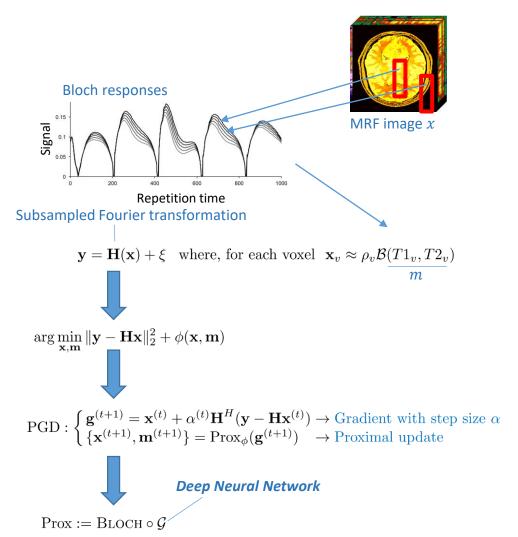
## Compressive MR Fingerprinting Reconstruction with Neural Proximal Gradient Iterations (accepted to MICCAI'20)





Dongdong Chen<sup>1</sup> (d.chen@ed.ac.uk), Mike E. Davies1<sup>1</sup>, Mohammad Golbabaee<sup>2</sup>

<sup>1</sup> School of Engineering, the University of Edinburgh, <sup>2</sup>Department of Computer Science, University of Bath



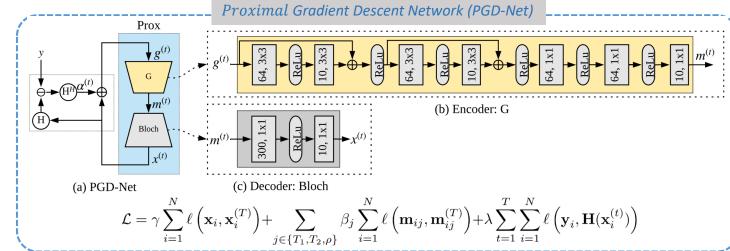


Table 1: Average errors (NRMSE, SSIM, MAE), memory (for storing a dictionary or a network) and runtimes (per image sclice) required for computing T1, T2 and PD maps using the MRF baselines and our PGD-NET algorithm.

## The proposed PGD-Net:

- ☐ State-of-the-art MRF reconstruction
- ☐ interpretable and Physics-engaged
- ☐ Cheaper computation
- ☐ Cheaper memory

	NRMSE			SSIM MAE (msec) time (sec) memory (M			mory (MB)		
	T1	T2	PD   T1	T2	PD   T1	T2			
FGM	0.475	0.354	1.12   0.614	0.652	0.687   350.0	14.6	1.29		8.81
BLIP+FGM	0.230	0.545	0.073   0.886	0.880	0.984   91.7	8.0	79.28		8.81
MRFCNN	0.155	0.158	0.063   0.943	0.972	0.987   80.3	5.4	0.083		4.72
SCQ	0.172	0.177	0.064   0.929	0.967	0.984   91.7	6.1	0.132		464.51
$\overline{\mathcal{G}}$ (encoder alone)	0.142	0.155	0.065   0.948	0.973	0.987   77.1	5.6	0.067		0.55
$\overline{\text{PGD-Net } (T=2)}$	0.104	0.138	0.050   0.973	0.979	0.991   59.9	5.0	0.078		0.57
$\overline{\text{PGD-Net } (T=5)}$	0.100	0.132	0.045   0.975	0.981	0.992   50.8	4.6	0.103	I	0.57

