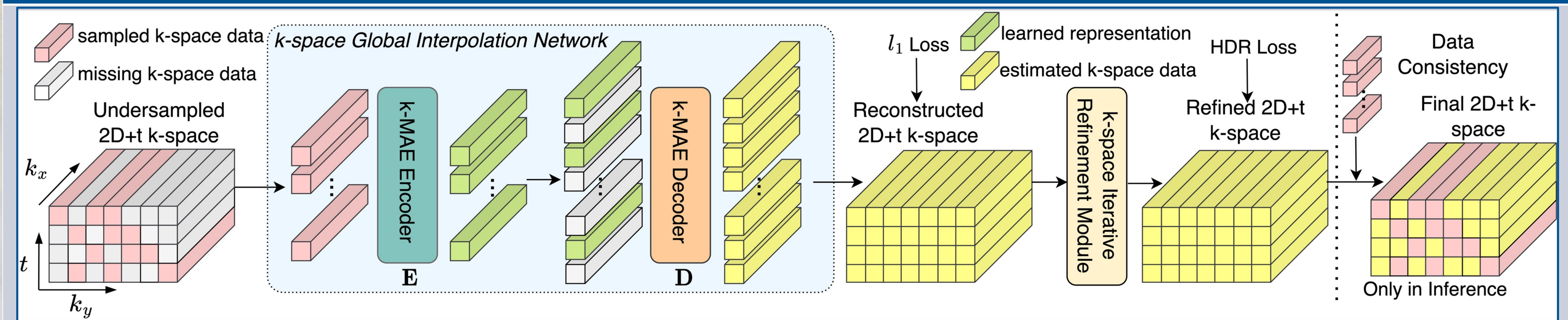
Typical Works:
 GRAPPA [Griswold MRM18]
 SPIRiT [Lustig MRM2010]
 ALOHA [Han TMI2019]
 RAKI [Akçakaya MRM19]
 ...

Limitations of current k-Space Interpolation methods:

1. **Limited Flexibility** due to Calibration Signals
2. Only **Local** Operators
3. **No Representation** of the Sampled Data

In this work, we proposed k-Space interpolation method -- k-GIN 🍷

(Btw. after GRAPPA, RAKI, Caipirinha, SAKE etc., We have one more alcoholic beverage in MR Reconstruction) 😎



The Tool We apply – Masked Autoencoders (MAE)

He et al. CVPR2022

Gist of the work:

- **First usage of MAE** in MR reconstruction and in k-space
- **Global** Interpolation based on **learned representation** from the sampled data
- **More robust** than image domain-based methods at different acceleration rates

Some details

- k_x is used as channel dimension
- Iterative refinement applied on every plane (x-y, y-t, x-t)
- High-Dynamic Range (HDR) loss handle the large magnitude difference in k-space

Reconstruction Results (training at R4, infer at R4 and R8 (unseen))

More robust to unseen data than image domain-based method

Outlook

1. Enable Multi-coils setting
2. Flexible input
3. Computational efficient model
4. Downstream task e.g. disease classification, cardiac volume regression and segmentation