

# Robust PCA Unrolling Network for Super-Resolution Vessel Extraction in X-Ray Coronary Angiography

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**Abstract**—Although robust PCA has been increasingly adopted to extract vessels from X-ray coronary angiography (XCA) images, challenging problems such as inefficient vessel-sparsity modelling, noisy and dynamic background artefacts, and high computational cost still remain unsolved. Therefore, we propose a novel robust PCA unrolling network with sparse feature selection for super-resolution XCA vessel imaging. Being embedded within a patch-wise spatiotemporal super-resolution framework that is built upon a pooling layer and a convolutional long short-term memory network, the proposed network can not only gradually prune complex vessel-like artefacts and noisy backgrounds in XCA during network training but also iteratively learn and select the high-level spatiotemporal semantic information of moving contrast agents flowing in the XCA-imaged vessels. The experimental results show that the proposed method significantly outperforms state-of-the-art methods, especially in the imaging of the vessel network and its distal vessels, by restoring the intensity and geometry profiles of heterogeneous vessels against complex and dynamic backgrounds. The source code is available at <https://github.com/Binjie-Qin/RPCA-UNet>

**Index Terms**—Algorithm unrolling, RPCA unrolling network, X-ray coronary angiography, vessel extraction, sparse feature selection, super-resolution.

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## I. INTRODUCTION

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## II. RELATED WORKS

### A. XCA Vessel Extraction

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29, 30 2D/3D 29, 31

An CA

32 CA

33, 34, C



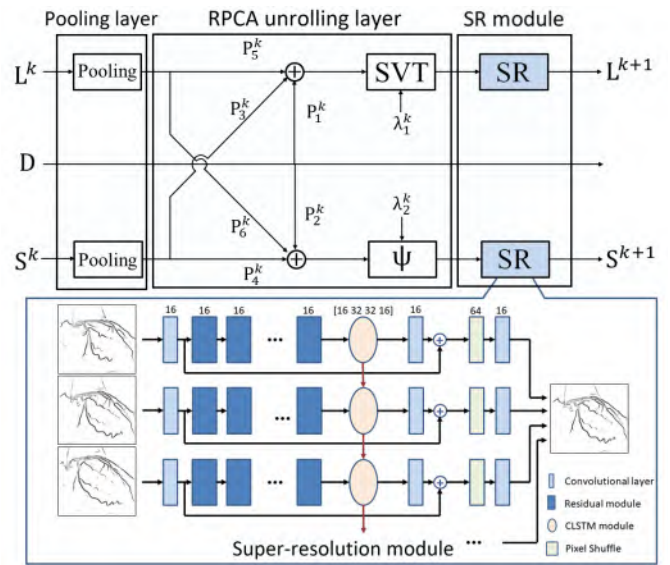


Fig. 1. The architecture of a single iteration/layer of RPCA-UNet for decomposing XCA data  $D$  into vessel ( $S$ ) and background ( $L$ ) components, which consists of a pooling layer, an RPCA unrolling layer, and an SR module. The SR module is mainly built upon the convolutional layer, residual module and CLSTM network.

$$D = H_1 + H_2 + H_3 + \dots + H_m + \lambda_1 L + \lambda_2 S \quad (1)$$

A. RPCA Modelling

The RPCA model is used to decompose the XCA data  $D$  into vessel ( $S$ ) and background ( $L$ ) components. The RPCA model is defined as follows:

$$D = H_1 + H_2 + H_3 + \dots + H_m + \lambda_1 L + \lambda_2 S \quad (1)$$

$$D = H_1 + H_2 + H_3 + \dots + H_m + \lambda_1 L + \lambda_2 S \quad (2)$$

III. METHOD

A. RPCA Modelling







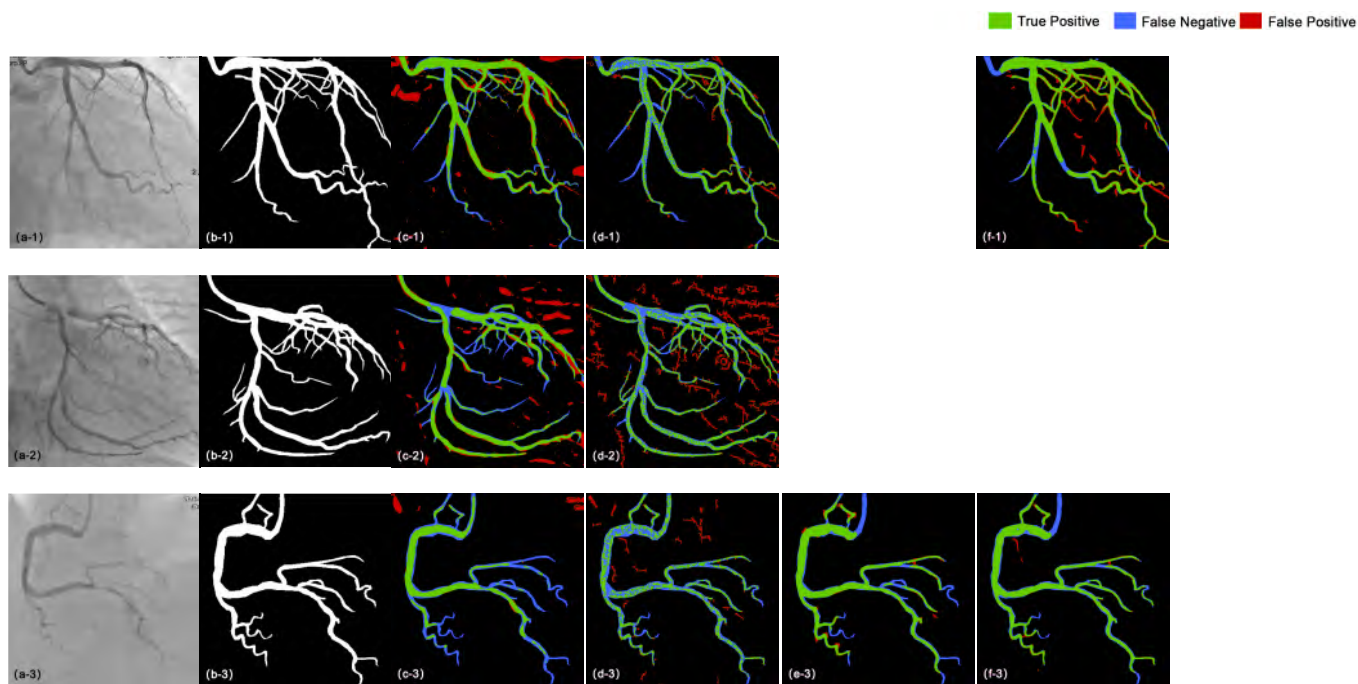


Fig. 3. XCA vessel segmentation results. Pixels labelled with green, blue, and red colours represent true positive pixels, false negative pixels, and false positive pixels, respectively. (a) Original XCA image; (b) Ground-truth vessel mask; (c) Frangi's; (d) Coye's; (e) SVS-net; (f) CS<sup>2</sup>-Net; (g) RPCA-UNet.

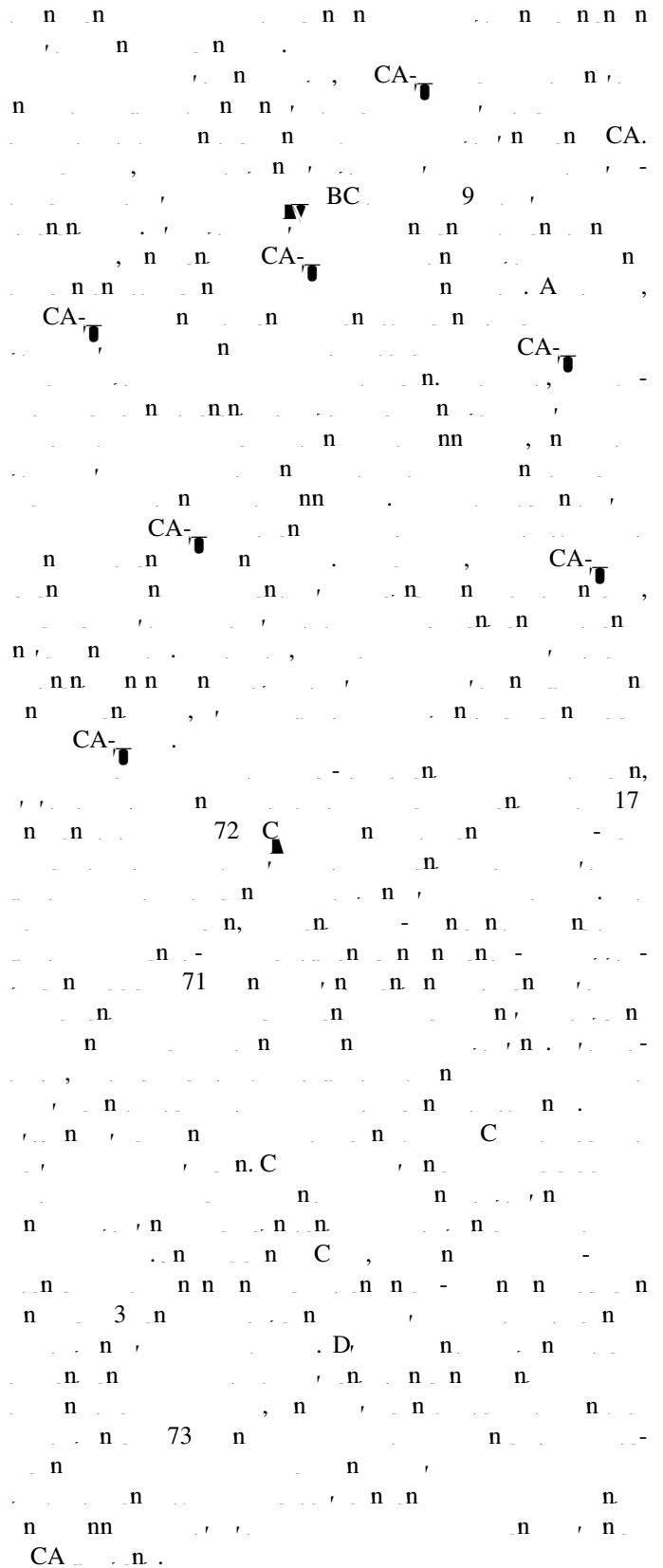
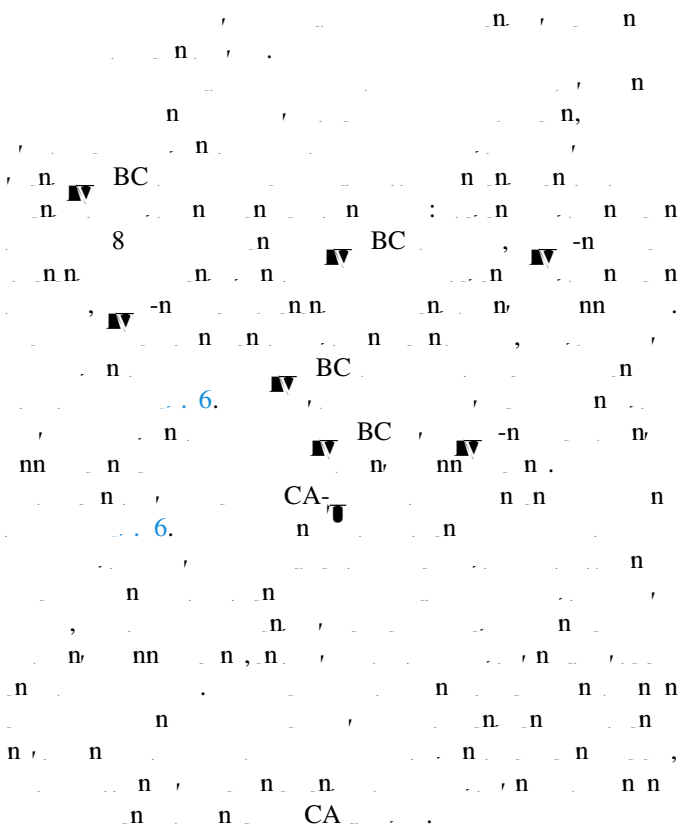
TABLE I

PERFORMANCE OF DIFFERENT VESSEL EXTRACTION METHODS  
IN TERMS OF CNR VALUES (MEAN  $\pm$  STANDARD DEVIATION)





Fig. 6. The effect of coarse versus fine vessel label on the result of weakly supervised learning. The first row are the coarse and fine grey value labels automatically generated by the VRBC combined with different binary vessel mask segmentations, i.e. from left to right being original segmentation method in the VRBC, SVS-net with training data generated by the original segmentation method, SVS-net with training data by manual annotation; the second row of results are test cases of the corresponding networks trained with different grey value labels.



V. CONCLUSION AND DISCUSSION

ACKNOWLEDGMENT



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