

AMIR Supplement

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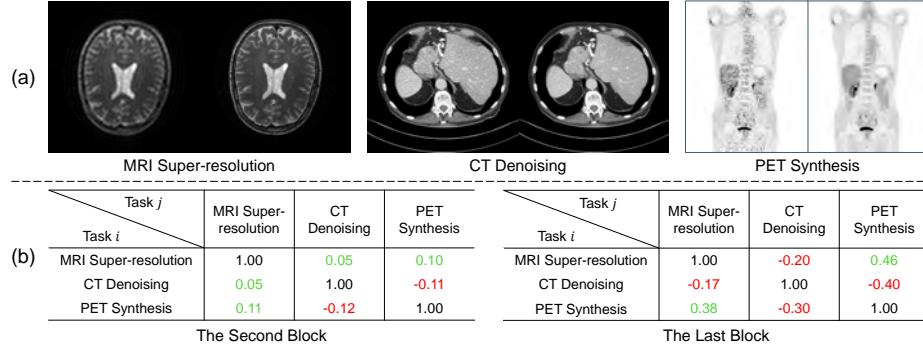


Fig. I. (a) Examples of LQ/HQ pairs for three different MedIR tasks. (b) The interference metric of task *j* on task *i* at the second and last blocks in Restormer. Red values indicate that task *j* negatively impacts task *i*, while green values indicate a positive impact.

Table I. Single task medical image restoration results.

| Method | MRI Super-Resolution | | | Method | CT Denoising | | | Method | PET Synthesis | | |
|-----------|----------------------|---------------|----------------|----------|----------------|---------------|---------------|-------------------|----------------|---------------|---------------|
| | PSNR↑ | SSIM↑ | RMSE↓ | | PSNR↑ | SSIM↑ | RMSE↓ | | PSNR↑ | SSIM↑ | RMSE↓ |
| SRCNN | 28.8067 | 0.8919 | 41.3488 | CNN | 32.7600 | 0.9075 | 9.3928 | Xiang's | 35.9268 | 0.9167 | 0.0980 |
| VDSR | 30.0446 | 0.9140 | 36.0508 | REDCNN | 33.1889 | 0.9113 | 8.9427 | DCNN | 36.2710 | 0.9243 | 0.0954 |
| SwinIR | 31.5549 | 0.9334 | 30.5788 | Eformer | 33.3496 | 0.9175 | 8.8030 | ARGAN | 36.7272 | 0.9406 | 0.0902 |
| Restormer | 31.8474 | 0.9378 | 29.7005 | CTFormer | 33.2506 | 0.9134 | 8.8974 | Spach Transformer | 37.1371 | 0.9456 | 0.0871 |
| AMIR | 31.9923 | 0.9393 | 29.2095 | AMIR | 33.6738 | 0.9183 | 8.4773 | AMIR | 37.2121 | 0.9473 | 0.0863 |

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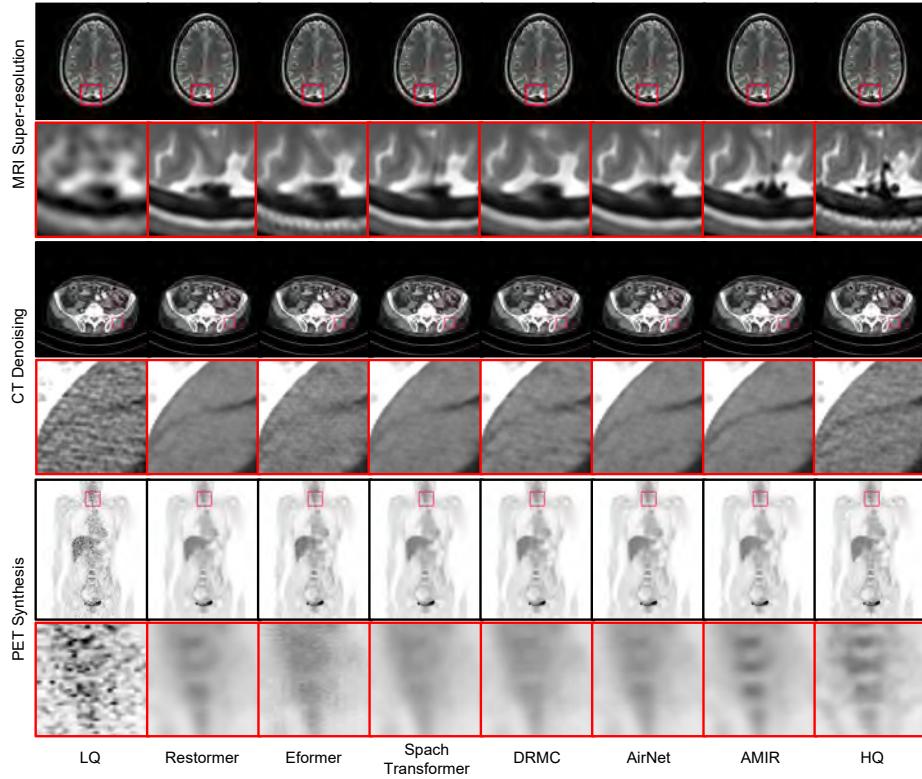


Fig. II. Visual comparison in three different medical image restoration tasks

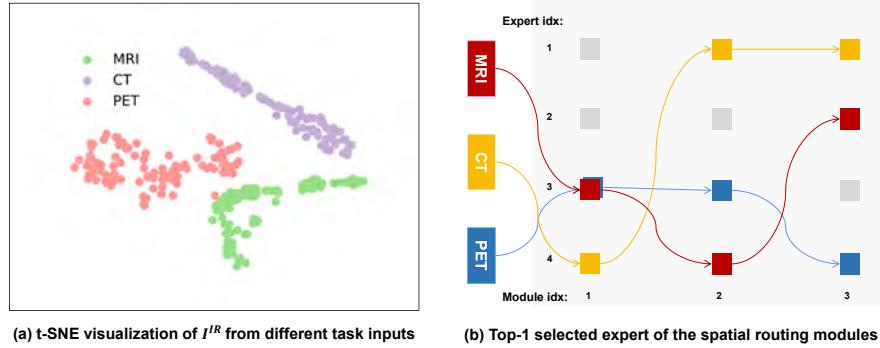


Fig. III. (a) t-SNE visualization of I^{IR} from different tasks, indicating a clear clustering of different task inputs. (b) Top-1 selected expert in each spatial routing module (SRM). In our AMIR network setting, there are 3 SRMs, each incorporating a mixture of experts (MOE) with 4 experts. Within these SRMs, the top-1 selected expert is identified across the 3 SRMs for each task. Remarkably, the top experts selected across SRMs form distinct paths, with variations observed across different tasks.