

MOTIVATION

3D cardiac MR imaging requires long acquisition time and breath-hold. For this reason, the clinical routine is still dominated by multi-slice 2D imaging, which hamper the visualization of anatomy and quantitative measurements as relatively thick slices are acquired. As a solution, we propose a novel image super-resolution (SR) approach that is based on a residual convolutional neural network model. It reconstructs high resolution 3D volumes from 2D image stacks for more accurate image analysis. Our model allows the use of multiple input data acquired from different viewing planes for improved performance.

IMAGE SUPER RESOLUTION PROBLEM

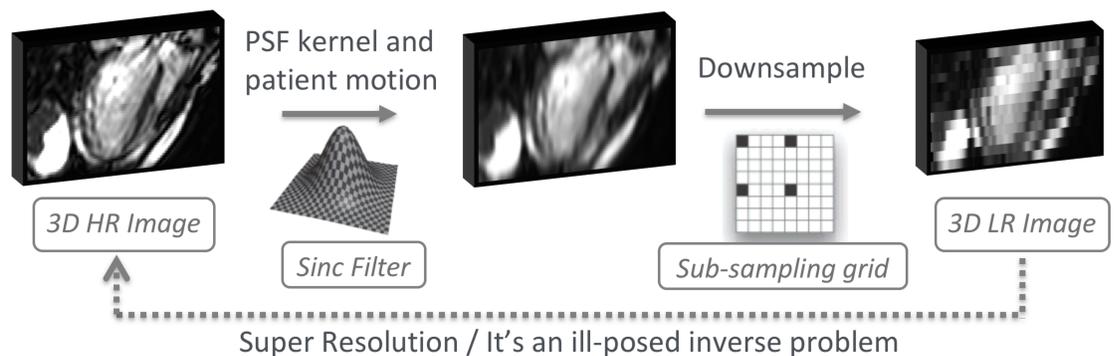


Figure 1: Low and high resolution image model

QUALITATIVE ASSESSMENT

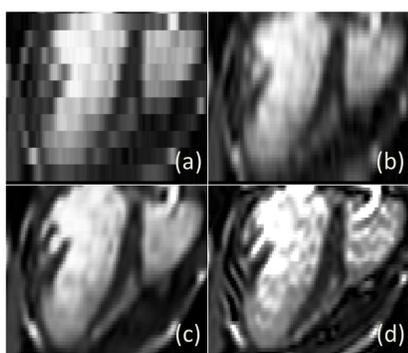


Figure 3: LR image (a), cubic-spline interpolation (b), our model (c), and ground-truth HR image (d).

PROPOSED SUPER RESOLUTION SINGLE INPUT MODEL

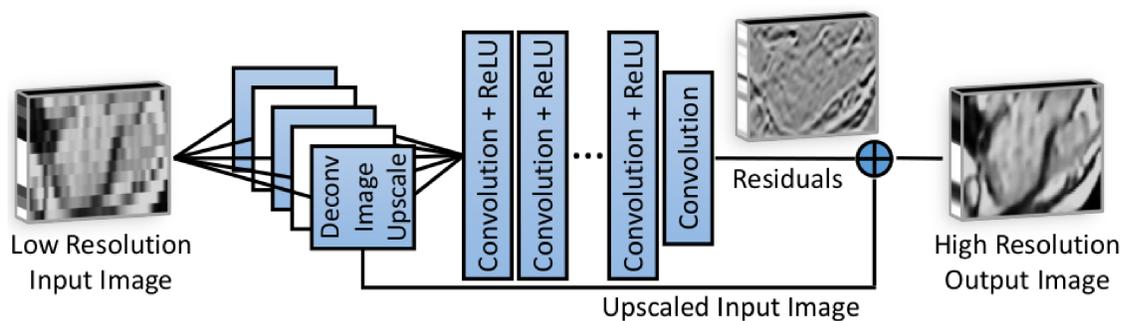


Figure 2: The proposed single input residual super-resolution network model

RESIDUAL LEARNING IN SR

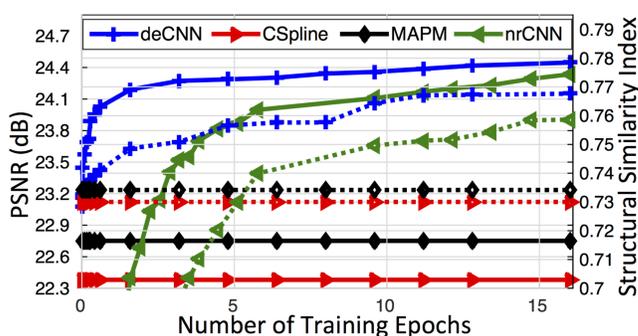


Figure 4: CNN based methods can learn better HR models, and it can be seen that the model without the residual learning (nrCNN) underperforms and requires a large number of training iterations.

We extend the SR model in [1] with:

- (I) Residual learning of 3D kernels
- (II) End-to-end training of upsampling layer
- (III) Multi-input model extension (SAX-LAX)

The proposed model training details:

- (I) 930 LR-HR training image pairs
- (II) Batch-normalization [Ioffe and Szegedy'15]
- (III) Smooth ℓ_1 norm is used as a cost function

QUANTITATIVE RESULTS

Table 1: Quantitative comparison of results obtained with different image upsampling methods on 150 testing images

	PSNR (dB)	SSIM	# Filters/Atlases
Linear	20.83±1.10	.70±.03	-
CSpline	22.38±1.13	.73±.03	-
MAPM	22.75±1.22	.73±.03	350
sh-CNN	23.67±1.18	.74±.02	64,64,32,1
CNN	24.12±1.18	.76±.02	64,64,32,16,8,4,1
de-CNN	24.45±1.20	.77±.02	64,64,32,16,8,4,1

⇒ Image qualities are evaluated based on peak signal-to-noise ratio (PSNR) and structural similarity index measure (SSIM).

⇒ The proposed method achieves better image quality compared to multi-atlas patch match (MAPM) [2], shallow network with 4 layers (sh-CNN) [1], and network without a deconvolution layer (CNN).

Table 2: Quantitative comparison of multi-input network models (A separate testing dataset is used 150 SAX/LAX image pairs)

	PSNR (dB)	SSIM	p - values
de-CNN (SAX)	24.76±0.48	.807±.009	0.005
Siamese (SAX/4CH)	25.13±0.48	.814±.013	0.016
MC (SAX/4CH)	25.15±0.47	.814±.012	0.017
MC (SAX/2/4CH)	25.26±0.37	.818±.012	-

⇒ Multi-input models improve high-resolution image synthesis results. Siamese and multi-channel (MC) input models are compared. 2/4 chamber images (2/4 CH) are processed together with short axis (SAX) stacks.

IMAGE SR & LV TRACKING

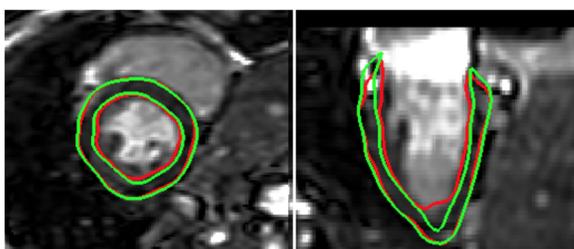


Figure 5: Left ventricular (LV) tracking results (end-systole phase) obtained with linearly interpolated (red) and super-resolved images (green).

IMAGE SR BASED LEFT VENTRICLE SEGMENTATION

Table 3: As a subsequent image analysis, 18 SAX upsampld images are automatically segmented using a state-of-the-art multi-atlas patch fusion method. Those segmentations are compared with the manual annotations performed on HR images. Segmentation results for CSpline ($p = .007$), MAPM ($p = .009$), and deCNN are compared in terms of mean and Hausdorff distances (MYO) and LV cavity volume differences.

	Linear	CSpline	MAPM	de-CNN	High Res
LV Vol Diff (ml)	11.72±6.96	10.80±6.46	9.55±5.42	9.09±5.36	8.24±5.47
Mean Dist (mm)	1.49±0.30	1.45±0.29	1.40±0.29	1.38±0.29	1.38±0.28
Haus Dist (mm)	7.74±1.73	7.29±1.63	6.83±1.61	6.67±1.77	6.70±1.85

REFERENCES

- [1] Dong, C., Loy, C.C., He, K., Tang, X.: Image super-resolution using deep convolutional networks. IEEE PAMI 38(2), 295–307 (2016)
- [2] Shi, W., et al.: Cardiac image super-resolution with global correspondence using multi-atlas patchmatch. In: MICCAI (2013)