RFA-Cut: Semi-automatic Segmentation of Radiofrequency Ablation Zones with and without Needles via Optimal s-t-Cuts

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Abstract— In this contribution, we present a semi-automatic segmentation algorithm for radiofrequency ablation (RFA) zones via optimal s-t-cuts. Our interactive graph-based approach builds upon a polyhedron to construct the graph and was specifically designed for computed tomography (CT) acquisitions from patients that had RFA treatments of Hepatocellular Carcinomas (HCC). For evaluation, we used twelve post-interventional CT datasets from the clinical routine and as evaluation metric we utilized the Dice Similarity Coefficient (DSC), which is commonly accepted for judging computer aided medical segmentation tasks. Compared with pure manual slice-by-slice expert segmentations from interventional radiologists, we were able to achieve a DSC of about eighty percent, which is sufficient for our clinical needs. Moreover, our approach was able to handle images containing (DSC=75.9%) and not containing (78.1%) the RFA needles still in place. Additionally, we found no statistically significant difference (p < 0.423) between the segmentation results of the subgroups for a Mann-Whitney test. Finally, to the best of our knowledge, this is the first time a segmentation approach for CT scans including the RFA needles is reported and we show why another state-of-the-art segmentation method fails for these cases. Intraoperative scans including an RFA probe are very critical in the clinical practice and need a very careful segmentation and inspection to avoid under-treatment, which may result in tumor recurrence (up to 40%). If the decision can be made during the intervention, an additional ablation can be performed without removing the entire needle. This decreases the patient stress and associated risks and costs of a separate intervention at a later date. Ultimately, the segmented ablation zone containing the RFA needle can be used for a precise ablation simulation as the real needle position is known.

I. INTRODUCTION

Mainly because of hepatitis infections and alcohol abuse, liver cancer is on the rise worldwide. Especially patients with Hepatocellular Carcinomas (HCC) have a poor prognosis because of its late symptomatic onset resulting in a median survival time of four to six months from the time of diagnosis when untreated. However, according to the recent treatment guidelines [1], Radiofrequency ablations (RFA) serve as a first

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line therapy approach for early HCC in patients with liver cirrhosis. In addition, the local usage of ablation therapies increases for metastatic liver disease. Originally developed for patients who were not eligible for surgery, the usage has now expanded to patients as a bridge to liver transplantation and even as an alternative to surgical resection in the early stages of the disease [2]. In the early 1990s, RFA was first described followed by huge technical advances throughout the last decades. In summary, the underlying principle is based on a high frequency alternating current which is delivered through one or more electrodes placed inside the lesion [3] (Fig. 1.).

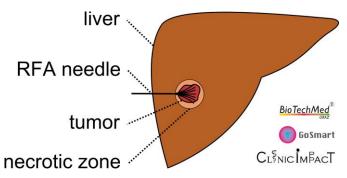


Fig. 1. Sketch of a liver (brown) with an umbrella-shaped and fully expanded radiofrequency ablation (RFA) needle in black. The needle tips are located inside a liver tumor (red) which is surrounded by the necrotic zone (light brown).

The heat that is generated around the electrodes destroys the cancer cells by inducing a coagulative necrosis as a result of denaturation of cellular proteins. In general, tissue necrosis already begins at approximately 60°C, but in general, temperatures over 100° are needed to achieve satisfying results. Also the placement of the needle and the individual impedance determine the amount of destroyed tissue. Further, it is inversely proportional to the square of the distance from the electrode, and as a result, tissue cools rapidly away from the tip of the needle probe. Thus, the proximity to large blood vessels plays also a major role in the heat transmission.

Normally, the blood flow protects the vessel wall from getting damaged, however, on the flipside acts as a heat sink by cooling down nearby tissue limiting the methods overall success [4]. As a consequence, a significant mismatch between expected and truly induced lesion size and geometry has been observed in many liver RFAs. On one side, it can lead to overtreatment with severe injuries (up to 9% major complications [5]) or, on the other side, to under-treatment with tumor recurrence (up to 40% [5]). Cohort patient studies have shown the evidence of a significant reduction in the recurrence rate, if there is a safety rim generated around the tumor via the RFA [6]. This elicits the need for (1.) a reliable method for the comparison of localization, geometry and size of the tumor in the preoperative images, and (2.) a thermally induced lesion after ablation. Tumor recurrence can be diagnosed by the detection of typical alteration in tissue enhancement, but an increase of the size or a change of the geometry of the lesion seems to be a more sensitive indicator of early recurrence in follow-up imaging. Thus, a reliable and feasible determination of the ablation zone at baseline and follow-up may contribute to a positive outcome for the patient and can therefore lead to a better understanding of the cause of new tumor growth. And the additional knowledge might lead to an improvement of ablation protocols or even in the long term to new treatment strategies. However, the determination of therapeutically induced lesions after minimally invasive cancer treatment can be performed by segmentation, which is usually done by a very time consuming manual (slice-by-slice) procedure – and thus not yet part of the clinical routine. But a validated segmentation algorithm may have the potential to increase the acceptance of the method in the medical community and finally lead to a benefit in patient treatment. The overall segmentation (and partially also registration) field in Computer Vision deals with the computer-aided analysis and classification of image data in a broad range of applications [7]-[19]. This can be the automatic detection of humans in videos [20] or the volumetry of brains from medical images [21]. The aim of a segmentation algorithm is mainly to support and speedup a time-consuming manual selection and contouring process. In the literature, a number of algorithms have been proposed, like Active Contours (ACM) [22]-[25], Deformable Models [26]-[32], Active Appearance Models (AAM) [33]-[36], graph-based approaches [37]-[40], fuzzybased approaches [41], or neural networks [42], which often base on a mathematical model from another discipline, like Physics or Electrical Engineering, or a combination of several mathematical models. Furthermore (or alternatively), the segmentation methods can be classified as fully-automatic or semi-automatic algorithms. A research group working in the specific area of (semi-)automatic segmentation of ablation zones in RFA scans of the liver are Passera et al. [43]. They apply a Live-Wire algorithm (implemented within the MeVisLab platform [44], www.mevislab.de) and clustering for the segmentation, but they did not use data where the RFA needle is still present within the scan. However, this is the case when the interventional radiologist (or interventionalist) particularly wants to assess the size of the induced lesion, to check if an additional ablation is needed. And to avoid the repositioning or the replacement of the needle, it remains inside the liver while performing the control scan. In addition, the introduced segmentation method worked only in 2D, which can be very time-consuming in case of tumors or

ablation zones that extend over many slices and the reported segmentation time was ten minutes. McCreedy et al. [45] presented another 2D livewire-based method, a radio frequency ablation registration, segmentation, and fusion tool called RFAST. However, the (segmentation) process has not been described in detail and no quantitative segmentation results have been presented. Keil et al. [46] presented the semiautomated segmentation of liver metastases treated by radiofrequency ablation with an algorithm that consists of six steps including a three-dimensional region growing and morphologic operations, like erosion, dilation. In addition, the user has to draw a diameter across larger lesions or provide a single click inside smaller lesion as initialization. A workflow oriented software support for image guided RF ablations of focal liver malignancies has been introduced by Weihusen et al. [47]. Thereby, they also segment coagulation necrosis in the - post-interventional - control scans (after the ablation). After the user has provided a seed point inside the ablation zone, a morphology-based region growing algorithm is started, which has been proposed by Kuhnigk et al. [48]. Finally, the user has the option to correct the segmentation outcome towards more "irregular" or "roundish" geometry by manual interaction. Initial results of a 3D shape-based analysis of CT acquisitions for the detection of local recurrence of liver metastases after RFA treatment, have been reported by Bricault et al. [49]. In doing so, they applied a semi-automated 3D segmentation method that uses a "tagged" watershed algorithm. On average, the segmentation process took about four minutes and the minimum required user interaction included two mouse clicks: the first one in the ablated tumor and the second one in the surrounding liver parenchyma. Elsewhere, Stippel et al. [50] introduced a volumetric evaluation study of the variability of the size and the shape of necrosis that have been induced by RF ablations in human livers. Therefore, the study has been performed with the software package VA 40C from Siemens on a Leonardo workstation. For the overall segmentation process of the ablation-induced lesions, a region of interest had to be defined in each slice by manually tracking the approximate lesion borders. Then, the precise border of the lesion was determined by a filter algorithm from the software package, which was based on density differences between the ablated and the liver tissue. In addition to this section, we want to point the interested reader to a state of the art publication computer-assisted planning, intervention about and assessment of liver tumor ablation from Schumann et al. [51].

Finally, to the best of our knowledge, there is no work that has studied the semi-automatic 3D segmentation of postinterventional RF ablation zones with clinical data that has the ablation needles still in place.

II. MATERIAL AND METHODS

A. Data Acquisition

We used twelve datasets from ten patients, who underwent radiofrequency ablation in the liver, for this retrospective study [52], [53]. The number of slices in z direction ranged from 52 to 232 and all datasets had a matrix size of 512x512 in x- and y-direction. The slice thickness was either one or two mm, and the pixel spacing ranged from 0.679 to 0.777, with spacing between the slices of one to three mm. The ablation needle was still present in six datasets, and all datasets have been acquired on multislice CT scanner (Philips Brilliance or Mx8000, Philips Healthcare, Netherlands). The data analysis and publication, was approved by the Institutional Review Board (IRB) of the Leipzig University Hospital under the reference number: 381-14-15122014. In addition, we plan to provide the datasets to the research community on our ClinicImppact project page:

http://www.clinicimppact.eu/

Medical ablation datasets from a comprehensive pig study can already be found on the webpage of our European Project named *GoSmart* [54]:

http://www.gosmart-project.eu/

B. Manual Segmentation

The Ground Truth of the ablation zones has been generated by setting up a simple segmentation framework for manual contouring under MeVisLab. This allowed the physicians to manually outline the RFA lesions slice-by-slice without any algorithmic support (to avoid distortions). Then, the single contours were voxelized and merged to a 3D mask, which represented the manual segmented ablation zone. Afterwards, the 3D masks were used for comparison and quantitative evaluation with the RFA-Cut segmentation results.

C. Evaluation Metric

For evaluation we used the Dice Similarity Coefficient (DSC) metric [55]-[58], widely used in medical image analysis and describing the agreement between two binary volumes M and S:

$$DSC = \frac{2 \cdot V(M \cap S)}{V(M) + V(S)} \tag{1}$$

M and S are the binary masks from the manual (M) and the semi-automatic RFA-Cut (S) segmentations, $V(\cdot)$ denotes the volume (in cm³) and (\cap) denotes the intersection. The volume was computed by counting the number of voxels and multiplying them with the physical size of voxels. However, in addition to the DSC, we measured also the time it took an experienced radiologist to manually outline the ablation zones and compared it with the computation time of the semi-automatic RFA-Cut segmentations.

D. Semi-automatic Segmentations with the RFA-Cut

The semi-automatic segmentation algorithm named RFA-Cut uses a spherical template to set up a three-dimensional graph G(V,E) around the ablation zone [59]. Altogether, the Graph G consists of nodes $n \in V$ and edges $e \in E$ connecting these nodes, and a source s and a sink t [60]-[63]. The nodes n are sampled along rays whose origin resides at the user-defined seed point (inside the image), while their direction points towards the surface of a polyhedron [64]. After the graph has been constructed (for more details please see [65]-[69] and its origin for a segmentation and stent simulation [70]-[79]), the segmentation result is calculated by dividing the graph into two sets of nodes via a Min-Cut/Max-Flow algorithm [80]. In doing so, one set of nodes represents the ablation zone (called foreground) and the other set represents the surrounding structures (called background). Note that the basic segmentation scheme has already been successfully applied to pituitary adenomas [81] and prostate central glands [82].

However, the rigid segmentation scheme was later turned into an interactive real-time approach [83]-[87], and for this study it was enhanced by an additional refinement option [88]-[91]. To initialize the interactive segmentation process, the user has to place only one seed point roughly in the middle of the ablation zone on a 2D slice of the (CT) scan (note: the segmentation, however, performs in 3D). From the seed point position, the graph is generated in the background (not visible to the user) and segmentation result is automatically calculate (min-cut) and displayed to the user in 2D and 3D views. Nevertheless, the user can drag the seed point around to interactively generate new segmentation results, which always depend on the new/current seed point positions. Moreover, the user can drop the seed point (by means it remains at a fixed position in the image) and add an arbitrary amount of additional seed points along the border of the ablation zone. In doing so, the algorithm gets additional input about the location of the border and steers the behavior of the min-cut. Though, the user can always come back to the initial seed point, grab it and start dragging it around the image again. Note that the additional placed seed points along the ablation boarder will not get lost then, they stay fixed and continue to influence/restrict the min-cut calculation. The schematic workflow of the RFA-Cut is shown in Fig. 2.

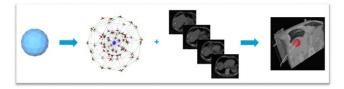


Fig. 2. Schematic workflow of the RFA-Cut segmentation: a sphere (left, blue) is used to construct a graph (second image from the left). Then, the graph is constructed (note: not visible to the user) at the user-defined seed point position within the image (third image from the left). Finally, the segmentation result of the RFA-Cut (red) corresponding to the current seed point is shown to the user (rightmost image).

III. RESULTS

Our proposed interactive segmentation algorithm RFA-Cut has been implemented as an own C++ module (under Microsoft Visual Studio 2008) for the medical prototyping platform MeVisLab. The overall computation of the segmentation result – including the graph construction from the current user-defined seed point position and the optimal min-cut calculation – could be performed within one second on a laptop (with Intel Core i5-750 CPU, 4×2.66 GHz, 8 GB RAM, Windows 7 Professional x64 Version with Service Pack 1 installed). As a consequence, this enabled real-time feedback of our algorithm presented to the user and the immediate response and feedback of the segmentation allowed user guidance of the algorithm to a satisfying outcome.

Fig. 3. shows the result of a semi-automatic segmentation of a post-interventional ablation zone for visual inspection. Because the computed tomography (CT) data has been acquired immediately after the treatment, the ablation needle is still in place and thus visible in the scan. The image on the left shows the axial slice with a user-defined seed point (blue) and the red dots are the border of the segmentation in this slice (note: these dots represent the last nodes of the graph that are still bound to the source *s* after the min-cut calculation). The image in the middle shows the segmentation result in a 3D view, and the red dots show the last nodes of the graph which are still connected to the source after executing the min-cut calculation. The rightmost image presents a closed surface in green, which represents the segmentation result, and that has been generated from the graph's nodes. Later, this closed surface is used to generate the solid 3D mask for the Dice Similarity Coefficient calculation with a pure manual slice-by-slice segmentation.

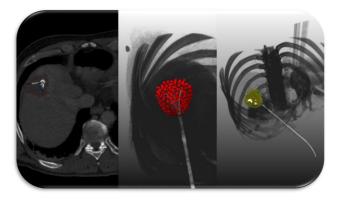


Fig. 3. RFA-Cut segmentation result of an ablation zone in a postinterventional CT scan (still including the ablation needle). The axial slice where the user-defined seed point (blue) has been placed, is shown in the left image. In addition, the corresponding segmentation result (red dots) is presented (note: the red dots are the last nodes of the graph that are still bound to the source s after the min-cut calculation). A 3D view of all nodes (red dots) that describe the surface of the segmented ablation zone after the mincut calculation is shown in the middle image. Moreover, the rightmost image displays a closed surface (green) of the segmentation result of the RFA-Cut (red dots from the previous image). Finally, the closed surface of the segmentation result is used to generate a solid mask for the DSC calculation.

Table 1 and Table 2 present the direct comparison of manual slice-by-slice segmentations from physician 1 and 2 with RFA-Cut segmentation results for twelve ablation zones using the Dice Similarity Coefficient. In addition, Table 3 presents the direct inter-observer comparison of manual slice-by-slice segmentations from physician 1 and physician 2 via the DSC.

| | Volume (cm ³) | | Number of Voxel | | DSC |
|----------|---------------------------|-------|-----------------|---------|------|
| - | manual 1 | semi | manual 1 | semi | (%) |
| | 10.0 | 6.3 | 5866 | 3689 | 71.8 |
| Range | - | _ | - | _ | _ |
| | 122.6 | 104.0 | 70806 | 70208 | 83.5 |
| μ | 35.9 | 33.0 | | | 77.0 |
| ± | ± | ± | 31294.8 | 30756.3 | ± |
| σ | 30.0 | 25.1 | | | 4.7 |

Tab. 1. This table presents the direct comparison of manual slice-by-slice segmentations from physician 1 and semi-automatic RFA-Cut segmentation results for twelve ablation zones via the DSC.

| | Volume (cm ³) | | Number of Voxel | | DSC |
|----------|---------------------------|-------|-----------------|---------|------|
| - | manual 2 | semi | manual 2 | semi | (%) |
| Range | 11.1 | 6.3 | 6543 | 3689 | 68.1 |
| | - | - | - | - | - |
| | 117.7 | 104.0 | 67963 | 70208 | 85.3 |
| μ | 36.2 | 33.0 | | | 77.1 |
| <u>+</u> | ± | ± | 31240.1 | 30756.3 | ± |
| σ | 28.7 | 25.1 | | | 5.8 |

Tab. 2. Similar to the first table, this table presents the direct comparison of the RFA-Cut segmentation results with physician 2.

| | Volume (cm ³) | | Number of Voxel | | DSC |
|----------|---------------------------|----------|-----------------|----------|------|
| | manual 1 | manual 2 | manual 1 | manual 2 | (%) |
| Range | 10.0 | 11.1 | 5866 | 6543 | 82.4 |
| | _ | _ | _ | _ | _ |
| | 122.6 | 117,7 | 70806 | 67963 | 92.6 |
| μ | 35.9 | 36.2 | | | 88.8 |
| ± | ± | ± | 31294.8 | 31240.1 | ± |
| σ | 30.0 | 28.7 | | | 3.3 |

| Tab. 3. This table presents the direct comparison of the manual slice-by-slice |
|--|
| segmentations from physician 1 and physician 2 (from table 1 and table 2) |
| for the twelve ablation zones also via the DSC. |

Furthermore, we differentiated between the acquisitions where the RF electrodes where still in place and the cases where the RF electrodes have already been removed. However, we found no significant differences between these cases, and the DSC values between experts were significantly higher (p < 0.01) than those between automatic and manual processing (88.8% vs. 77.0%). This was also independent of whether the needle was still included (86.8% vs. 75.9%, p < 0.05) in the dataset or not (90.9% vs. 78.1%, p < 0.05), respectively. For automatic processing the segmented volumes appeared to be smaller than for experts (33.03 ml vs 36.02 ml), but the differences were not statistically significant: p=0.308. This was also the case for scans with the needle included (25.76 ml vs. 25.93 ml, p=0.917) and without (40.30 ml vs. 46.10 ml, p=0.249). The overall mean DSC value of both experts appeared to be smaller when the needle was present (75.9% vs. 78.1% without, p=0.423), however, the difference was not significant. On the other hand, the inter-observer DSC was significantly higher when the needle was not present (90.9% without vs. 86.8%, p=0.025). The statistical differences in the DSC values and segmentation volumes between the methods were analyzed with Wilcoxon signed-rank tests. And differences between both subgroups (with and without needle present) with a Mann-Whitney test. All analyses were performed with Version 20 of SPSS from IBM (using a level of significance of 0.05). However, in summary, the main difference between the automatic and manual segmentation is the computation time. Making use of the introduced automatic tool, a segmentation could be performed in a few seconds, whereas a pure manual segmentation took on average between 48 seconds and 8 min 16 seconds (mean 3 min 13 seconds).

IV. CONCLUSIONS

Radiofrequency ablations of liver tumors induces areas of tissue necrosis, which can be visualized reliably in contrast enhanced CT scans. In this paper, an interactive segmentation algorithm was applied to routine control CT scans after RFA of liver cancer for the semi-automatic determination of thermally induced lesions. Semi-automatic segmentation accuracy was found to be sufficient for most of the cases, although a manual slice-by-slice segmentation still provided the best accuracy. However, speed was the main advantage of the proposed tool over manual segmentation, which makes it an appealing alternative for physicians. As stated in the introduction, a minimally invasive RFA can be an alternative to open surgery and might also be suitable for inoperable patients. In addition, when the interventional radiologist presumes that continuing of the therapy might be necessary, the RFA needle may still reside in the target organ during image acquisition (note that post-interventional imaging is regularly performed to document the success of the RFA treatment). Furthermore, hardening artefacts can compromise the image quality significantly, but such cases were also segmented and included in this study. Manual slice-by-slice segmentations have been performed by two radiological experts for the evaluation of the RFA-Cut; this enabled the DSC calculation between the manual and the semi-automatic RFA-Cut segmentation outcome. In conclusion, the achieved highlights of this publication are:

- Applying an interactive segmentation algorithm to RFA acquisitions from the clinical routine;
- Using post-interventional patient scans with and without RFA needles still in place;
- Clinical experts performed manual slice-by-slice segmentations for an evaluation;
- DSC calculation for a statistical validation of the RFA-Cut;

For a comparison of our employed method with an existing segmentation approach, we applied the GrowCut implementation in (3D) Slicer (www.slicer.org) [92], [93] to our data. This GrowCut implementation is user friendly, because it does not require any precise parameter setting. Rather the user initializes the algorithm by marking areas in the image with several simple strokes (Fig. 4.). Moreover, we've had already good experiences with certain types of brain tumors (Glioblastoma multiforme (GBM) [94] and pituitary adenomas [95]) and vertebral bodies [96]. But especially the cases where the needles were still in place and therefore visible within the images caused segmentation problems (Fig. 5.). In more detail, GrowCut leaks along the needles, because it cannot handle the large gray value differences between the ablation zone (dark) and the RFA needle (bright). In addition, the initialization of GrowCut (marking parts of the lesion and the background) takes a trained user between thirty and sixty seconds, in contrast to our method that needs only a single seed point. Furthermore, the user has to wait several seconds for the GrowCut segmentation result (about ten seconds on the PC we used also for interactive method). In contrast, our method provides the segmentation result immediately, which also makes refinement much more convenient. The sharp edges of the GrowCut segmentation result (green) in the rightmost image of Fig. 5. occur, because the (3D) Slicer implementation restricts the segmentation area with a bounding box - note that the size of the bounding box depends on the manual GrowCut initialization of the user.

There are several areas for future work, like the integration of the interactive segmentation algorithm into a medical application framework for supporting ablation therapies. The framework is currently under development within a project funded by the European FP7 program and is based the continuation on the European project IMPPACT (Image-based Multi-scale Physiological Planning for Ablation Cancer grant Treatment, agreement no. 223877) [97] (http://www.imppact.eu/). More precise, we plan to use our semi-automatic algorithm for the segmentation of difficult cases where automatic methods fail. Furthermore, we plan to provide the datasets acquired during the project over the next

years to the public. Finally, we want to investigate the possibilities for liver tumor tracking after one or several RFA interventions and support the needle placement [98] with our real-time segmentation algorithm and motion tracking [99] in an Augmented Reality (AR) setting [100].

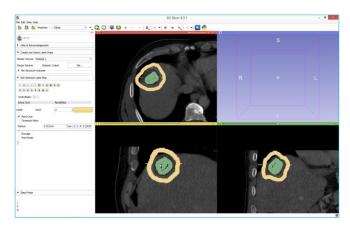


Fig. 4. Manual GrowCut initialization for the segmentation of the RF ablation zone: the ablated zone is marked in green and the background is marked in yellow on three 2D slices, respectively.

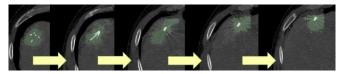


Fig. 5. The GrowCut segmentation result (green) for the manual initializing from Fig. 4. The GrowCut segmentation leaks along the RF ablation needle, because it cannot handle the large gray value differences between the ablation zone (dark) and the needle (bright). The sharp edges of the segmentation result in the rightmost image occur, because the GrowCut implementation under Slicer automatic restricts the segmentation area with a bounding box that depends on the user initialization.

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https://www.youtube.com/c/JanEgger/videos

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