Weakly Supervised Deep Learning-based Intracranial Hemorrhage Localization

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Abstract: Intracranial hemorrhage is a life-threatening disease, which requires fast medical intervention. Owing to the duration of data annotation, head CT images are usually available only with slice-level labeling. However, information about the exact position could be beneficial for a radiologist. This paper presents a fully automated weakly supervised method of precise hemorrhage localization in axial CT slices using only position-free labels. An algorithm based on multiple instance learning is introduced that generates hemorrhage likelihood maps for a given CT slice and even finds the coordinates of bleeding. Two different publicly available datasets are used to train and test the proposed method. The Dice coefficient, sensitivity and positive predictive value of 58.08\%, 54.72\% and 61.88\%, respectively, are achieved on data from the test dataset.

1 INTRODUCTION

Intracranial hemorrhage (ICH) is a relatively common life-threatening disease (25 cases per 100,000 people per year) that may develop after physical trauma or non-traumatically. The significance of this event is given by a high 30-days mortality rate (up to 52\%) and a large risk of lasting consequences among survivors (Caceres and Goldstein, 2012). For these reasons, fast diagnosis of the disease is crucial for the early initiation of treatment. If the treatment is delayed (even within minutes), the risk of permanent brain dysfunction or even death is increased. Modern algorithms, which analyse brain CT scans, can provide fast and effective support for computer-aided diagnosis and can therefore be very useful for physician’s decisions, especially in acute cases.

Nowadays, most state-of-the-art methods are focused on deep learning approaches, especially using convolutional neural networks (CNN) and their modifications or combinations. Published detection algorithms (Ker et al., 2019) and (Arbabshirani et al., 2018) use 3D CNN-based classification at the level of the whole CT scan that provide a decision about the occurrence of ICHs in the patient. A combination of 2D CNN and LSTM (Long short-term memory) algorithm for ICH detection in CT slices was designed by the authors of (Nguyen et al., 2020). Another combination of CNN and recurrent neural network was published in (Ye et al., 2019) that includes classification into its subtypes.

The algorithm for 2D ICH segmentation including its type classification based on cascade CNN model was applied by the authors of (Cho et al., 2019). For the same task, the authors of (Chang et al., 2018) suggested a hybrid 2D/3D approach using Mask Regional-CNN algorithm. The well-known U-net architecture of CNN was used in (Majumdar et al., 2018) for 2D ICH segmentation. A similar 3D approach was introduced by the authors of (Patel et al., 2019).

One of the first published mentions of the possible utilizing of attention maps for ICH detection or segmentation is in (Lee et al., 2019). Here, the authors validated the ICHs center detection to a certain degree, using simple thresholding of the attention maps. However, it was done only as a brief appendix of their manuscript. The main aim of their attention maps was to enhance the explainability of the deep learning algorithm decision. In (Ye et al., 2019), the attention maps were displayed for mere visualization of the network’s field of view.
The approach of ICH localization presented in this paper is based on a detection of local extrema in generated attention maps, which can be interpreted as maps of ICH occurrence likelihood for each pixel. These maps are produced by the proposed weakly supervised approach based on multiple instance learning (MIL) (Ilse et al., 2018). Its advantage is a position-free learning, thus precise position annotations are not needed for training, and slice-level annotations (healthy/ICH) are quite sufficient. In addition, unlike (Lee et al., 2019), we apply a special local maxima detector (Koenderink and van Doorn, 1992) to the attention map that directly leads to the precise localization of ICHs. The whole design of our approach is to our knowledge original, as most of the published papers did not deal with the precise ICH position’s localization. Besides, our approach is trained and tested using datasets different from each other.

2 METHODS

2.1 Experimental Data

The head CT data from two publicly available datasets is used in this work. More than 25,000 non-contrast brain CT scans manually labeled at the level of each slice have been released for the purposes of a challenge held by the Radiological Society of North America (RSNA) (Flanders et al., 2020). Besides, almost 500 head scans are available within the CQ500 dataset (Chilamkurthy et al., 2018) together with patient-level annotations of different pathologies, including ICHs. For the purposes of this work, CT images with soft-tissue reconstructions, kernel including ICHs or healthy patient’s scans are used (other pathologies are included only in case of co-occurrence with ICH). An extension of CQ500 annotation has been made public on the PhysioNet (Goldberger et al., 2000) in the BHX dataset (Reis et al., 2020). The authors released bounding boxes (BBs) which label the position of ICHs and are gained either manually by three radiologists for slices with the lower axial resolution or by an extrapolation. The reason of using CQ500 is that the ICHs positional labels are required for the evaluation of the proposed method. Moreover, the evaluation of the proposed approach on the dataset completely different from the training and validation set proves the ability to perform well even on new data.

2.2 Weakly Supervised Detection

The aim of this work is to build a CNN-based detector which can accurately localize ICHs in axial CT slices, using only slice-level annotations. A classification CNN model is trained to predict both the probability of ICH’s presence in the image and an attention map providing information about the likelihood of the appearance of ICHs at particular positions. Using this heat map, a detector is proposed that provides exact positional information about the location of ICHs.

The block scheme of the algorithm is shown in Fig 1. To form an input of the model, the axial slices from CT scans are taken as the first channel. To form the other two channels, the contrast enhancement is applied with respect to two radiological windows, i.e., brain (L=40, W=80) and subdural (L=50, W=130), which has been proven to be effective in previous work. Besides, all data are standardized by the mean and standard deviation of the whole dataset.

ResNet-like architecture (He et al., 2016) is chosen for the classification CNN. The feature extractor has a total of 18 convolution layers and undersamples the feature maps to a quarter of the original size. The resulting feature map is input into either the max-pooling or attention layer. Both layers result in creating outputs (for details see section 2.3) that are processed by the fully connected layer providing a binary classification (pathological or non-pathological slices). In parallel, either the activation maps ahead of global max-pooling or the weights of the attention layer are resized to the size of the original image to form the attention maps, which are further processed by the proposed ICH detector described in the section 2.4.

To obtain the desired results, the CNN classifier is trained using CT scans from the RSNA dataset and a weighted cross-entropy loss function. The data was divided in proportion 97:3 into training and validation sets to maximize the amount of training data while ensuring sufficient validation of the training process.

2.3 Max-pooling and Attention Layer

In the case of max-pooling, the forward propagation can be described by:

\[
z = \max_{k=1,...,K} (h_k)
\]

where \( z \) denotes the value of the output and \( h_k \) is the value of the feature image on the network output. In our case, global pooling over all positions \( k \) in the image is applied. By taking the maximal value from the image, the output captures the strongest feature position \( h_k \). Feature image \( h_k \) shows the prediction of ICH localization.
at individual positions, and thus, it can be used as an attention map. However, it is limited to the scalar feature map $h_k$, which limits the classification capability of the network (Ilse et al., 2018).

The MIL attention layer (Ilse et al., 2018) computes the value of the output feature vector $z$ as a weighted average of low-dimensional embeddings $h_k$:

$$z = \sum_{k=1}^{K} a_k h_k.$$  \hfill (2)

The weights $a_k$ are determined by a two one-layer fully connected neural networks applied to each position of the feature image, where outputs are element-wise multiplied and transformed with softmax to achieve that $\sum_{k=1}^{K} a_k = 1$:

$$a_k = \frac{\exp\{w^T(tanh(Vh_k^T) \odot sigm(Uh_k^T))\}}{\sum_{j=1}^{K} \exp\{w^T(tanh(Vh_j^T) \odot sigm(Uh_j^T))\}},$$  \hfill (3)

where $w \in \mathbb{R}^{L \times 1}$, $V \in \mathbb{R}^{L \times M}$ and $U \in \mathbb{R}^{L \times M}$ are parameters of the small neural network, $\odot$ is an element-wise multiplication, $\tanh()$ and $\text{sigmoid}()$ are activation functions representing hyperbolic tangent and sigmoid nonlinearity. In the case of image, the fully connected neural network can be realized by 1x1 convolution. The attention map $a_k$ represents the importance of individual positions to form the network’s decision (Ilse et al., 2018).

### 2.4 Detector

Generated attention maps mark ICHs as the areas with higher pixel values in comparison to the background. Hence, the detector finds local maxima via comparison of the original image with a grayscale-dilated (using the maximum filter) image (Koenderink and van Doorn, 1992), and returns their coordinates representing individual ICHs. To suppress irrelevant peaks with peak prominence smaller than $h$, h-maxima transform (Thiruvasagam et al., 2013) is performed. Besides, to avoid multiple and false findings, the value of local maxima must be higher than the threshold $T$ and the minimal allowed distance between peaks is $d$. The optimal detector’s parameters $h$, $T$ and $d$ were found by Bayesian optimization (for details see section 2.5).

### 2.5 Implementation Details

Network was trained using Adam optimizer (Kingma and Ba, 2014) with weight decay of $10^{-6}$, 1st and 2nd moment estimates were set to 0.9 and 0.999, respectively. Learning rate 0.01 decaying to 10 % after 20, 10 and 5 epochs was used. Network from epoch with optimal accuracy on validation set was used to generate results. Batch size 128 was used, where images were randomly cropped to size 256. Furthermore, augmentation was performed using: random mirroring, random affine transformation (max. 30° rotation, max. 10 % scaling, max. 5 % shearing), brightness addition (max. ±0.2), blurring/sharpening (max. 0.5 Gaussian sigma and max. subtraction of 0.5×Laplacian). The network contains 18 convolutional layers with 3 levels separated by pooling layers, where each level contains 3 residual blocks (He et al., 2016). The code is available at https://github.com/tomasvicar/ICH-MIL-attention-based-detector.

Parameters of the detector were determined by Bayesian optimization (Snoek et al., 2012) (implementation from (Nogueira, 2014)). The Dice coefficient (DC) was maximized using 2/5 of CT data from the CQS00 dataset. For the attention map given by max-pooling layer and attention layer, the optimal parameters are: $h = 0.024$, $T = 0.76$, $d = 10$, and $h = 0.0038$, $T = 0.024$, $d = 58$, respectively.
Figure 2: Top: Examples of ICH localization results (red ×-markers) together with annotated BB (red rectangle). Bottom: Generated attention maps; yellow color denotes to higher pixel values in contrast to blue. A – Correct localization of single subdural hemorrhage. B – Precise localization of individual subarachnoid bleedings with one FP finding. C – Multiple intraventricular and subarachnoid hemorrhages localization resulting from multiple peaks in attention map. D – Bleedings on the right are localized correctly; other bleedings are displayed in the attention map, however their maximal values are insufficient to be localized. E – Multiple bleedings act like one massive high intensity area with two local maxima in the attention map, hence only ICH in the middle is localized. F – False localization of unannotated high intensity region.

3 RESULTS AND DISCUSSION

The proposed detection method was tested on the remaining 3/5 of data from the CQ500 dataset (i.e., 254 scans, more than 55,000 slices) using slice-level BB annotations. DC together with sensitivity (Se) and positive predictive value (PPV) were chosen to evaluate the algorithm. To compute the metrics, the total number of true positive (TP), false positive (FP), and false negative (FN) results were calculated. TP is defined as an intersection of the localized position and the annotated BB (e.g., Fig. 2 C shows 2 TPs); FP is the detection outside a BB (e.g., the isolated lower finding in Fig. 2 B); FNs are the undetected BBs (e.g., Fig. 2 D shows 3 FNs). The overall results for the test dataset are shown in Tab. 1.

The classification accuracy of the CNN on the validation part of the RSNA dataset (circa 750 scans) was 89.54 % and 91.33 % for the model with max-pooling and attention layer, respectively.

Table 1: The results of ICH localization algorithm on the test dataset and comparison of both methods that were used to generate likelihood maps: max-pooling and attention layer. PPV (positive predictive value) indicates probability, that localized position is truly ICH, SE (sensitivity) reflects the percentage of correct localizations out of all ICHs.

| Method  | PPV [%] | SE [%] | Dice [%] |
|---------|---------|--------|---------|
| Pooling | 61.88   | 54.72  | 58.08   |
| Attention | 38.19   | 62.13  | 47.30   |

In this study, we proposed a weakly supervised deep-learning-based ICH localization algorithm. Having only slice-level annotations, the detector was trained to predict the precise position of ICHs in axial CT slices, which is the main advantage of the proposed method. Classification CNN was used to predict likelihood maps giving information about possible areas of bleeding. Maps highlight the regions of CNN attention to predict the classification result. Hence, likelihood maps may be considered as the interpretation of the final classification.

Precise ICH positions were found by peak detection of the likelihood map. The detector was optimized and tested on a publicly available dataset (different to training data) to test the generalization ability of the proposed approach to the new data and to make the results potentially comparable. However, considering the design of our experiment, any comparison of our approach with other authors would be out of place at present owing to its uniqueness given by the original and specific localization approach and testing.

According to the evaluation results, the attention maps given by the max-pooling layer seem more appropriate for the localization. The detector can find any type of ICHs (also in case of multiple bleedings) despite their large size, shape, and location variability. Even difficult-to-recognise bleedings are detected – e.g., subdural hemorrhage in the higher intensity region of the cerebral falx (Fig. 2 A) or small bleeding such as little intraparenchymal or subarachnoid hemorrhages (Fig. 2 B, C, D).

FN results might occur in some cases of small IPH surrounded by large edema. Both FN and FP results
sometimes originate in the detected positional coordinates closing aboard the BB. Besides, unannotated high-intensity regions subjectively similar to an ICH cause FP detections (Fig. 2 F).

Despite the possibility of false results, the detection ability denotes the high potential of the algorithm to minimize the probability of missing an ICH by oversight in clinical practice. Incorporation of the method in a computer-aided diagnostic system might warn a radiologist by highlighting the possible locations of ICHs while examining axial slices of a CT scan. Besides, the algorithm might significantly decrease the examination time as the processing of a slice takes only a few seconds. Considering the aforementioned arguments, the clinical use of the algorithm could help to prevent permanent disability or even death.

4 CONCLUSIONS

This paper demonstrates a fully automated, weakly supervised method for the localization of ICHs in axial head CT slices. The proposed algorithm is based on local extrema detection in attention maps that are obtained by a deep learning model and represent the likelihood of ICH’s presence in the given slice. The main advantage of the algorithm is the proposed MIL position-free learning method, which is used for the attention map generation. Our approach showed the ability to localize ICHs using only slice-level annotations. The Dice coefficient of 58.08 % was achieved on data from publicly available dataset.

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