

# BDCN: Semantic Embedding Self-explanatory Breast Diagnostic Capsules Network

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## Abstract

Building an interpretable AI diagnosis system for breast cancer is an important embodiment of AI assisted medicine. Traditional breast cancer diagnosis methods based on machine learning are easy to explain, but the accuracy is very low. Deep neural network greatly improves the accuracy of diagnosis, but the black box model does not provide transparency and interpretation. In this work, we propose a semantic embedding self-explanatory Breast Diagnostic Capsules Network (BDCN). This model is the first to combine the capsule network with semantic embedding for the AI diagnosis of breast tumors, using capsules to simulate semantics. We pre-trained the extraction word vector by embedding the semantic tree into the BERT and used the capsule network to improve the semantic representation of multiple heads of attention to construct the extraction feature, the capsule network was extended from the computer vision classification task to the text classification task. Simultaneously, both the back propagation principle and dynamic routing algorithm are used to realize the local interpretability of the diagnostic model. The experimental results show that this breast diagnosis model improves the model performance and has good interpretability, which is more suitable for clinical situations.

## 1 Introduction

Breast cancer is an important killer threatening women's health because of rising incidence. Early detection and diagnosis are the key to reduce the mortality rate of breast cancer and improve the quality of life of patients. Mammary gland molybdenum target report contains rich semantic information, which can directly reflect the results of breast cancer screening (CACA-CBCS, 2019), and AI-assisted diagnosis of breast cancer is an important means. Therefore, various diagnostic models were born. Mengwan (2020) used support vector machine(SVM) and Naive Bayes to classify morphological features with an accuracy of 91.11%. Wei (2009) proposed a classification method of breast cancer based on SVM, and the accuracy of the classifier experiment is 79.25%. These traditional AI diagnoses of breast tumors have limited data volume and low accuracy. Deep Neural Networks (DNN) enters into the ranks of the diagnosis of breast tumor. Wang (2019) put forward a kind of based on feature fusion with CNN deep features of breast computer-aided diagnosis methods, the accuracy is 92.3%. Zhao (2018) investigated capsule networks with dynamic routing for text classification, which proves the feasibility of text categorization. Existing models have poor predictive effect and lack of interpretation, which can not meet the clinical needs.

Based on the above pain points, we propose a semantic embedding self-explanatory Breast Diagnostic Capsules Network (BDCN), which diagnoses breast tumors based on the mammary gland molybdenum target report. Our contributions are as follows:

- Semantic segmentation algorithm is used to segment breast cancer lesions.
- Semantic tree is integrated into Bidirectional Encoder Representation from Transformers(BERT) pre-training to obtain word vectors.

- A capsule network with multi-head attention mechanism was proposed to predict breast tumors.
- Using Back Propagation to Realize Local Interpretation of BDCN Model.

## 2 Related Works

**A Self-interpretation Method based on Capsule Network:** LIME (Ribeiro et al., 2016) realizes partial interpretability of the model based on the idea of perturbation. SHAP (Lundberg and Lee, 2017) treats all features as "contributors" and produces a predicted value. These can be explained in hindsight, they were a unified interpretation method for all models. We do not consider the model independent interpretation method, through back propagation and weight sharing (Wang et al., 2020) to construct a semantic embedding self-explanatory Breast Diagnostic Capsules Network. Capsule network (Sabour, 2017) with the dynamic routing algorithm to dynamically determine rights, itself from a certain extent. It has provided the edge explanatory power to determine, which is an interpretable model based on network node association analysis.

**Fusion Embedded for Semantic Vector:** word2vec (Mikolov et al., 2013) is a representative of the embedded word paradigm, but the word that produces word2vec is static, regardless of the relationship between context and connection. BERT (Devlin et al., 2018) provides many pre-training models, which are excellent in various evaluation indexes. However, since BERT is trained in open data sets, the word vectors directly extracted by Bert are not accurate for specific fields.

## 3 Model

BDCN is based on semantic embedding, multi-head attention and capsule network to realize text diagnosis of breast cancer examination report. The overall architecture is shown in Figure 1. It is mainly composed of three modules: the semantic segmentation layer of inspection report, the semantic tree knowledge embedding extraction word vector layer (Sem-Bert), and the capsule network assisted target feature multi-head attention representation classification layer (Muti-Cap).

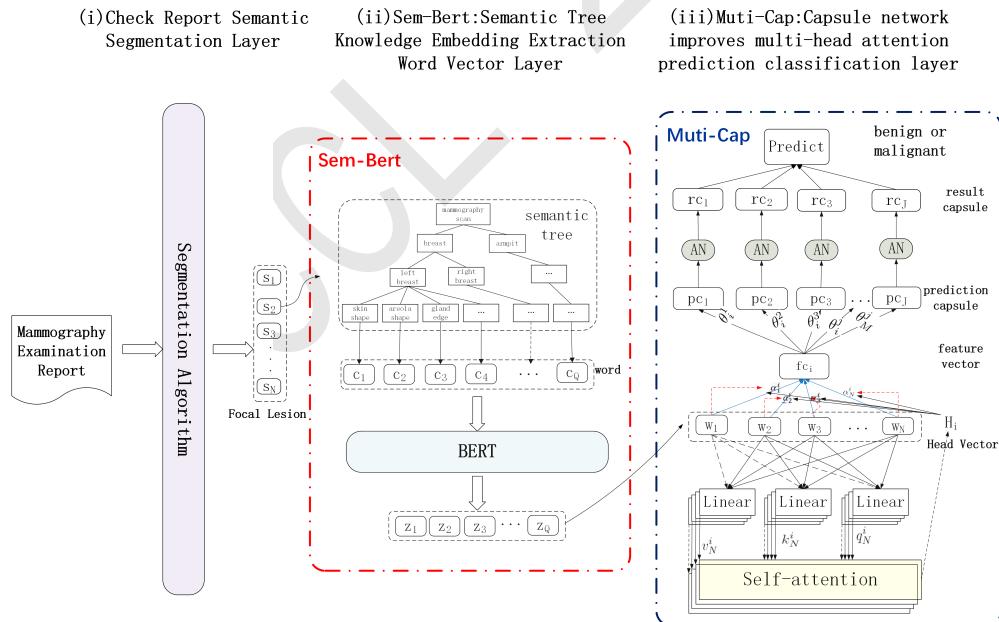


Figure 1. Overall architecture of the BDCN.

The input is an original mammography report. We use segmentation algorithm to attain preprocessing and segmentation of lesions and the implementation details are shown in Section 3.1. Then we propose Sem-Bert method, which uses semantic tree to improve BERT and pre-training to obtain the word vector for the medical field. The details are discussed in Section 3.2. The word embedding  $z_1, z_2, \dots, z_N \in R^{dz}$

is converted into the phrase embedding  $w_1, w_2, \dots, w_N \in R^{d_w}$  in one-dimensional convolution. Next, the feature capsule  $fc_1, fc_2, \dots, fc_M \in R^{d_f}$  is transformed into a prediction capsule  $pc_j, j \in [1, 3]$  based on the dynamic routing algorithm, and the result capsule  $rc_j, j \in [1, 3]$  is obtained by activating the network. Finally, the benign and malignant diagnosis of breast tumors was predicted, and the implementation details are shown in Section 3.3.

### 3.1 Check the Report Semantic Segmentation Layer

The main role of this layer is to achieve semantic segmentation of the report. The clinician can diagnose the breast tumor by analyzing the mammography report to judge the pathological condition of the patient's breast and its surrounding tissue. However, a single report may contain multiple focal lesions in the same location or even in different gland background tissues, which may lead to lower prediction accuracy if generalized. Therefore, segmentation of breast report is an important link. The core steps of the segmentation algorithm are as follows:

- Rough segmentation. Divide the report into sentences based on " " or " ", then according to different keywords such as skin, mass and axilla, they were divided into corresponding glandular background, focal lesions and axilla.
- Further subdivision. Most focal lesions mixed in the background part of the gland contain the keywords "nodules" or "densified shadow". The sentence of the background part of the gland is further subdivided to screen out the information of focal lesions and axillary parts. The sentences on the glandular background were further subdivided to screen for information on focal lesions and axillary lesions.
- Distinguish the left and right sides according to the position described. According to " " is divided into short sentences, when the "left" or "right" keyword appears, the short sentences are divided according to the position words; When "double" appears, it is divided into left and right sides.

### 3.2 Sem-Bert Layer

This layer is mainly to extract word vectors with BERT combined with semantic tree (Chen et al., 2019; Jiang and He, 2020) and the process diagram of Sem-BERT to obtain word vectors is shown in Figure 2. The Sem-Bert method constructs a semantic tree first. Semantic tree has obvious advantages of context hierarchy, so the construction of semantic tree can help to solve the problem of unreasonable word segmentation and context incoherence. According to the rule of "segment - organization description sentence - attribute description sentence", relational extraction is carried out, and dependency syntax is used to construct Extensible Markup Language(XML) semantic tree. The concrete content of report semantic tree construction includes the following five parts.

- Chinese word segmentation. Jieba word segmentation tool is the best choice to ensure the accuracy of word segmentation, and breast molybdenum target dictionary can be customized according to the knowledge of breast molybdenum target terminology combined with clinicians' guidance.
- Synonym conversion. Different doctors have different habits of describing mammograms. Replacing the words with the same general meaning in medical science into unified words can effectively reduce the redundancy of semantic tree.
- Organization description sentence acquisition. Find the organization description sentence of the corresponding part of the paragraph, scan the report from left to right, and the description before each organization word A is encountered until the next new organization word B is encountered will be classified as the description of A.
- Organization segment subtree path acquisition. Each organization descriptive sentence is converted into an attribute description sentence and the attribute value is extracted. Taking the current organization as the root node, we find finer attributes through dependency syntax and extract the attribute value of each attribute in the organization word.

- Mammary report dictionary construction and XML transformation. The information extracted from different parts was added to the branches of the semantic tree, and the duplicated information was pruned to obtain the mammary gland report dictionary.

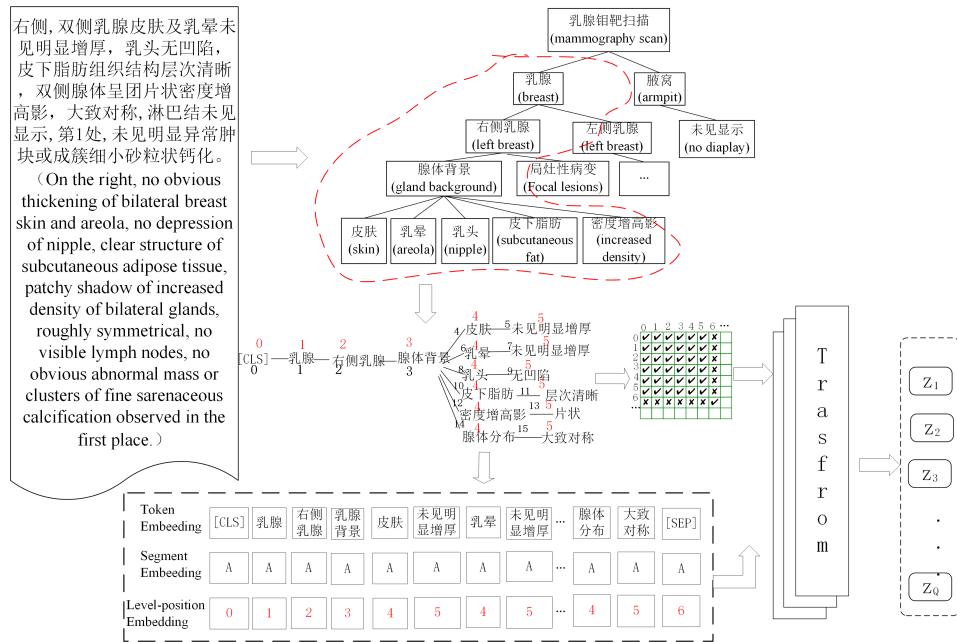


Figure 2. The structure diagram of word vector was obtained by Sem-Bert method.

Taking the semantic tree as input, by configuring the BertConfig class, setting the Tokenizer word slicer, then numbering the words in the lexicon and converting them into dictionaries, selecting the word that makes the likelihood function increase the most, and selecting the word slicer according to the frequency. The structural diagram of the Sem-Bert method is shown in Figure 2, taking the gland background description information as an example (circled by a red dotted line). Similar to Bert, the mammography report semantic tree needs to be converted into a sequence by means of token embedding, position embedding and segment embedding, while preserving its structural information. However, unlike traditional BERT, because the input is a semantic tree rather than a sequence of tokens, the positional embedding of the BERT input needs to be changed in order to preserve the structural information of the breast examination report. Position bedding is changed to level-position embedding and original-position embedding, marked by red and black numbers in Figure 2, respectively. Level-position id represents the position of the same branch in the semantic tree, and gives each Token the same branch to scale the hierarchical order information of the semantic tree starting from 0. The Original-position id is represented in the same way as the position id in BERT. [SEP] represents a special marker for multiple sentences. However, in order to preserve the structural information, a relational matrix is introduced to record whether it is reachable under the same branch, reachable under the same branch, or not. Lastly, multiple self-attentions in Transform are stacked with each other to code, and the final word vector  $z_1, z_2, \dots, z_N \in R^{d_z}$  is obtained by pre-training.

### 3.3 Muti-Cap Layer

This layer converts the word vectors of the pre-training layer into capsules, uses the capsule network to obtain the required prediction capsules, and combines effective information from multiple attention heads to achieve better classification. As one of the three most powerful semantic feature extractors, transform's self - attention mechanism is superior to CNN and neural network in word sense ambiguity resolution (Long et al., 2015). However, when the vector dimension is too high, the self-attention in each component represents different features, which leads to the fact that all attention can not fully capture the features (Tang et al., 2018). Multi-head attention can learn the features of sequences from different

aspects, which is helpful for the network to capture more abundant features. Each of these head vectors points to a feature capsule, performing attention independently. Long attention to extract the feature layer will get  $w_1, w_2, \dots, w_N \in R^{d_w}$  as its input in the layer. By way of generating M features capsule  $fc_1, fc_2, \dots, fc_M \in R^{d_f}$ , we need to have M long since attention vector  $h_1, h_2, \dots, h_M \in R^{d_h}$ , and for each  $H_i \in R^{d_f}, m \in [1, M]$  can obtain different key vectors  $k_n^i$ , value vectors  $v_n^i$  and query vectors  $q_n^i$  through linear transformation, and then get more attention by the vector series. The generation process of the head vector is as follows:

for  $n = 1, 2, 3, \dots, N$  do

$$v_n^i = W_i^V w_n \quad (1)$$

$$k_n^i = W_i^K w_n \quad (2)$$

$$q_n^i = W_i^Q w_n \quad (3)$$

$$H_i = \text{attention}(q_n^i, k_n^i, v_n^i) \quad (4)$$

where  $W_i^V \in R^{d_f \times d_w}, W_i^K \in R^{d_h \times d_w}$  are parameters. The weight of attention and the output feature capsule of multi-head attention can be calculated. The formulas are as following:

$$\alpha_n^i = \frac{\exp(\frac{H_i^T k_n^i}{\sqrt{d_p}})}{\sum_{n'} \exp(\frac{H_i^T k_{n'}^i}{\sqrt{d_p}})} \quad (5)$$

$$fc_i = \sum_n \alpha_n^i v_n^i \quad (6)$$

The feature capsule is obtained by the sum of attention weights. As a parameter, the number of feature capsules is adjustable, which solves the problem that the intermediate parameters of three prediction capsules and M feature capsules are too large, and we need to normalize the attention weight:

$$\sum_n \alpha_n^i = 1 \quad (7)$$

With CNN classification, some important information will be lost in the operation of the aggregation layer, while capsule network represents a group of neurons by capsule, replacing the neuron output vector (Sabour, 2017). Therefore, the capsule network can effectively represent the location and semantics of features, and each upper capsule is the high-level semantics of the lower capsule. In addition, it can improve the information aggregation of multiple attention, so as to obtain more effective features and improve the ability of text representation. The structural schematic diagram is shown in Figure 3.

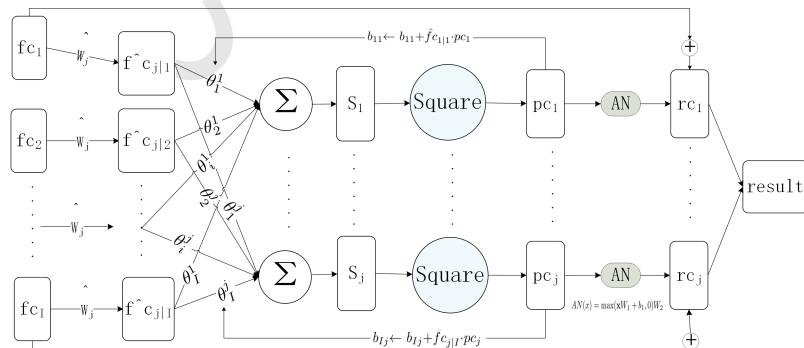


Figure 3. Structure diagram of improved capsule network.

The dynamic routing algorithm of the capsule network (Zhao et al., 2019) is used to calculate the prediction capsule  $pc_i$ . Through multiple iterations of the routing process, Muti-Cap can determine the number of characteristic capsules flowing into the prediction capsules, which plays an significant role in the prediction capsules. Firstly, the feature capsule vector is calculated by inputting the feature capsule

$pc_i$  and multiplying the learning transformation matrix  $\hat{W}_j$ . The dynamic routing weight  $\theta_i^j$  is then determined by calculating the "routing softmax" of the initial logits  $b_{ij}$ , the formulas are as following:

$$\hat{f}c_{j|i} = \hat{W}_j f c_i + \hat{b}_{ij} \quad (8)$$

$$\theta_i^j = \frac{\exp(b_{ij})}{\sum_j \exp(b_{ij})} \quad (9)$$

where  $\hat{W}_j$  here is shared among each feature capsule to obtain the feature capsule vector, and  $b_{ij}$  is initialized to 0.

The final output is normalized activation by the function of squash (Zhao et al., 2018) to make the whole model nonlinear and get the predicted capsule, as shown in Equation 10 and 11.

$$s_j = \sum_i \theta_i^j \hat{f}c_{j|i} \quad (10)$$

$$Squash(s_j) = \frac{\|s_j\|^2}{1 + \|s_j\|^2} \frac{s_j}{\|s_j\|} \quad (11)$$

where  $s_j$  is the original output capsule vector, which is the predicted capsule value before the square function is activated.

We introduce the activation network based on the dynamic routing algorithm in the traditional capsule network. The obtained prediction capsule was input into the activation network of linear transformation and ReLU activation function, and the final output was obtained by connecting with the residual  $f c_j$  feature capsule. The formula is shown in 12 and 13.

$$AN(x) = \max(\mathbf{x}W_+c_1, 0)W_2 + c_2 \quad (12)$$

$$predict = f c_j + AN(pc_i) \quad (13)$$

In order to match the attention weight of multi head attention, we need to softmax the dynamic routing weight.

### 3.4 Model Interpretability

We use the principle of back propagation (Wang et al., 2020) to reach local interpretability. That is, a sample of mammography report to explain why the report was diagnosed as benign or malignant breast cancer. The model defines two interpretable parameters: attention weight  $\alpha_i^j$  and dynamic routing weight  $\theta_i^j$ . Attention weight indicates whether attention right perform an important function in the formation of feature capsule. The calculation method is shown in Equation 5 above. The dynamic routing weight determines the higher level classification capsule to which the current feature capsule will output it. The formula is described in the Equation 9. As a classifier, the weight matrix of the traditional full-connection layer is fixed after training (Zhang et al., 2019), which is not conducive to interpretation. However, in our model, all layers are fully connected. The capsule network part and the multi attention part are interdependent, which has an important impact on the research of local interpretation based on back propagation. As shown in the figure 4, prediction capsules are divided into three categories  $j \in [1, 3]$ : benign, suspected malignant, and malignant breast cancer. By setting the parameter  $P1$  of the capsule classification layer and the parameter  $P2$  of the feature layer extracted by the multi-head self-attention.

$P1$  The largest routing weight was selected from  $\theta_1^j, \theta_2^j, \dots, \theta_M^j$ , and the largest  $\theta_i^j$  indicated that the  $i$ th feature capsule made the largest contribution to the  $j$ th prediction capsule, indicating that  $P1$  the largest routing weight had an important impact on the prediction of benign breast cancer. Meanwhile, based on the principle of backpropagation, for  $P1$  corresponding feature capsules,  $P2$  phrase embedding  $w_i$  which contributed the most to the largest feature capsule was found in the multi-head self-attention according to the weight of attention  $\alpha_1^j, \alpha_2^j, \dots, \alpha_N^j$ . Then the keywords are searched in the breast examination report After one-dimensional convolution layer. When words are colored, it is important to predict the results. The model can be visualized in the report to provide simple and clear help for doctors.

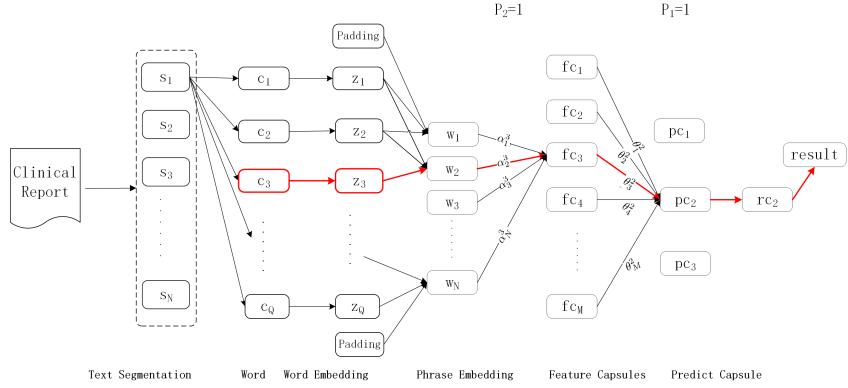


Figure 4. Model interpretability process display diagram.

## 4 Experiment

In this section, we mainly discuss the validity and interpretability of our model. Firstly, the selection of methods in different stages is considered, such as the performance differences of Sem-Bert, BERT and word2vec in acquiring word vectors. Secondly, the model in this paper is compared with some unexplainable text classification models such as TextCNN (Rakhlin, 2016) and LSTM (Kalchbrenner et al., 2015). Finally, we demonstrate the interpretability of our model through experiments.

### 4.1 Dataset

This dataset is the molybdenum mammography examination report of Shanghai Ruijin Hospital. We selected 1600 preoperative mammography data from 34 million original reports and included 2857 data after segmented pretreatment. The small sample dataset is classified into three categories, including benign, suspected malignant and malignant. The analysis of the data set is shown in Table 1.

Dataset	Classes	Benign	Suspected of Malignant	Malignant
Training Set	3	1344	390	268
Test Set	3	734	58	63
Mammography	3	2078	448	331

Table 1. Summary of mammography report dataset

### 4.2 Parameter Settings

The model was implemented by Tensorflow and trained by Adam Optimizer (Bock and Weiβ , 2019). The multi-head attention part activation function uses a ReLU nonlinear function, the capsule network layer activation function Squash function. We set  $P1 = P2 = 3$ , the number of class capsules is 3, the number of head vectors is 18.

### 4.3 Different stage selection

**Evaluation indexes:** The main evaluation indexes of the experiment are Micro-Precision(Mi-P), Micro-Recall(Mi-R), Micro-F1-score(Mi-F1), Macro-Precision(Ma-P), Macro-Recall(Ma-R), Macro-F1-score(Ma-F1), Receiver Operating Characteristic(ROC).

**Comparative experiment:** The main purpose of this section is to verify the validity of the 'Sem-Bert + Muti-Cap' model, including the advantages of Sem-Bert in obtaining word vectors and the classification accuracy of Muti-cap. In this experiment, six other models are selected as the baseline. Comparing experiment 'word2vec+Muti-Cap' and 'BERT+Muti-Cap' in table 2 mainly verifies whether texts in specific fields such as medical care will affect the acquisition of word vectors, thus affecting the prediction

performance. The comparative experiments textCNN, textRNN, LSTM, and Capsule in table 3 mainly verify the influence of different classification models on the prediction performance.

**Experimental Results:** Figure 5 shows the ROC curves of the model in this paper obtained according to different evaluation criteria under three classifications. For three classifications, the area under the curve and AUC values of micro and macro methods are different. In this paper, the dataset of benign, malignant and suspected malignant are large difference between three kinds of sample size, and there are obvious characteristics of malignant samples. So the ROC curve of class 2 malignant tag is more left, and the AUC area is larger. Meanwhile, the average (micro) AUC is larger than that of macro AUC. Since the samples in our dataset are unbalanced, so to give equal attention to the categories with small samples and those with large sample data. The ROC curves in subsequent comparative experiments were obtained by macro method.

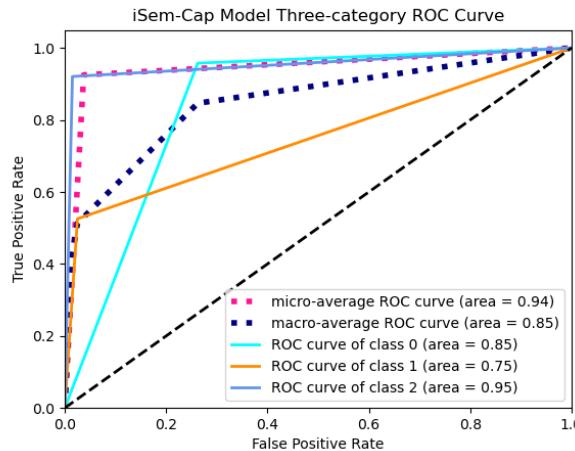


Figure 5. ROC curve of three classifications of BDCN model

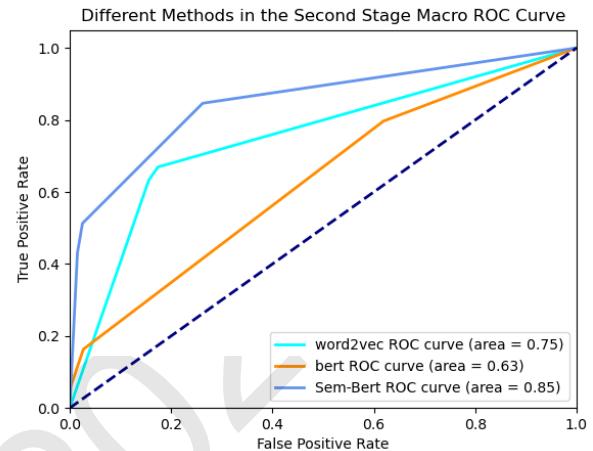


Figure 6. In the second stage, ROC curves of word vectors are obtained by different methods

Comparing (1) (2) with (3) in Table 2 and Figure 6, we can basically draw a conclusion: under the premise of using Muti-Cap method in the third stage, the evaluation index of the word vector method obtained by Sem-Bert in the second stage is higher, the ROC curve covers the other two comparative experiments, and the AUC value of the area under the ROC is significantly larger than the other two. Although there may be a loss of text information in the process of constructing semantic tree, the experiment effectively proves that the method of acquiring word vectors through semantic tree combined with BERT is more accurate than traditional word2vec and BERT method directly, which is very meaningful.

Model	Evaluation index					
	Mi-P(%)	Mi-R(%)	Mi-F1(%)	Ma-P(%)	Ma-R(%)	Ma-F1(%)
(1)word2vec+Muti-Cap	83.25	83.25	83.25	44.02	61.08	47.83
(2)BERT+Muti-Cap	87.37	87.37	87.37	78.26	42.37	45.56
(3)BDCN(our)	<b>91.58</b>	<b>91.58</b>	<b>91.58</b>	<b>75.95</b>	<b>79.73</b>	<b>77.14</b>

Table 2. Model selection at second stage

Table 3 compares the current mainstream deep learning models of text classification: CNN family textCNN, RNN family textRNN, LSTM family LSTM and capsule of Muti-Cap. The baselines are word2vec method to obtain the word vector, whereas in BDCN model, term vectors are obtained by Sem-Bert method. Comparing experiments (1), (2),(3) and (4) in table 5, the capsule network from the view on the text classification task to task, effect and less textCNN this classic mode, but better results than simple LSTM, which shows that the attempt is meaningful. Moreover, through the improvement of the

traditional capsule network and word vector acquisition method, our model has favorable performance results in each index. Experiments show that the selection of the second stage of this model has greatly improved the accuracy of the model.

Model	Evaluation index					
	Mi-P(%)	Mi-R(%)	Mi-F1(%)	Ma-P(%)	Ma-R(%)	Ma-F1(%)
(1)textCNN	88.79	88.79	88.79	73.94	78.43	75.15
(2)textRNN	74.24	74.24	74.24	70.59	71.78	72.74
(3)LSTM	69.93	69.93	69.93	66.69	64.05	69.12
(4)Capsule	77.56	77.56	77.56	67.57	71.57	69.37
(5)BDCN(our)	<b>91.58</b>	<b>91.58</b>	<b>91.58</b>	<b>75.95</b>	<b>79.73</b>	<b>77.14</b>

Table 3. Comparison to multimodal baselines

#### 4.4 Interpretability

Based on the back propagation principle, the BDCN can be interpreted to find the top three important weights in the dynamic routing weights according to the prediction results. Then the three important weights in the attention weights can be deduced in reverse, so as to find the three important features of judging the benign and malignant in this sample.

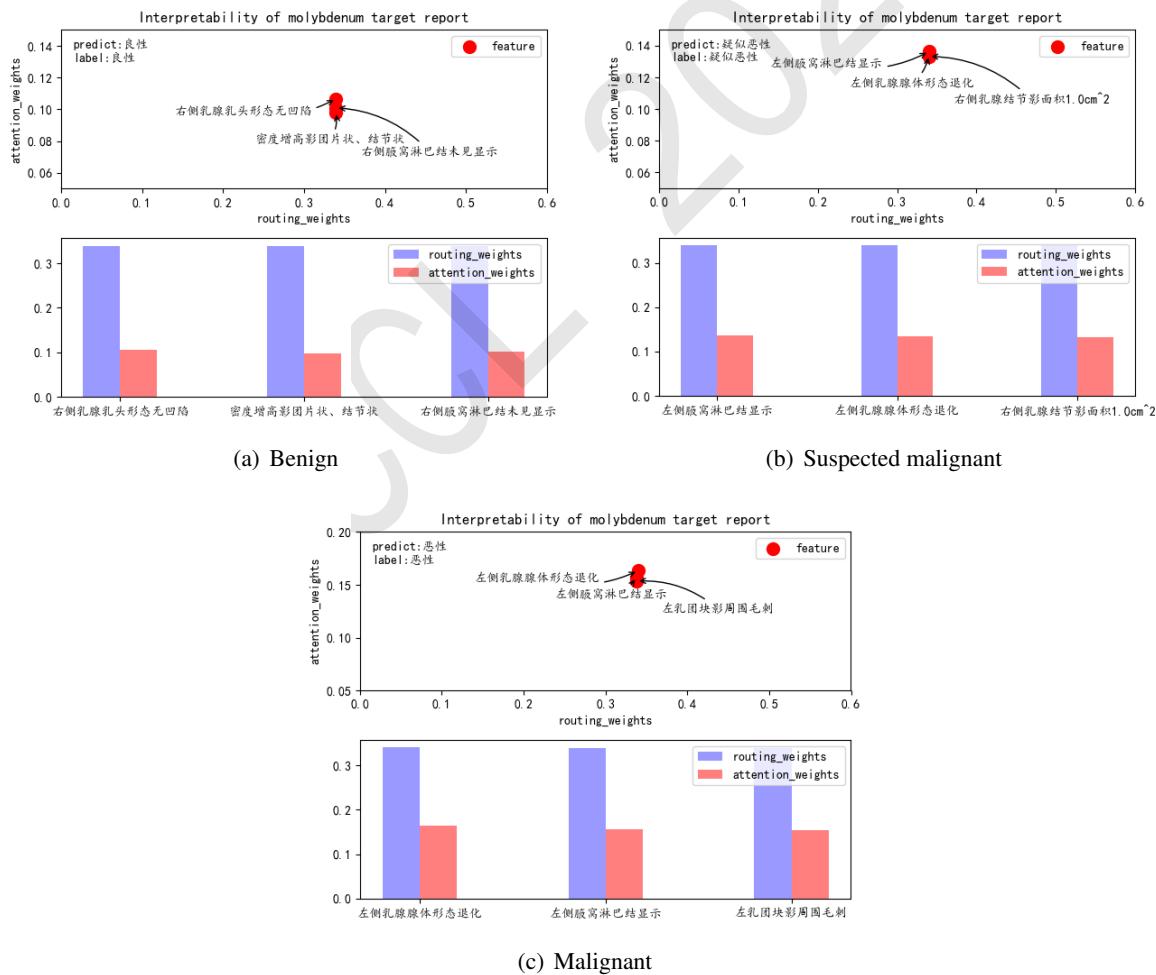


Figure 7. Analysis of benign and malignant characteristics of breast molybdenum target

We plot a scatter plot and a bar chart, as shown in Figure 7. They respectively explained the important features of benign, suspected malignant and malignant cases as well as the attention weight and routing weight. For example, according to the ranking of importance, the three important features that appear for the first time of each sample were selected. In Figure (c), it can be seen that 'the shape of the burr', 'glandular degeneration' and 'lymph node display' are all important features of malignant tumors. In Figure (a) 'roughly symmetrical glands', 'density increase', 'no depression of the nipple' and 'no lymph nodes displayed' are all important indicators of benign breast tumors. Suspected malignancy is a grading result of BI-RADS4, the more similar the characteristics are to malignant, the greater the probability of malignant tumors, and it is convenient for the doctor to remind the patient for further examination.

More intuitively show the local interpretability of the model, the extracted important features were labeled and displayed on the segmented breast molybdenum target samples. As shown in Figure 8 (a) 'no inverted nipples', 'no lymph nodes' and 'increased density'. Benign and malignant breast tumors can be determined by directly observing the words marked yellow, and the prediction accuracy of the BDCN is also verified from the side. On the basis of concise and accurate text report, we add the annotation of important words to make up for the lack of interpretation of text classification better.

**(a) Benign**

```

predict: 良性
label: 良性
00227613420151030090427双侧乳腺皮肤及乳晕未见明显增厚, 乳头无凹陷, 皮下脂肪组织结构层次清晰, 双侧腺体较丰富, 呈团片状、结节状密度增高影, 边缘膨隆, 大致对称; 淋巴结未见显示。右侧第2处: 右乳钙化灶。

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**(b) Suspected malignant**

```

predict: 疑似恶性
label: 疑似恶性
10035575420110218090601左乳下部皮肤略内凹, 余双侧乳腺皮肤及乳晕未见明显增厚, 乳头无凹陷, 皮下脂肪组织结构层次清晰, 双侧腺体部分退化, 呈条索状改变, 大致对称; 淋巴结显示。左侧第1处: 左乳内下象限深部偏高密度小结节, 直径约1cm, 边缘浅分叶, 局部似见浸润毛刺。

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**(c) Malignant**

```

predict: 恶性
label: 恶性
00154885620110317092257左乳外侧皮肤增厚凹陷, 乳头无凹陷, 皮下脂肪组织结构层次清晰, 双侧腺体部分退化, 呈条索状改变, 大致对称; 淋巴结显示。左侧第1处: 左乳外上可见团块影, 周围可见毛刺, 直径约为2.0cm, 周围浸润性改变。

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Figure 8. Analysis of benign and malignant characteristics of breast molybdenum target

## 5 Conclusion

We proposed a semantically embedded self-interpreted breast diagnostic capsule network model. Semantic segmentation algorithm was used to segment the report, Sem-Bert method was used to obtain word vectors in medical field with hierarchical relationship, and capsule network with multiple attention was used to achieve prediction and classification of breast tumors. The validity of our model is better than other models in breast molybdenum target dataset. In addition, local self-interpretation method was used to provide intelligibility analysis, which was in line with doctors' clinical expectations. In the future, we will further study the global interpretability of the model and we hope to apply our technology to other diseases.

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