MULTI-MODALITY BREAST MRI SEGMENTATION USING NNU-NET FOR PREOPERATIVE PLANNING OF ROBOTIC SURGERY NAVIGATION

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ABSTRACT
Segmentation of the chest region and breast tissues is essential for surgery planning and navigation. This paper proposes the foundation for preoperative segmentation based on two cascaded architectures of deep neural networks (DNN) based on the state-of-the-art nnU-Net. Additionally, this study introduces a polyvinyl alcohol cryogel (PVA-C) breast phantom based on the segmentation of the DNN automated approach, enabling the experiments of navigation systems for robotic breast surgery. Multi-modality breast MRI datasets of T2W and STIR images were acquired from 10 patients. Segmentation evaluation utilized the Dice Similarity Coefficient (DSC), segmentation accuracy, sensitivity, and specificity. First, a single class labeling was used to segment the breast region. Then it was employed as an input for three-class labeling to segment fat, fibroglandular (FGT) tissues, and tumorous lesions. The first architecture has a 0.95 DSC, while the second has a 0.95, 0.83, and 0.41 for fat, FGT, and tumor classes, respectively.

Keywords: MRI segmentation, convolutional neural networks, breast surgery, surgery planning,
1 INTRODUCTION

Breast cancer is the most common cancer among women, except skin cancer (Torre et al. 2015). While the primary treatment is breast-conserving surgery (BCS), breast cancers can be spotted on a screening X-ray mammogram (XRM) as a nonpalpable abnormality, enabling less invasive local treatment. However, tumor localization is necessary to guide surgical excision (Frank, Hall, and Steer 1976); existing approaches have limited visualization in the procedure; a radiologist inserts a wire marker into the tumor prior to the surgical excision, which is determined by 2D guidance founded on a projection XRM (St John et al. 2017). Subsequently, the surgeon uses the guidewire to localize the tumor through XRM annotations by the radiologist and possibly aided by 2D ultrasound (US). Hence, the wire marker usually offers poor localization of the resection area, particularly for nonpalpable tumors, frequently leading to inaccurate localization and imprecise excision, which contributes to sizeable healthy tissue resections, tumor spillage, and spreading of the resection border (Torre et al. 2015).

Additionally, navigation toward complex 3D lesions visualization is frequently accomplished with 2D freehand US (FUS) (Krücker et al. 2011). Yet, localization cannot be adequately visualized due to the poor image contrast of the US. Although highly sensitive imaging modalities (such as MRI) can be used in a preoperative procedure to obtain precise positions of the lesions, finding a method to update and align these lesions’ positions from real-time US images can benefit current surgical procedures. Therefore, many research and commercial platforms have applied image fusion techniques that align preoperative and intraoperative images, utilizing rigid or affine registration techniques (Guo et al. 2018). Hence, the breast is hyperelastic anatomy; applying compression forces by the US probe would entail a considerable nonlinear deformation challenge. Therefore, an applicable alignment procedure of the preoperative and intraoperative US images is vital for an actual probe-tissue coupling and sufficient image quality. However, a proper soft tissue modeling of breast elasticity in real-time has not been solved.

This paper describes a deep learning (DL) approach to multi-modality breast MRI segmentation that can be used as the groundwork for preoperative imaging in a future combination with intraoperative imaging in a surgery navigation system. Consequently, magnetic resonance imaging (MRI) of the breast is beneficial for diagnosing and screening breast cancer (Sinha and Sinha 2009). Furthermore, due to its high soft-tissue contrast, MRI can detect the discrimination between different structures in the breast and enable 3D visualization (Giess et al. 2014). However, breast MRI images consist of other organs such as the lung, heart, and pectoral muscles. As a result, it is crucial to segment the breast region from the other organs.

Moreover, segmentation is essential in many clinical applications (Aerts et al. 2014, Nestle et al. 2005), such as interventional treatments and intra-operative surgery navigation systems (Bernard et al. 2018). However, MRI manual segmentation is time-demanding and user variability-dependent (Granzier et al. 2020). Therefore, numerous techniques have been developed to help radiologists diagnose, detect, and enhance the analysis efficacy of breast lesions (Pang et al. 2015). Semi-automated methods (Chen, Giger, and Bick 2006) need less time than manual methods, but they still require user involvement, yielding varying results depending on different users. Yet, fully automated segmentation of breast tissue and lesions continues to be a challenge even by using computer-aided diagnosis (CAD) systems (Gubern-Mérida et al. 2015).

Various traditional algorithms have been utilized to address this problem. For example, Fuzzy c-mean (FCM) has been used for lesion detection, where voxels are assigned into classes based on their distance in the feature space (Bezdek 1981, Chen, Giger, and Bick 2006). Wu et al. (2013) developed an edge technique in the 3D sagittal T1-weighted MRI scans to segment the breast region out from other organs and achieved a DSC of 0.95. In recent years, deep learning techniques have been largely used in medical image segmentation tasks, for instance, Fully Convolutional Network FCN (Shelhamer, Long, and Darrell 2017), SegNet (Badrinarayanan, Kendall, and Cipolla 2017), U-Net (Ronneberger, Fischer, and Brox 2015), and V-Net (Milletari, Navab, and Ahmadi 2016). These DNNs can extract detailed features and
accomplish end-to-end segmentation. Zhang et al. (2019) developed a DL approach using U-Net for breast region and FGT segmentation, achieving a DSC of 0.86 and 0.83, respectively.

However, most traditional and DL segmentation models require modification for specific datasets. Thus, the DNN parameters are often fine-tuned for particular MRI scanner characteristics and protocols. Therefore, different modalities and scanning protocols yield different breast MRI scans. Although existing approaches have shown an acceptable performance on specific task optimization problems, they may not handle the variability of MRI data.

Therefore, we propose a fully automated approach by applying two consecutive nnU-Net architectures for the breast region, inner breast tissues, and tumor masses segmentation by using multi-modality breast tumor images. This will cope with the variability of MRI data and avoid manual intervention. nnU-Net has appeared as a state-of-the-art biomedical segmentation architecture (Isensee et al. 2021). It is a self-adopted network architecture method for every particular image dataset. Without manual tuning or user intervention, nnU-Net configures all the segmentation task stages, resulting in performance optimization for every dataset. This architecture exhibited an outstanding performance for the task-specialized DL pipelines of 33 international public segmentation competitions (Isensee et al. 2021). Likewise, it has been widely applied to many segmentation competitions, but it has not been applied to multi-modality breast cancer MRI datasets yet. To the best of our knowledge, this study is the first to explore and test nnU-Net for breast region and 3-class breast tissue segmentation for fat, FGT, and tumor mass in a routine MRI dataset that consists of 10 T2-weighted and STIR fat-suppressed multi-modality images gained from an open-access breast MRI database.

Furthermore, we are also presenting a patient-specific breast-mimicking phantom based on that automated segmentation approach. It will allow the experiments of developing and validating a calibration-assisted preoperative MRI to intraoperative 3D US registration technique utilizing an open-source PLUS toolkit (Lasso et al. 2014). This registration will integrate later with the optical tracking of a breast surgery robot. Validation will use surgical fiducials markers inserted in a breast phantom.

2 MATERIALS AND METHODS

2.1 Datasets

The datasets of this study contain 10 patients with a median age of 40 years; and a range of 40-70 years (Bloch, 2015). All subjects are acquired from an open-source cancer imaging archive (www.cancerimagingarchive.net) under the breast-diagnosis study category (Bloch, 2015). Our selection of each subject in the dataset contains two MR modalities: T2-weighted and STIR fat-suppressed images. Each breast image volume consists of 82 to 95 axial slices with a slice thickness of 2 mm. The variants of pixel resolution and image sizes are noted on the selected dataset. As a result, we resample to the pixel resolution of 0.65 mm along x and y, and the image size of 512 × 512. Patients are imaged in the prone position by a 1.5 T Philips MRI scanner coil while their breasts are hanging in the two holes. Thus, we randomly divide the dataset to 80:20, yielding 8 image volumes for training/validation and 2 image volumes for testing. Therefore, a trained biomedical engineer has manually produced the ground truth (GT) segmentation to generate a reference for breast region, fat, FGT, and tumor since there is insufficient data analysis in online repositories. This manual segmentation is revised and validated by a radiologist from Eastern Virginia Medical School (EVMS) in Norfolk, VA. The manual segmentation on 3D Slicer 4.11 (Fedorov et al. 2012) platform is used for this purpose.

2.2 Segmentation based on nnU-Net

Our pipeline network is developed based on nnU-Net, two consecutive nnU-Net architectures with multimodal inputs of STIR fat-suppressed MRI and T2-weighted modalities. Our consecutive nnU-Net approach is first to segment the whole breast region and then use it as mask input for the second network to perform three-class segmentation of fat tissue, FGT tissue, and tumor mass within the breast region.
Figure 1 shows the proposed segmentation framework where the left rectangle represents the breast region segmentation, and the right rectangle represents the inner breast tissues segmentation.

The nnU-Net architecture is adapted from 3D U-Net (Ronneberger, Fischer, and Brox 2015, Isensee et al. 2021, Çiçek et al. 2016). It comprises a contracting network, encoder, followed by an expansive network, decoder, which constructs a U-shaped architecture.

The encoder network, corresponding to the downslope of the U, embeds the repetitive implementation of a convolution, a Leaky Rectified Linear Unit (Leaky ReLU), and a max-pooling process, where the feature-based representation is increased, and the spatial information is reduced. Therefore, the decoder network, which indicates the upslope, combines spatial and feature data by deconvolutions series for high-resolution features from the encoder over successive layers (Ronneberger, Fischer, and Brox 2015). As a result, the skip connection between the corresponding encoder and decoder layers maintains the precise feature information crucial for the up-sampled output image. Thus, at the final layer of the decoder network, a convolution of a $1 \times 1 \times 1$ kernel conducts an up-sampling so that the segmentation voxel results correlate to the voxel input image (Isensee et al. 2021). Figure 2 shows the network parameters and their datasets. Hence, nnU-Net combines dice loss and cross-entropy loss functions to train one class label and three class labels for the first and second network, respectively, instead of a regular cross-entropy loss function in the original U-Net architecture. As a result, the segmentation accuracy and training stability are improved (Isensee et al. 2021).

Additionally, eight operations of data augmentation are employed by nnU-Net to handle the limitation of training data, such as rotation scaling, mirroring, Gaussian blur, and Gaussian noise (Isensee et al. 2021). It is noteworthy that nnU-Net implants some modifications to the U-Net architecture baseline (Ronneberger, Fischer, and Brox 2015, Isensee et al. 2021), specifically: (1) convolution padding to maintain the exact image size for inputs and outputs; (2) utilizing of instance normalization (IN) as a
replacement of batch normalization; (3) as a substitute of ReLU, Leaky ReLU is employed to address the dying neuron matter. Hence, nnU-Net is a self-adoptable algorithm where normalization, resampling, and cropping are performed to the dataset parameters, such as resolution and slice thickness, in the nnU-Net preprocessing step (Isensee et al. 2021).

The nnU-Net model employs the stochastic gradient descent with an initial learning rate of 0.01 and Nesterov Momentum (0.9) to optimize the loss function (Isensee et al. 2021). The patch sizes of the two nnU-Net networks for the breast region and breast tissue segmentations are $40 \times 192 \times 256$ and $48 \times 160 \times 256$, respectively. In this study, the minimum feature map size is at $4 \times 4 \times 4$, the maximum number of feature map is at 320, and the minimum batch size is at 2. Therefore, the down-sampling is 6 layers. Each epoch contains 250 batches, and training runs for 1000 epochs. We apply 5-fold cross-validation (CV) for training and validation to exploit the manual GT segmentations we analyze and obtain from the breast MRI dataset's public repositories. The training process of the segmentation model takes place in Old Dominion University’s (ODU) High-Performance Computing cluster. A virtual environment is created on python 3.8.5 in the cluster using PyTorch 1.6.0 (Paszke et al. 2019) as a framework. Additionally, Batchgenerators 0.21 (Isensee 2020) and other necessary python libraries are installed in the virtual environment. Finally, the nnU-Net code is available for public access at https://github.com/MIC-DKFZ/nnUNet.

2.3 Evaluation

We evaluated the performance of the segmentation network of consecutive architecture with standard statistical metrics of segmentation, DSC, segmentation accuracy, segmentation sensitivity, and segmentation specificity (Tharwat 2021, Popovic et al. 2007).
PVA powder is combined in an Erlenmeyer flask. Hence, we write down the flask weight and its contents. Then, we use a magnetic stir plate to blend the mixture for 30 minutes. Then, the solution is taken to a 95°C temperature bath for 2 h. to break down any masses. After that, we lightly stir PVA for 1 hour to promote dissolution and solution homogeneity and gradually cooled down to room temperature. Lastly, the flask is weighed one last time to restore any weight loss by deionized water, so the solution weight remains 10 wt. % PVA.

In order to make an anthropomorphic elastic phantom, the PVA liquid must be poured into the proper molds. Hence, these molds are 3D printed from surfaces extracted from our segmentation results. Therefore, we make molds for tissue-mimicking fat, FGT, and tumors. Accordingly, we use open-source 3D Slicer 4.11 (Fedorov et al. 2012) to extract surfaces from segmented tissue.

30 ml of liquid PVA are distinguished with red enamel paint (Testor's 1105tt, red metallic paint) and 320 ml with yellow paint (Testor's 1115TT, amber metallic paint). We then dispense the red-colored liquid PVA into its mold and let it rest for 12 hours, allowing air bubbles to rise and dissipate. We keep it well-sealed at room temperature during this time. Soon later, the freezing phase begins in a standard chest freezer, starting from room temperature till gradually reaching -20°C over 12 h. Accordingly, the thawing phase is initiated by turning off the freezer, and gradually the temperature goes to room temperature over 12 hours. In this way, one FTC is finished. Then, a second FTC is initiated for the tumor-mimicking volume. The yellow-colored liquid PVA is poured into its mold for its first FTC.

Later, the tumor and FGT components are separated from their molds and placed inside the fat-mimicking mold to complete the making of the breast phantom. Therefore, we sew a nylon thread over the FGT and tumor mimicking volumes so that they are hanging in the fat mimicking volumes so that they are hanging in the fat mimicking volume mold to make certain of the distance and spatial coherence of the three components. Next, the fat-mimicking volume container is loaded with PVA liquid. Finally, we place it in the freezer for a last FTC. Once it is finished, we cut out the threads. Thus, the phantom is separated from its mold as we have a breast-mimicking phantom that contains 1 FTC for Fat-mimicking tissue, 2 FTCs for the FGT component, and 3 FTCs for the tumor. Therefore, we can store it in deionized water at 5°C (Surry et al. 2004).

\[
\begin{align*}
    \text{DSC} &= \frac{2\times TP}{2\times TP+FP+FN} \\
    \text{Jaccard} &= \frac{TP}{TP+FP+FN} \\
    \text{Sensitivity} &= \frac{TP}{TP+FN} \\
    \text{Specificity} &= \frac{TN}{TN+FP}
\end{align*}
\]

TP, TN, FP, and FN are true positive, true negative, false positive, and false negative, respectively.

### 2.4 Segmentation-Guided Elastic Breast Phantom Application

We build a breast phantom based on the DL automated breast segmentation approach for our robotic surgery application navigation experiments. We have created Polyvinyl Alcohol Cryogel (PVA-C) breast phantoms started on our segmentation results. We opt for PVA-C because it shows elastic fidelity, mimicking soft tissue deformations and medical imaging properties (Surry et al. 2004). PVA-C is a blended mixture of deionized water and PVA powder first heated to stimulate a total dissolution. Then it becomes an elastic solid by frozen and thawed cycles one or more times. Additionally, we can control imaging and elastic properties by controlling the PVA concentrations and the number of freeze-thaw cycles (Surry et al. 2004). It is worth noting that 3D printing is a crucial component of the PVA-C molds; the planning and navigation experiments need a highly compatible soft tissue phantom, which disqualifies most 3D-printed options in our case.

Polyvinyl alcohol (PVA) powder with an average molecular weight (MW) of 130,000 and the hydrolysis of over 99% (Sigma-Aldrich, SKU 563900) is utilized for the preparation of aqueous liquids. Our method and approach are generated from the published methods of (Surry et al. 2004) and (Kharine et al. 2003). Therefore, a 90 wt. % deionized water and 10 wt. % PVA powder is combined in an Erlenmeyer flask. Hence, we write down the flask weight and its contents. Then, we use a magnetic stir plate to blend the mixture for 30 minutes. Then, the solution is taken to a 95°C temperature bath for 2 h. to break down any masses. After that, we lightly stir PVA for 1 hour to promote dissolution and solution homogeneity and gradually cooled down to room temperature. Lastly, the flask is weighed one last time to restore any weight loss by deionized water, so the solution weight remains 10 wt. % PVA.
3 RESULTS

We used the segmentation evaluation metrics described in section 2.3 to examine our pipeline method. Table 1 details all results as mean ± SD. Hence, the highest model performance out of 5-fold CV corresponds to the reported results in Table 1. The DSC scores for breast region, fat, FGT, and tumor segmentation are 0.95 ± 0.07, 0.95 ± 0.00, 0.83 ± 0.04, and 0.41 ± 0.58, respectively.

<table>
<thead>
<tr>
<th>Network/ Seg. task</th>
<th>DSC</th>
<th>Jaccard</th>
<th>Sensitivity</th>
<th>Specificity</th>
</tr>
</thead>
<tbody>
<tr>
<td>1st nnU-Net</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Breast region</td>
<td>0.95±0.00</td>
<td>0.90±0.01</td>
<td>0.95±0.03</td>
<td>0.99±0.00</td>
</tr>
<tr>
<td>(Test dataset)</td>
<td>(0.95±0.00)</td>
<td>(0.91±0.01)</td>
<td>(0.96±0.02)</td>
<td>(0.99±0.00)</td>
</tr>
<tr>
<td>Fat</td>
<td>0.95±0.00</td>
<td>0.91±0.00</td>
<td>0.98±0.02</td>
<td>0.99±0.00</td>
</tr>
<tr>
<td>(Test dataset)</td>
<td>(0.96±0.00)</td>
<td>(0.93±0.00)</td>
<td>(0.96±0.04)</td>
<td>(0.99±0.00)</td>
</tr>
<tr>
<td>2nd nnU-Net</td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>FGT</td>
<td>0.83±0.04</td>
<td>0.72±0.07</td>
<td>0.77±0.01</td>
<td>0.99±0.00</td>
</tr>
<tr>
<td>(Test dataset)</td>
<td>(0.84±0.03)</td>
<td>(0.73±0.05)</td>
<td>(0.86±0.18)</td>
<td>(0.99±0.00)</td>
</tr>
<tr>
<td>Tumor</td>
<td>0.41±0.58</td>
<td>0.35±0.49</td>
<td>0.45±0.64</td>
<td>0.99±0.00</td>
</tr>
<tr>
<td>(Test dataset)</td>
<td>(0.35±0.48)</td>
<td>(0.27±0.37)</td>
<td>(0.33±0.45)</td>
<td>(0.99±0.00)</td>
</tr>
</tbody>
</table>

*Results displayed were from the fold 2 model among 5-fold CV.

Figure 3 illustrates a case of an image prediction of our DL method in comparison to the ground truth. In addition, an applicable phantom made based on the segmentation results is shown in Figure 4.

Figure 3. (a) shows a prediction example of the breast region mask in white overlaid on its ground truth. (b) shows the GT of 3-class segmentation where fat tissue is represented in yellow, FGT in blue, and tumor in red. (c) shows the prediction of the case (b).

Our breast segmentation task is implemented by using two consecutive networks. We utilized 5-fold CV models for every network. The training time needed an average of 145 s and 170 s for one epoch during model training for one model of breast region and 3-class inner tissue segmentation networks, respectively. Accordingly, 40 h and 47 h are taken to complete every model's training process.

4 DISCUSSION AND FUTURE WORK

We exhibited a DL approach to address the clinical need to segment breast tissues and form a groundwork for image navigation since traditional methods do not provide good performance to separate the breast region and label breast tissues. Thus, our segmentation pipeline has two consecutive networks of nnU-Net architecture. Accordingly, we used the trained model and the segmentation results to develop a PVA-C breast phantom to support and experiment with navigation surgery studies.
Figure 4. illustrates the produced PVA-C breast phantom, where, at (a), the yellow corresponds to the segmented volume of FGT, and the red corresponds to the tumor volume. (b) A complete breast phantom of PVA-C mimics fat, FGT, and tumor. (c) MRI image of PVA-C breast phantom.

We used an open-source for our dataset that included T2W/STIR MRI images for 10 patients. As shown in Table 1, our results demonstrated high sensitivity, and specificity, avoiding over-segmentation and preserving segmentation sensitivity.

We compared our outcomes with other deep-learning methods, as shown in Table 2. (Dalmış et al. 2017, Jiao et al. 2020). Therefore, our pipeline had accomplished exceptional performance with a smaller dataset, specifically for breast region and fat segmentation, with a DSC value of 0.95 and 0.95, respectively. However, the DSC values of literature (Dalmış et al. 2017, Jiao et al. 2020) in Table 2 were near to our results, but the training cases were much more than ours, and their segmentation object is for two regions.

Table 2. DSC values comparison of our method and other literature*.

<table>
<thead>
<tr>
<th>Author</th>
<th>Method, number of the training dataset</th>
<th>Segmentation tasks</th>
<th>Tumor sensitivity</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Breast Region</td>
<td>Fat</td>
<td>FGT</td>
</tr>
<tr>
<td>Dalmış et al. (2017)</td>
<td>2-D U-Net</td>
<td>66</td>
<td>-</td>
</tr>
<tr>
<td>Jiao et al. (2020)</td>
<td>U-Net ++</td>
<td>75</td>
<td>-</td>
</tr>
<tr>
<td>Ours</td>
<td>3-D nnU-Net</td>
<td>8</td>
<td>0.95 ± 0.00</td>
</tr>
</tbody>
</table>

*Tumor sensitivity results were displayed for comparison purposes.

The tumor segmentation showed DSC and sensitivity of 0.41 and 0.45, respectively, implying that a considerable tumor mass was wrongly unsegmented (i.e., high false negative). Yet, expanding the training dataset is expected to support overcoming this challenge and improve the segmentation performance even more.

This work presents the foundation of preoperative images to develop a breast surgery navigation system by introducing a DL segmentation framework assembled on the nnU-Net for breast region, fat, FGT, and tumor segmentation. Our group is working on a real-time tracking system for breast tumors using DCNN for pre-and intra-operative imaging. Our pipeline plan will allow us to update in real-time the boundary position of breast tumors based on a patient-specific model by relating preoperative scans to US images, enabling constant visualization. As we showed in this study, the stage of the preoperative image, our intraoperative stage, consists of MRI-US image registration, which also leverages a nnU-Net architecture, this time for synthesizing finite elements. Moreover, the intraoperative stage will employ a combination of MRI-US image registration, optical tracking, and surgical robotic integration. It will be completed in three steps. 1) MRI-US calibration 2) MRI-US fiducials-based point-cloud affine registration 3) DL-based elastic registration Figure 5. Our navigation system will create a real-time soft-tissue tracking system to improve tumor localization during surgical procedures. The position of lesions localized on preoperative images segmentation will be updated from intraoperative ultrasound data and visualized by the surgeon in real-time. Our robotic surgery design is proposed by a co-author (K.K); it is a hand-held grasping robot
equipped with multiple claws that are embedded in a balloon-like covering to perform tumor resection spillage-free.

Figure 5. intraoperative DL-based MRI-US registration stage.

Our study has some limitations. Our small dataset of 10 subjects will grow in future work to increase the performance and reliability. Another limitation is that we have no control over MRI scanning protocols since images are from an online repository. However, even with these limitations, our pipeline allowed us to represent remarkable results and exhibit the potential capabilities of consecutive nnU-Net architectures.

5 CONCLUSION

This paper described the preoperative DL segmentation pipeline stage for preoperative breast surgery planning from MRI, built on the nnU-Net. We also introduced a mimicking breast phantom made on these segmentation results. We took advantage of multi-modality breast MRI datasets acquired from a public archive. The nnUNet-based framework exhibited high segmentation accuracy among everyday breast MR images without post-processing or fine-tuning steps. In short, this study establishes the base for developing and validating robotic experiments of the intra-operative navigation system based on 3DUS. Elastic MRI-3DUS registration, as well as based on DNNs, for finite elements synthesis, will be emphasized in the near future.

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