

# MULTI-HEAD FEATURE PYRAMID NETWORKS FOR BREAST MASS DETECTION

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## ABSTRACT

Analysis of X-ray images is one of the main tools to diagnose breast cancer. The ability to quickly and accurately detect the location of masses from the huge amount of image data is the key to reducing the morbidity and mortality of breast cancer. Currently, the main factor limiting the accuracy of breast mass detection is the unequal focus on the mass boxes, leading the network to focus too much on larger masses at the expense of smaller ones. In the paper, we propose the multi-head feature pyramid module (MHFPN) to solve the problem of unbalanced focus of target boxes during feature map fusion and design a multi-head breast mass detection network (MBMDnet). Experimental studies show that, comparing to the SOTA detection baselines, our method improves by 6.58% (in AP@50) and 5.4% (in TPR@50) on the commonly used INbreast dataset, while about 6-8% improvements (in AP@20) are also observed on the public MIAS and BCS-DBT datasets.

**Index Terms**— Breast Mass Detection, Feature Pyramid Networks, Multi-head Integration, Faster RCNN

## 1. INTRODUCTION

Breast cancer has become a major disease threatening women's health in terms of morbidity and mortality. According to the statistics [1], in 2018 alone, the number of new cases reached 2,088,849 (11.6%) and deaths were 626,679 (6.6%). By comparing the morbidity and mortality rates of breast cancer, it is clear that timely diagnosis and treatment at an early stage can significantly reduce the mortality rate [2]. Currently, the detection of breast masses is mainly divided into: ultrasound image detection, X-ray image detection and biopsy. Among them, X-ray has become the main modality for the early detection of breast cancer due to its outstanding imaging performance. However, X-ray images require specialized radiologists to screen a large number of Digital Imaging and Communications in Medicine (DICOM) files, and the process of resolving X-ray images is very tedious and time-consuming [3]. In order to improve the accuracy and efficiency of breast mass detection, many computer-aided

detection methods based on deep learning have been rapidly developed and achieved good results [4–7].

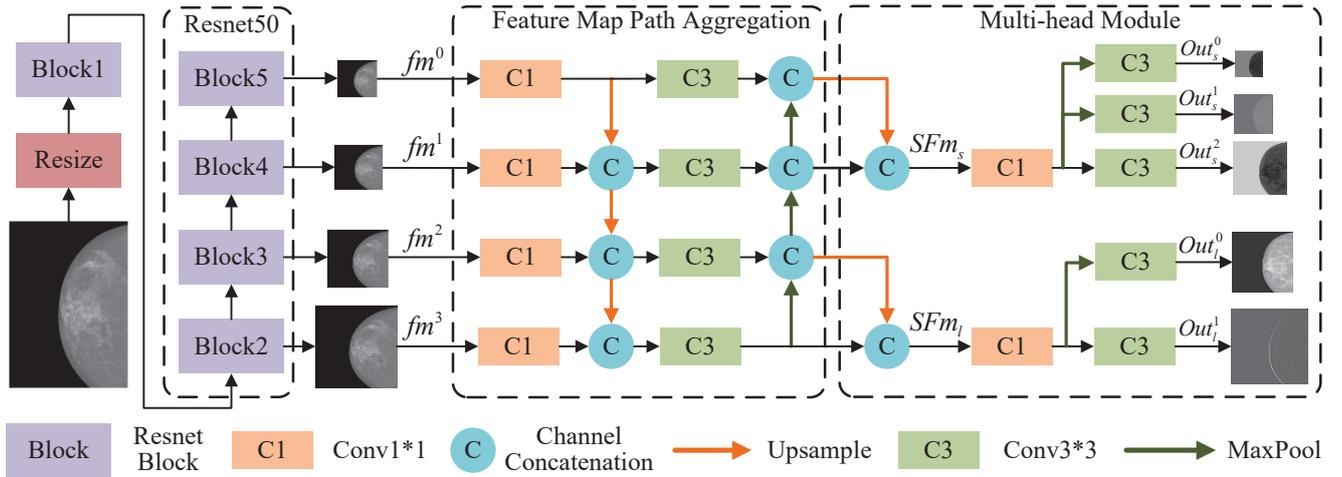
Most of the breast mass detection studies are based on the classical network to adapt the network structure to the characteristics of X-ray images. L. Zhang et al. [4] proposed an Anchor-free YOLOv3 network to alleviate the imbalance between positive boxes and negative boxes that occurs in breast mass detection. B. Ibromkhimov et al. [5] proposed a slider-cutting image method based on Faster RCNN to generate small square patches to solve the problem of excessive X-ray image size and resolution. Y. Wu et al. [8] first proposed the use of the Faster RCNN network based on hierarchical candidate frames for breast mass detection. The approach not only reduces the amount of data to be processed but also improves the accuracy of detection. H. Cao et al. [9] then proposed a new normalization approach and image enhancement algorithm to further improve the detection accuracy of breast masses based on the FSAF network structure. The aforementioned methods have achieved excellent performance on public datasets such as INbreast [10] or private datasets; however, they have a common shortcoming in that they focus more on the features of large masses and ignore those of small masses during feature fusion, which thus leads to serious medical misdiagnosis and may endanger the health of patients in clinical practices.

To solve the unequal focus in the feature fusion of breast X-ray images in target detection, We propose a parallel multi-head feature pyramid module (MHFPN). By adding two additional aggregated focus heads, the network can more easily focus on breast masses that make up a smaller percentage of the image, resulting in improved detection accuracy. Based on MHFPN, Faster RCNN is selected as the detection network. By testing on three publicly available breast mass datasets, our method can achieve the same performance as the previously mentioned X-ray breast image detection studies with a simpler network structure.

In general, our contributions are three-folded: (1) We design a general MHFPN on the feature map fusion layer to alleviate the problem of unequal focus in the feature fusion stage. (2) We propose a multi-headed breast mass detection network (MBMDnet) to improve the accuracy of mass target detection

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**Fig. 1.** Overview of multi-head high resolution FPN network. Since the head part of the detection network follows the original structure, the head part has been omitted from the overall structure diagram.

and reduce the miss detection rate. (3) Experiments show that our method improves by 6.58% (in AP@50) and 5.4% (in TPR@50) on the commonly used INbreast dataset, and 6%-8% (in AP@20) on the MIAS and BCS-DBT datasets.

## 2. METHODS

### 2.1. Overall Structure

Inspired by PANet [17], HRFPN [18] and multi-scale in the segmentation [19], we propose a parallel multi-head feature pyramid network to improve the accuracy of breast mass detection in X-Ray images. The network consists of three parts, the backbone for extracting features, the feature map path aggregation, and the multi-head module. The overall structure is shown in Fig 1, where the network is Faster RCNN, and the backbone is selected as Resnet50. At the model input, the image size of the INbreast dataset is  $3328 \times 4084$ , which is oversized for the existing target detection network breast X-ray images. Due to the limitation of computational volume and operation speed, MHFPN uses resized images ( $1333 \times 800$ ) as the input of the network. Then, the images are fed into the structure of the backbone to obtain feature maps (Denoted as  $fm^i, i \in (0, 3)$ ) at different scales. After that, the feature map is sent through a path aggregation network of successive up and down paths to obtain the richer feature maps (Denoted as  $P(fm), f \in (fm^0, fm^3)$ ).

### 2.2. Multi-head Module

After the multi-resolution feature maps are aggregated by the feature map paths composed of multiple pooling and convolution layers, the feature information flow is fully fused. Compared with the feature-extracted  $fm^i$ ,  $P(fm^i)$  not only carries the precise location information in the bottom information, but also retains the useful information in each feature level. Because of the long path way of information experienced, the X-ray breast feature map suffers from a distraction problem, making the focus of masses with a small image ratio much smaller than that of masses with a large image ratio. Therefore, we designed the multi-head module to alleviate the inequality of focus. Through two feature aggregation head-

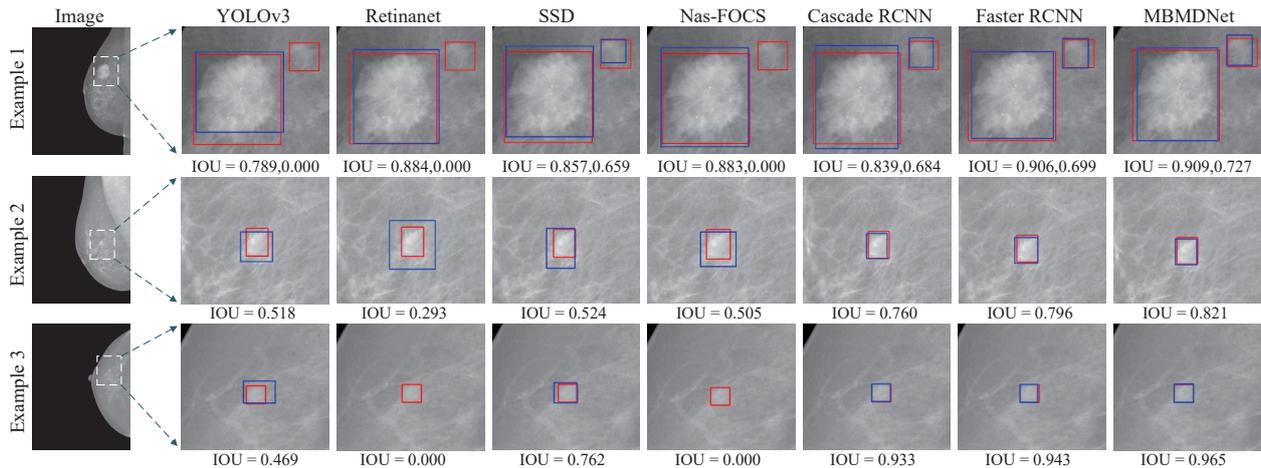
ers, the multi-resolution feature maps are assigned focus using the re-convolution to obtain the aggregated feature maps (Denoted as  $SFm_s$  and  $SFm_l$  in Eq 1). Finally, according to the demand of the network, the feature map resolution is pooled to acquire the output of the corresponding resolution (Denoted as  $Out_s^i, i \in (0, 2)$  and  $Out_l^j, j \in (0, 1)$  in Eq 2).

$$\begin{cases} SFm_s = (P(fm^0) + P(fm^1)), \\ SFm_l = (P(fm^2) + P(fm^3)), \end{cases} \quad (1)$$

$$\begin{cases} Out_s^k = (c^*(p_{2^k}^*(SFm_s)) & k \in (0, 2), \\ Out_l^j = (c^*(p_{2^j}^*(SFm_l)) & j \in (0, 1), \end{cases} \quad (2)$$

where  $c^*$  and  $p^*$  indicate convolution layer and pooling layer, respectively.  $2^k$  and  $2^j$  indicate the size of the dimensions in the pooling layer. Eq 1 expresses the calculation process of the feature map in the two parallel feature aggregation heads. More specifically, the multi-head module obtains two aggregated feature maps by measuring the output feature maps in four different dimensions, interpolating and concatenating the feature maps in adjacent dimensions to the same size. Eq 2 illustrates the process of differentiation from the semantic information-rich  $SFm$  into the corresponding output feature maps. The heads are pooled several times to get the corresponding size.

In particular, in order to achieve targeted recovery of the network's focus on multi-resolution feature maps, we use parallel two-headed modules for high-resolution and low-resolution features separately. Compared with the single-head module (HRFPN), although the multi-head module increases the calculation volume, it can attenuate the effect on focus inequality due to the difference in the ratio of breast mass images. Moreover, the approach can shorten the path of information flow insertion to the final resolution and prevent the feature information from being scattered again. On the feature maps of neighboring dimensions in the same group, the higher-level semantic information is calculated by focus reorganization with neighboring features using up-sampling to retain and improve the influence of higher-level semantic information on  $SFm$ . MHFPN has correspondingly improved the detection effect both on small targets and on large targets, and it works better on small targets.



**Fig. 2.** Visualized results of different models on INbreast, and blue (resp., red) boxes are detection results (resp., ground truths).

**Table 1.** Results of MBMDNet and classical object detection networks on the INbreast dataset

| Model             | AP@50                  | AP@75                  | TPR@50                 | FLOPS   | Params |
|-------------------|------------------------|------------------------|------------------------|---------|--------|
| YOLOv3 [11]       | 0.3700 (0.0242)        | 0.1288 (0.0544)        | 0.8160 (0.0717)        | 19.39G  | 61.53M |
| Retinanet [12]    | 0.3774 (0.0652)        | 0.0920 (0.0325)        | 0.8226 (0.0823)        | 81.89G  | 36.14M |
| SSD [13]          | 0.4892 (0.0889)        | 0.1526 (0.1106)        | 0.7590 (0.1604)        | 87.86G  | 24.53M |
| Nas-FCOS [14]     | 0.4170 (0.1210)        | 0.0872 (0.0367)        | 0.8336 (0.0587)        | 201.57G | 41.13M |
| Cascade RCNN [15] | 0.5212 (0.0509)        | 0.3092 (0.1135)        | 0.7354 (0.0867)        | 244.11G | 68.93M |
| Faster RCNN [16]  | 0.5804 (0.0730)        | <b>0.3160 (0.1577)</b> | 0.7716 (0.0633)        | 216.30G | 41.13M |
| <b>MBMDNet</b>    | <b>0.6462 (0.0934)</b> | 0.3034 (0.1035)        | <b>0.8390 (0.0915)</b> | 291.36G | 44.80M |

**Table 2.** Detection results on three breast masses datasets.

| Dataset      | Method         | AP@20 | TPR@20 |
|--------------|----------------|-------|--------|
| INbreast     | Faster RCNN    | 0.725 | 0.933  |
|              | <b>MBMDNet</b> | 0.806 | 0.962  |
| BCS-DBT [20] | Faster RCNN    | 0.350 | 0.881  |
|              | <b>MBMDNet</b> | 0.412 | 0.878  |
| MIAS [21]    | Faster RCNN    | 0.312 | 0.535  |
|              | <b>MBMDNet</b> | 0.396 | 0.619  |

**Table 3.** Compare to advanced detection models on INbreast.

| Methods                     | TPR@20      | FPPI         |
|-----------------------------|-------------|--------------|
| Amit et al. [22]            | 0.87        | 1.423        |
| Wu et al. [8]               | 0.88        | 0.750        |
| Akselrod-Ballin et al. [23] | 0.93        | 0.560        |
| BMassDNet [9]               | 0.93        | <b>0.495</b> |
| <b>MBMDNet</b>              | <b>0.96</b> | 0.636        |

### 3. EXPERIMENTAL STUDIES

#### 3.1. Experimental Environment and Dataset

All experiments were run on a server with 2 Nvidia GeForce 2080 GPUs with 8192M RAM per card. The server has an Intel(R) Xeon(R) Silver 4110 CPU with 2.10GHz and 16G RAM. For the software configuration, all code was implemented in Python based on the mmdetection framework.

The main dataset used for the experiment is the INbreast dataset<sup>1</sup>, which is more commonly used for breast mass detection. The dataset has 107 images containing 116 masses.

<sup>1</sup><http://medicalresearch.inescporto.pt>

Each image had an average of 1.1 masses, with the smallest mass being  $15 \text{ mm}^2$  and the largest mass measuring  $3689 \text{ mm}^2$ . Meanwhile, in order to test the performance of the network on other different types of datasets, the earlier MIAS dataset<sup>2</sup> and the latest BCS-DBT dataset<sup>3</sup> are also chosen. For the accuracy of the experimental data, every dataset is divided equally into five parts, of which three parts are the training set, one part is the validation set, and one part is the test set. Five replicate experiments are conducted by randomizing permutations of the five data parts. The data are processed slightly differently depending on the experimental objectives. Please refer to the corresponding experimental notes for detailed processing.

#### 3.2. Main Results

In the paper, a variety of classical target detection networks are selected for training and testing on the INbreast dataset. The best-performing network is selected by comparing the common parameter metrics such as  $AP@50$  and  $AP@75$ . In the test metrics,  $AP$  represents the average accuracy rate.  $TPR$  is the True Positive Rate (i.e., Recall).  $@x$  denotes the calculation of the relevant evaluation metric under the restriction with the  $IOU > x$ .  $FLOPS$  and  $Params$  represent the computational and parametric quantities of the model, respectively. As can be seen from Table 1 (The data in the table are the mean (standard deviation)), compared to other classical networks, Faster RCNN achieves 58.04% and 31.60% in  $AP@50$  and  $AP@75$ , respectively. Thus MBMDnet is based

<sup>2</sup><http://peipa.essex.ac.uk/info/mias.html>

<sup>3</sup><https://wiki.cancerimagingarchive.net/>

**Table 4.** Results of ablation experiments on the INbreast dataset

| Net   | AP@50                  | AP@50S                 | AP@50L                 | TPR@50                 | FLOPS   | Params |
|-------|------------------------|------------------------|------------------------|------------------------|---------|--------|
| FPN   | 0.5804 (0.0730)        | 0.1470 (0.0210)        | 0.6020 (0.0540)        | 0.7850 (0.0585)        | 216.30G | 41.13M |
| HRFPN | 0.5701 (0.0929)        | 0.0840 (0.0801)        | 0.5530 (0.0290)        | 0.8006 (0.0982)        | 274.24G | 41.72M |
| PANet | 0.5974 (0.1070)        | -                      | 0.5560 (0.0390)        | 0.7964 (0.1505)        | 242.33G | 44.67M |
| MHFPN | <b>0.6462 (0.0934)</b> | <b>0.3260 (0.0740)</b> | <b>0.6345 (0.0295)</b> | <b>0.8390 (0.0915)</b> | 291.36G | 44.80M |

on Faster RCNN and achieves 64.62% in the  $AP@50$  and 83.90% in the  $TPR@50$ .

Fig 2 visualizes the inference of different networks on the INbreast dataset. In example 1, there are adjacent closer regions with larger size masses and smaller size masses. From the inference results, it can be observed that MBMDNet is more effective in detecting such targets with larger size differences. In examples 2 and 3, the detected breast mass is the only mass in the entire image and is small in size. Compared with others, the inference results of MBMDNet fit the ground truth box more closely.

To further test the generalizability of the network under different quality datasets, MBMDnet and Faster RCNN are evaluated on both MIAS dataset [21] and BCS-DBT dataset [20]. The above two datasets are treated in the same way as the INbreast dataset. Due to the early production of the MIAS dataset, the quality and annotation of the images differed significantly from the current approach. The dataset showed a large amount of blurred noise, which seriously affected the correct rate of breast mass detection. The BCS-DBT dataset is a recently proposed large dataset for breast lesions. The dataset file has more image data and less annotation data. The images are darker in general and there is interference from non-breast objects, which is more consistent with the real clinical situation.

From Table 2, it can be seen that there is a large difference in performance on different quality data sets. But compared to the Faster RCNN network, our approach still has a 6%-8% improvement in  $AP@20$ . In the  $TPR@20$  metric, our method also shows an improvement of about 3.7% without fine-tuning additional procedures.

To further compare the performance of MBMDNet, we chose the advanced detection models with the INbreast dataset of 107 images as the dataset for comparison.  $FPPI$  represents False Positive per image. As can be seen in Table 3 (The data in the table are the maximum values from five experiments), Although our method does not perform well in the  $FPPI$ , it reaches 0.96 in the  $TPR@20$  metric. In medical computer-aided systems, a missed test is more harmful to the judgment of a medical condition than a false test. And in the network training, we use Free-response Receiver Operating Characteristic Curves (FROC) to select the best TPR performance. Therefore, our method is not the best model in terms of performance on the  $FPPI$  metric.

### 3.3. Ablation Study

To test the effectiveness of each part of the MHFPN module, we performed an ablation analysis of the information flow channels during the feature fusion phase of the target detection network. Meanwhile, in order to target test the degree to the network’s equalization of focus to target boxes with

different size ratios in the feature fusion, the accuracy of different sizes of target boxes ( $AP@S$  and  $AP@L$ ) is added to the evaluation metrics of the ablation experiment. According to the format of the COCO data, the size of the breast mass is divided into large, medium and small boxes, and then the performance metrics are tested.  $AP@50S$  indicates the detection accuracy of the small box under the condition that  $IOU > 0.5$ .  $AP@50L$  is the detection accuracy of a large box under the same constraints.

According to Table 4, compared to the FPN network, both PANET and HRFPN had reduced detection accuracy for targets with a smaller ratio of x-ray breast masses. Thus, it is demonstrated that the detection of small targets in X-ray images by simply adding information flow paths or information aggregation does not perform well. Although MHFPN increases the model computation, the overall detection accuracy is improved by 6%  $AP@50$ . In particular, it improves the detection accuracy of small targets by 17.9%, thus proving that the parallel multi-head network structure can alleviate the problem of unbalanced focus of target boxes. More experimental data and visualization analysis can be obtained in the supplementary materials <sup>4</sup>.

## 4. CONCLUSIONS AND FUTURE WORKS

In the paper, we designed a general multi-head feature pyramid module (MHFPN) to address the problem of unequal focus to the target frame by multi-resolution feature maps in the feature fusion. Based on MHFPN, the detection network (MBMDNet) was designed for breast mass detection. Experiments showed that MBMDNet greatly outperformed the SOTA detection baselines on multiple breast X-ray datasets. In particular, the overall performance of the network was improved more in the detection accuracy of targets with a smaller image. However, this work can still be improved. We plan to explore semi-supervised methods [24, 25] using unlabeled data during model training, investigate the model’s application under imbalanced data conditions [26], and perform image pre-processing [27] to remove interference.

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<sup>4</sup><https://arxiv.org/abs/2302.11106>

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