

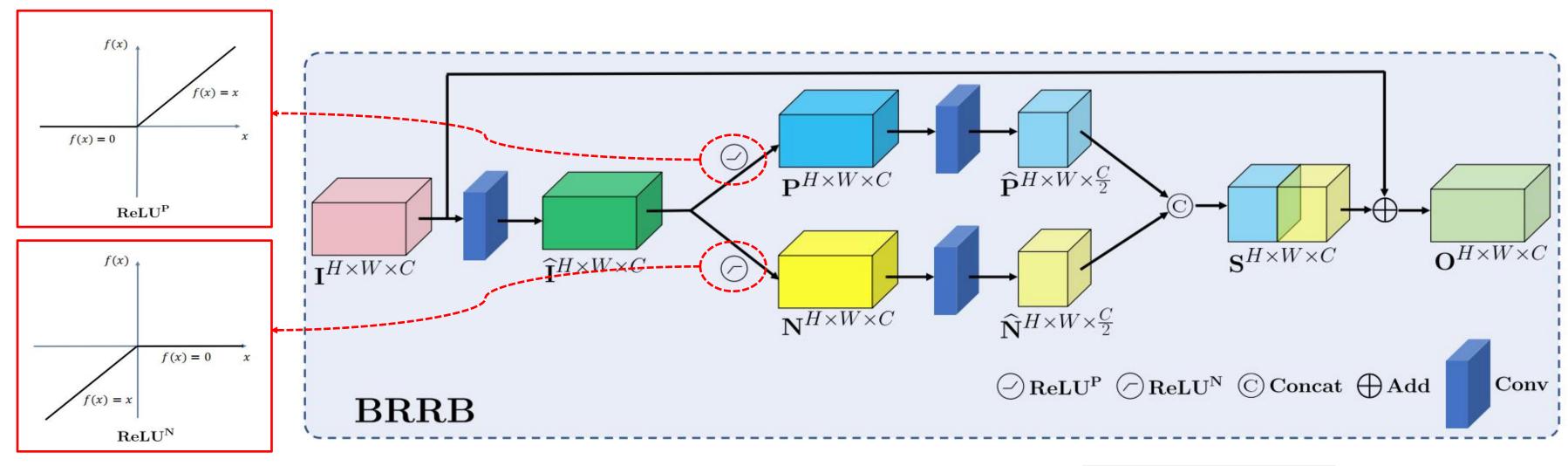
BAM: Bilateral Activation Mechanism for Image Fusion

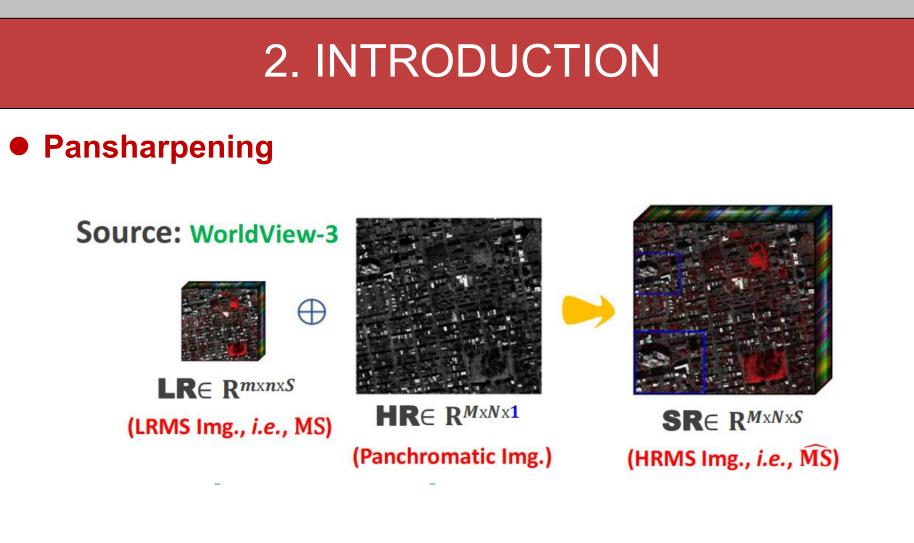
Zi-Rong Jin, Liang-Jian Deng, Tian-Jing Zhang, Xiao-Xu Jin University of Electronic Science and Technology of China, Chendu, China

1. ABSTRACT

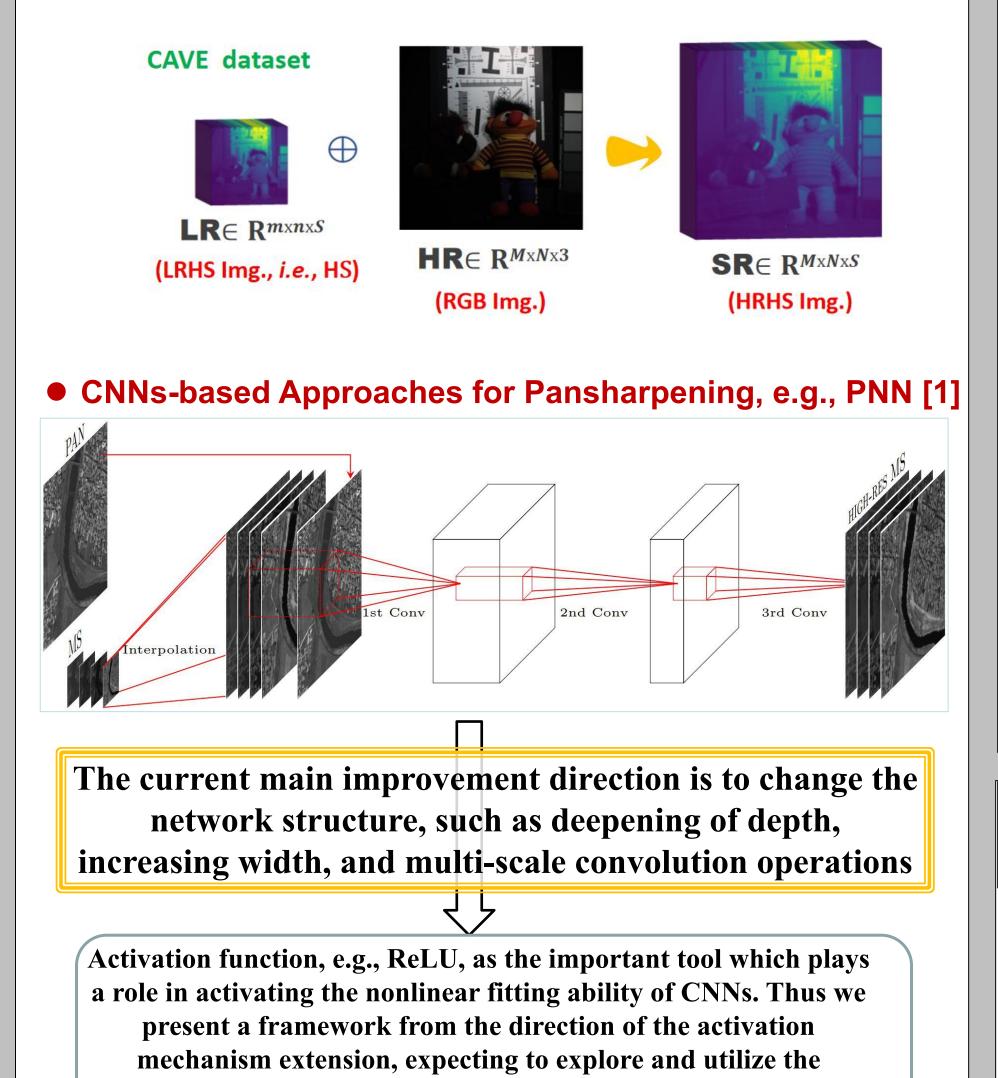
As the conventional activation functions such as ReLU, LeakyReLU, and PReLU, the negative parts in feature maps are simply truncated or linearized, which may result in unflexible structure and undesired information distortion. In this paper, we propose a simple but effective Bilateral Activation Mechanism (BAM) which could be applied to the activation function to offer an efficient feature extraction model. Based on BAM, the Bilateral ReLU Residual Block (BRRB) that still sufficiently keeps the nonlinear characteristic of **ReLU** is constructed to separate the feature maps into two parts, i.e., the positive and negative components, then adaptively represent and extract the features by two independent convolution layers. Besides, our mechanism will not increase any extra parameters or computational burden in the network. We finally embed the BRRB into a basic ResNet architecture, called BRResNet, it is easy to obtain state-of-the-art performance in two image fusion tasks, i.e., pansharpening and hyperspectral image super-resolution (HISR). Additionally, deeper analysis and ablation study demonstrate the effectiveness of BAM, the lightweight property of the network, etc.

o Bilateral ReLU Residual Block (BRRB)





• Hyperspectral Image Super-Resolution (HISR)





$\mathcal{L}(\theta) = \frac{1}{N} \sum_{N} \left\| \mathcal{F}_{\theta}(\widehat{\mathbf{LR}}^{(i)}, \mathbf{HR}^{(i)}) + \widehat{\mathbf{LR}} - \mathbf{GT}^{(i)} \right\|_{F}^{2}$

ACM multimedia

• Contribution

 \mathbf{SR}

Conv

 \bigcirc Concat \oplus Add

As a mechanism, BAM provides a more efficient feature extraction mode without increase the computational burden. Also, it has many variants and can be used as a substitution to replace any structure like "Activation + Convolution", giving us more flexibility in designing the network structure.

5. EXPERIMENT SETTING

M

\Box HISR:

1) Datasets:

 \mathbf{LR}

CAVE dataset

○ BRResNet

 \mathbf{HR}

ÎR

□ HAVARD dataset

2) Metrics: SAM, PSNR, SSIM, ERGAS

D Pansharpening:

1) **Datasets:**

• We introduce a simple but effective Bilateral Activation

Mechanism (BAM) that not only retains the nonlinearity of the activation function but also avoids information distortion caused by inactivation.

7. CONCLUSIONS

• A network with residual structure using BAM with ReLU (BRResNet) is proposed, which significantly improves the efficiency of feature extraction in image fusion tasks.

• More development history of CNN can be found from [2]

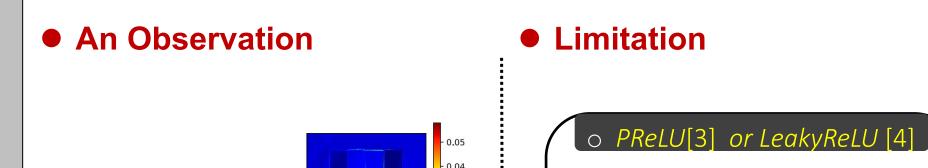
features that can not be activated while retaining the nonlinearity

[1] G. Masi, D. Cozzolino, L. Verdoliva, and G. Scarpa, "Pansharpening by convolutional neural networks," Remote Sensing, vol. 8, pp. 594, 2016.

[2] Vivone, G., et al. "A New Benchmark Based on Recent Advances in Multispectral Pansharpening: Revisiting pansharpening with classical and emerging pansharpening methods. IEEE Geoscience and Remote Sensing Magazine.

3. MOTIVATION

f(y)

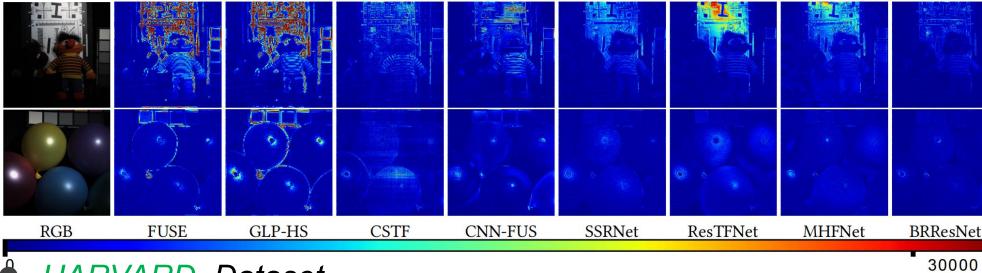


■ 8-band data: WorldView-3 (WV3)

- 1) reduced-resolution examples
- 2) full-resolution examples
- □ <u>4-band data</u>: Quikbird (QB), GaoFen2 (GF2)
- 2) Metrics:
- □ <u>Reduced-resolution</u>: SAM, ERGAS, SCC, Q8
- **\Box** <u>Full-resolution</u>: QNR, D_{λ}, D_S

6. RESULTS

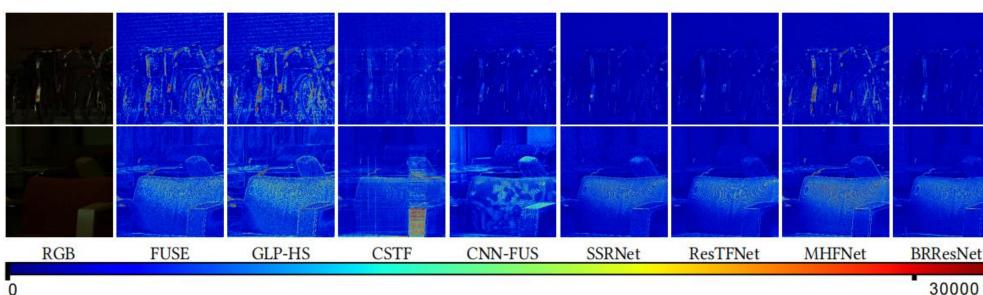
CAVE Dataset



HARVARD Dataset

BDSD

EXP



WV3 Reduced-Resolution Dataset

• Proposed BRResNet achieves state-of-the-art performance in two fusion tasks. Especially, the given BRResNet holds a large margin among other CNNs-based methods in terms of the parameters, thus can be viewed as a lightweight network.

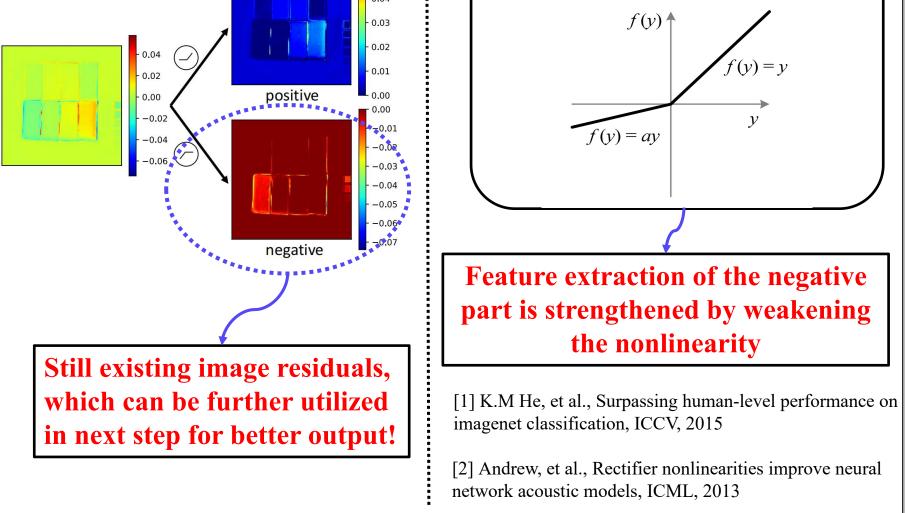
Table 1: Average quantitative comparisons on 11 CAVE examples (Red: the best; Blue: the second best).

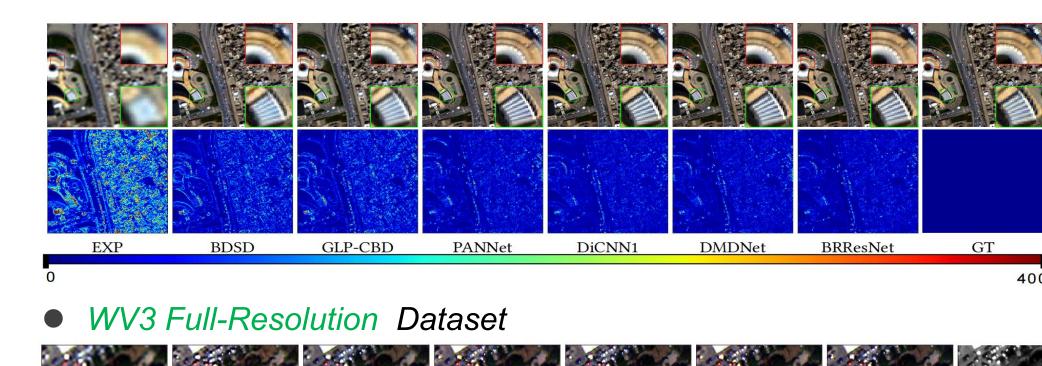
Method	PSNR	SAM	ERGAS	SSIM
FUSE	39.72 ± 3.52	5.83 ± 2.02	4.18 ± 3.08	0.975 ± 0.018
GLP-HS	37.81 ± 3.06	5.36 ± 1.78	4.66 ± 2.71	0.972 ± 0.015
CSTF	42.14 ± 3.04	9.92 ± 4.11	3.08 ± 1.56	0.964 ± 0.027
CNN-FUSE	42.66 ± 3.46	6.44 ± 2.31	2.95 ± 2.24	0.982 ± 0.007
SSRNet	45.28 ± 3.13	4.72 ± 1.76	2.06 ± 1.30	0.990 ± 0.004
ResTFNet	45.35 ± 3.68	3.76 ± 1.31	1.98 ± 1.62	0.993 ± 0.003
MHFNet	46.32 ± 2.76	4.33 ± 1.48	<u>1.74 ± 1.44</u>	0.992 ± 0.006
BRResNet (Our)	47.85 ± 3.56	2.96 ± 0.89	1.50 ± 1.18	0.995 ± 0.003
Ideal value	00	0	0	1

	Table 2: Average quantitative comparisons on 10 Harvard examples.
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Method	PSNR	SAM	ERGAS	SSIM
FUSE	42.06 ± 2.94	3.23 ± 0.91	3.14 ± 1.52	0.977 ± 0.009
GLP-HS	40.14 ± 3.22	3.52 ± 0.96	3.74 ± 1.44	0.966 ± 0.012
CSTF	42.97 ± 3.33	3.30 ± 1.25	2.43 ± 1.06	0.972 ± 0.021
CNN-FUSE	43.61 ± 4.73	3.32 ± 1.17	2.78 ± 1.64	0.978 ± 0.016
SSRNet	44.40 ± 3.49	2.61 ± 0.72	2.39 ± 1.02	0.985 ± 0.007
ResTFNet	44.47 ± 4.04	2.56 ± 0.68	2.21 ± 0.87	0.985 ± 0.008
MHFNet	43.10 ± 3.94	2.76 ± 0.77	3.28 ± 1.54	0.977 ± 0.009
BRResNet (Our)	45.74 ± 3.86	2.39 ± 0.66	1.94 ± 0.69	0.986 ± 0.009
Ideal value	8	0	0	1

Table 3: Average quantitative comparisons on 1258 reduced resolu-





PANNet

DiCNN1

GLP-CBD

DMDNet

BRResNet

PAN

tion WV3 examples.

Ideal value	0	0	1	1
BRResNet (Our)	3.5881 ± 1.2185	2.4618 ± 0.9306	0.9612 ± 0.0444	0.9183 ± 0.1099
DMDNet	3.9714 ± 1.2482	2.8572 ± 0.9663	0.9527 ± 0.0447	0.9000 ± 0.1142
DiCNN1	3.9805 ± 1.3181	2.7367 ± 1.0156	0.9517 ± 0.0472	0.9097 ± 0.1117
PanNet	4.0921 ± 1.2733	2.9524 ± 0.9778	0.9495 ± 0.0461	0.8942 ± 0.1170
GLP-CBD	5.2861 ± 1.9582	4.1627 ± 1.7748	0.8904 ± 0.0698	0.8540 ± 0.1144
BDSD	6.9997 ± 2.8530	5.1670 ± 2.2475	0.8712 ± 0.0798	0.8126 ± 0.1234
Method	SAM	ERGAS	SCC	Q8

Table 4: Average quantitative comparisons on 36 full resolution WV3 examples.

Method	QNR	D_{λ}	D_s
BDSD	0.9368 ± 0.0416	0.0170 ± 0.0137	0.0473 ± 0.0320
GLP-CBD	0.9107 ± 0.0518	0.0323 ± 0.0243	0.0597 ± 0.0325
PanNet	0.9605 ± 0.01551	0.0215 ± 0.0098	0.0184 ± 0.0074
DiCNN1	0.9454 ± 0.0268	0.0181 ± 0.0135	0.0374 ± 0.0159
DMDNet	0.9595 ± 0.0155	0.0201 ± 0.0098	0.0209 ± 0.0073
BRResNet	0.9671 ± 0.0095	0.0179 ± 0.0063	0.0152 ± 0.0050
Ideal value	1	0	0