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## Variabilities of MRI-, CT- and PET-CT-based tumor and lymph node delineations for lung cancer radiotherapy planning

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## Abstract

**Purpose**—Target delineation in lung cancer radiotherapy using CT and/or PET-CT is affected by large variability. MRI has excellent soft tissue visualization and better spatial resolution than PET-CT. The main purpose of this study is to analyze delineation variability for lung cancer using MRI.

**Methods and materials**—Seven physicians delineated the tumor volumes of ten patients for the following scenarios: (1) CT only; (2) PET-CT fusion images registered to CT ("clinical standard"); and (3) post-contrast T1-weighted MRI registered with diffusion-weighted MRI. To compute interobserver variability, the median surface was generated from all observers' contours and used as the reference surface. A physician labeled the interface types (tumor to lung, atelectasis (collapsed lung), hilum, mediastinum, or chest wall) on the median surface. Contoured volumes and bidirectional local distances (BLDs) between individual observers' contours and the reference contour were analyzed.

**Results**—CT- and MRI-based tumor volumes normalized relative to PET-CT-based volumes were31.62±0.76 (mean±SD) and 1.38±0.44, respectively. Volume differences between the imaging modalities were not significant. Between observers, the mean normalized volumes per patient averaged over all patients varied significantly by a factor of 1.6 (MRI) and 2.0 (CT and PET-CT) (p=4.10×10<sup>-5</sup> – 3.82×10<sup>-9</sup>). The tumor-atelectasis interface had a significantly higher variability

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than other interfaces for all modalities combined (*p*=0.0006). The interfaces with the smallest uncertainties were tumor-lung (on CT) and tumor-mediastinum (on PET-CT and MRI).

**Conclusions**—While MRI-based contouring showed overall larger variability than PET-CT, contouring variability depended on the interface type and was not significantly different between modalities despite of the limited observer experience with MRI. Multimodality imaging and combining different imaging characteristics might be the best approach to define the tumor volume most accurately.

## I. INTRODUCTION

Target delineation in lung cancer radiotherapy has, in general, large variability [1,2]. Reliable delineation of the target volumes is required for accurate radiation dose delivery [3,4]. PET-CT has been shown to improve target definition by improving the accuracy of determining tumor volume and localization [5,6]. Therefore, PET-CT is routinely used for lung cancer target delineation for radiotherapy planning. However, even with PET-CT target delineation remains highly variable [7–9]. Compared to PET-CT, MRI has good contrast between normal and cancerous tissues with higher spatial resolution [10]. MRI has been shown to be superior to CT for identifying tumor extensions to chest wall, mediastinum, vascular and neural structures, such as brachial plexus [10–14]. In conjunction with diffusion-weighted sequences (DW-MRI), MRI has been shown to be comparable or even superior to PET-CT in primary tumor and nodal staging [15–18]. MRI has rarely been evaluated for lung cancer target definition in radiotherapy. In a recent study, DW-MRI-based gross tumor volumes (GTVs) were reported to be as reliable as PET-CT-based GTVs [19].

Several methods have been used to evaluate interobserver variability in target volume delineation [4,20]. Typical approaches include assessment of centroid shifts or volumetric overlap measures, e.g., conformity index and dice similarity coefficient [3,20,21]. To understand the association of contour uncertainty with anatomical information presented on multimodality imaging, in the present study voxel-wise contouring variability on 3D volume surfaces was analyzed. The standard deviation (SD) of the distances of the multiple observers' contour surfaces from the reference contour surface, defined as the median contour, was assessed in relation to tumor-tissue interface types [22,23] using the bidirectional local distance (BLD) measure [24]. The purpose of this study is to investigate interobserver delineation variability for GTVs of primary lung tumors (PTs) and associated pathologic lymph nodes (LNs) and for their interfaces with surrounding tissues using MRI, and to compare the results to CT alone- and PET-CT-based delineations.

## **II. METHODS AND MATERIALS**

#### **II.A.** Patients

Multimodality imaging of ten patients with non-small cell lung cancer (NSCLC) obtained on an IRB-approved protocol (NCT xxx) was used for this study (Table 1). Patients underwent MRI, CT and 2-deoxy-2-(<sup>18</sup>F)fluoro-D-glucose (FDG) PET-CT before conventionally fractionated radiation therapy. The average time between CT and PET-CT was 17 days (range: 2–39 days), between CT and MRI 33 days (9–52), between PET-CT and MRI 19

days (10–36). Nine of the ten patients had primary tumors in lung and associated lymph nodes; one patient had pathologic LN involvement only.

#### **II.B.** Imaging modalities

Diagnostic CT: Diagnostic CT image sets with intravenous contrast were used for delineation in the CT only and PET-CT sessions, slice thickness was 3.00 mm and pixel size  $\sim 0.58 \times 0.58 \text{ mm}^2$ .

PET-CT: Fusion images of the attenuation-corrected PET images with the corresponding CT images of PET-CT scans were used for delineation after rigid registration to diagnostic CT images. The PET images had a slice thickness of ~4.25 mm and in-plane resolution of  $\sim$ 3.91×3.91 mm<sup>2</sup>.

MRI: Images from a 1.5 T MRI scanner included DW-MRI and post Gd-DTPA T<sub>1</sub>-weighted ultrafast gradient echo volume interpolated breath-hold examination (VIBE) images. Apparent diffusion coefficient (ADC) maps were obtained from the DW-MRI images of eight b-values =  $0-1000 \text{ s/mm}^2$ . VIBE images had a slice thickness of ~2.00 mm and an inplane pixel size of ~1.56×1.56 mm<sup>2</sup>. DW-MRI images had pixel size of ~1.98×1.98 mm<sup>2</sup>, slice thickness/gap = 6 mm/1.2 mm. Each frame of a DW-MRI scan had 7 slices. One or more frames were used to cover the whole tumor volume.

#### II.C. Target delineations

Five radiation oncologists and two chest radiologists were asked to independently delineate GTV of PT (GTV\_PT) and GTV of LN (GTV\_LN) using MIM Software (MIM Software Inc., Cleveland, OH, USA). During contouring, sagittal and coronal views were displayed. To avoid potential confounding errors for identifying and contouring different lesions, the physicians were informed on the locations of GTV\_PT and/or GTV\_LN by marking the location in a representative axial slice of the images. Seven lymph nodes were contoured in four patients. Only well demarcated, clearly visualized paratracheal and hilar nodes were selected for contouring. The following delineation sessions were created (see example in Figures 1 and S1):

- 1. CT only: The GTV\_PT contour was drawn on the lung/mediastinal window on diagnostic CT images, depending on tumor location. GTV\_LN structures were contoured in the mediastinal window.
- 2. PET-CT: Users were instructed to first use a gradient-based auto-segmentation tool (PET-Edge, MIM) [25]. Users then manually edited the volumes on the CT images.
- 3. MRI: Users were provided VIBE images rigidly registered with DW-MRI images and ADC maps (Fig. 1). DW-MRI images of the highest b-value (b = 1000 s/mm<sup>2</sup>) or, in case of very low signal to noise ratio, the second highest b-value (b = 800 s/mm<sup>2</sup>) were used as additional information for the delineation on VIBE images. Since most of the users had little or no experience with tumor delineation on VIBE, a presentation was prepared in collaboration with a specialized chest radiologist showing contouring examples for tumor and

atelectasis on VIBE and DWI scans for various clinical scenarios including contour recommendations.

#### II.D. Analysis

The median contour was generated from all observers' contours for each patient and modality separately and was selected as the reference contour. The median contours were generated for each GTV following the method of Steenbakkers et al. [23]. Briefly, a voxel was labeled as part of the median GTV if at least 50% of the observers (4/7 observers) labeled this voxel as GTV. All voxels labeled as median GTV were then combined into the median GTV contour, as shown in Fig. 1. A physician labeled the interface types (GTV\_PT or GTV\_LN to lung, atelectasis (collapsed lung), hilum, mediastinum, airways, blood vessels and/or chest wall) on the median surface as applicable to the respective GTV\_PT and GTV\_LN structures (Fig. 2).

The mask images of the 3D contours including individual observers' contours, median contours and contours identifying the interfaces were processed using MATLAB. A marching cubes algorithm was used to generate a surface contour mesh from the 3D volume of the binary mask images with a threshold value of 0.5 [26]. The BLD measures for the observers' contours from the reference median contour were obtained for the different interfaces and the whole 3D surface using the methods described by Kim et al [24]. BLD was employed in the presence of complex lung tumor contours to reduce errors of the minimum distance (MD) analysis and other conventional distance-based methods by taking into account the bidirectional characteristics with forward and backward directions.

BLD has been shown to be more appropriate than other distance measures, especially around concave or folding regions of either reference or individual observer's contours or both, where MD can be too short; normal distance either cannot be established or is erroneously very long; and ComGrad (method using the combined gradient of the signed distance transforms produced from the reference and observer's contours) cannot find corresponding points (see [24] for more details). The procedure for obtaining BLD is given below:

The forward minimum distance, d<sub>min</sub>(P, t), from a point P on a reference contour surface r to an individual observer's contour surface t is determined (see Figure S2):

Forward minimum distance= $d_{\min}(P, t) = \min\{d(P, T_i)\},\$ 

where points  $T_i$  (i = 1, 2, ..., N (total number of voxels on t)) are on contour t, and  $d_{min}$  represents the minimum distance. This step is the same as obtaining MD.

- ii. All possible backward minimum distances from the points  $T_j$  to r such that  $d_{min}(T_j, r) = d(T_j, P)$ , are identified.
- **iii.** BLD at P is identified as the greatest among these forward and backward minimum distances obtained in steps (i) and (ii) (in Figure S2, T<sub>3</sub>P is the BLD measure at P). Note that  $d_{min}(T_j, r) = d(T_j, P) \quad d_{min}(P, t)$ .

Backward minimum distances from a number of points  $T_j$  on contour t to contour r can be obtained at P. If no backward minimum distances to r can be identified at P, then the forward minimum distance, i.e. MD from P is established as the BLD measure. This process is repeated for all voxels on the reference contour surface for each observer's contour surface. The standard deviation of the distance from the observers' contours to a single point on the reference contour was calculated, which we termed the local SD. Root mean square of the local SD values (RMS SD) was obtained as an overall SD representing the variation in distance to all points as a measure of target delineation variability.

Plots of the local SD on the surface mesh (SDSM) were made for visualization of the regional variations of local SDs. For improved visualization (in Fig. 1) 3D Gaussian smoothing of the contour surface with  $\sigma = 1.0$  mm was applied, and a mean filter was then applied to the computed local SD values on the 3D smoothed surface. We tried to keep  $\sigma$  small to preserve the original variations of 3D GTV\_PT or GTV\_LN contours [24, 27]. Local SD and RMS SD were compared across the three imaging modalities and across the observers.

#### II. E. Statistical analysis

Comparisons of GTV\_PT and GTV\_LN volume differences across modalities and observers were performed using a repeated measures ANOVA model with modality and observers as main effects. To investigate the interobserver variability or systematic error made by the observers, the observers' contour volumes relative to the corresponding mean volume were compared for all modalities for both GTV\_PT and GTV\_LN using single factor ANOVA. Overall uncertainties for both GTV\_PT and GTV\_LN were computed for all interfaces and whole surfaces for all three modalities.

A random intercept clustered linear mixed effects model was used to model the regional RMSE in different imaging modalities and interface data using SAS v 9.4 and R v 3.2.1.

## III. RESULTS

#### III.A. Target volume variations between the modalities

The ranges of mean GTV\_PT and GTV\_LN volumes in PET-CT imaging modality were 3.6–284.9 and 5.2–30.4 cc, respectively. Because of a large range of volumes across the patients for each of the three modalities, mean volumes were normalized to the PET-CT volume per patient and averaged over all patients to compare volumes across the imaging modalities. CT- and MRI-based GTV\_PT normalized volumes were  $1.62\pm0.76$  (mean±SD) and  $1.38\pm0.44$  times larger than PET-CT volumes, respectively, whereas GTV\_LN volumes were  $1.02\pm0.21$  and  $1.30\pm0.42$  times larger, respectively (Figure 3). The CT-based absolute volume was larger than the PET-CT volume with an estimated difference of 77.28 cc (95% confidence interval (CI): -48.27, 202.83, p=0.21). The MRI-based volume was also larger than the PET-CT volume with an estimated difference of 35.84 cc (95% CI: -89.71, 161.4, p=0.56).

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#### III.B. Interobserver variability for whole GTV\_PT and GTV\_LN contours

The individual observers' normalized volumes relative to the mean volume per patient averaged over all patients (Fig. 4) were significantly different for all modalities for GTV\_PT (p=4.08×10<sup>-8</sup>, 3.82×10<sup>-9</sup> and 4.10×10<sup>-5</sup>, single factor ANOVA for CT, PET-CT and MRI, respectively) and associated pathologic GTV\_LN (p=0.00057, 0.09057 and 0.00024, respectively). Mean observers' normalized volumes per patient averaged over all patients for GTV\_PT and GTV\_LN varied by a factor of 1.6 and 2.2 for MRI and a factor of 2.0 and 1.7, respectively, for both CT and PET-CT, indicating better observer agreement with MRI for GTV\_PT, but less for GTV\_LN.

#### **III.C.** Delineation variability for interfaces

Using BLD analyses, RMS SD values of the interfaces are reported for both GTV\_PT and GTV\_LN (Fig. 5). The largest mean uncertainties of the whole surface for GTV\_PT were observed in CT (2.96 vs 2.06 and 2.77 mm for PET-CT and MRI, respectively) and for GTV\_LN in MRI (2.22 vs 1.90 and 2.01 mm for CT and PET-CT, respectively). The GTV\_PT -atelectasis interface, which was present in 7/10 patients, had a significantly higher variability than other interfaces for all modalities combined (*p*=0.0006). Also, for the three modalities individually, the GTV\_PT -atelectasis interface had the largest mean overall uncertainty (4.26, 2.53 and 4.13 mm for CT, PET-CT and MRI, respectively). The interfaces having the smallest uncertainty for CT, PET-CT and MRI were GTV\_PT -lung (1.92 mm), GTV\_PT -chest wall (1.61 mm) and GTV\_PT -mediastinum (2.25 mm), respectively. Differences for specific interfaces were not significantly different between modalities.

### IV. DISCUSSION

Contouring variability can exceed other geometric errors [28] with potentially detrimental effects on tumor control [29] and toxicity [30]. Hence, reducing contour variability has clinical implications for improving radiotherapy, particularly for highly conformal and adaptive treatment approaches. Van de Steene et al. [30] described several factors for large inter-observer variation such as methodological problems, inability to differentiate between pathologic or normal structures and tumor, as well as lack of knowledge. Our study aimed to identify tumor boundaries and differentiate between tumor and surrounding structures for various imaging modalities with particular focus on MRI.

As documented in the present study, observer variation in lung cancer delineation using CT only is a major source of geometric uncertainty [1,30–32]. Several studies [5,6,33–39] showed that use of FDG-PET has a large impact on the delineated tumor volume and localization. Bradley et al. [5] demonstrated that PET changes GTV contours in approximately two thirds of the cases compared to CT only. Steenbakkers et al's analysis [23] demonstrated that observer variation decreased with the use of PET-CT. Despite these advantages, the relatively low resolution of PET (compared to CT), improper registration, and motion blurring have been recognized as limitations for contouring. MRI provides excellent soft tissue contrast, and higher image resolution than PET, but has seen limited use in target definition for lung cancer radiotherapy [19]. Hence, radiation oncologists have generally less experience with MRI than CT for target definition. The growing use of MRI in

radiation oncology departments for treatment planning, image-guided adaptive radiotherapy, and treatment response assessment motivated us to compare the contouring variability for lung cancer on MRI to routine methods.

We selected the particular MRI sequences for several reasons. VIBE images provide anatomic details based on the different T1-relaxation values of water protons in tissues, can be acquired with sufficiently high spatial resolution, and motion artifact is minimized through breath hold acquisition. Analogous to PET-CT, incorporating functional imaging such as DW-MRI along with morphological VIBE images may assist in differentiating benign and cancerous tissue during delineation (e.g. within atelectatic lung). Variations in the apparent diffusion of water molecules due to different tissue cellularity provide contrast between tumor and the surrounding tissues in DW-MRI images and ADC maps [19].

Despite the time intervals described above for obtaining the different imaging modalities with CT acquired first, the overall tumor volume on CT only was the largest, whereas contouring on PET-CT resulted in the smallest average GTV\_PT volume. Similarly, contour variability for the whole tumor surface was largest for CT only, indicating that most of the volume differences are caused by the inability to determine the tumor boundary. However, four of the seven observers on average drew larger contours on PET-CT than CT only, potentially due to the blurring effect on PET-CT, and three observers were found to have their smallest average contours on MRI.

The ability to differentiate tumor from surrounding structures depends on tumor-specific factors such as tumor shape, size, location and its surrounding structures. Our results show that the variability of all GTV\_PT interfaces was smallest on PET-CT except for the interface with lung where CT contouring had the smallest variability. Functional information on PET-CT allowed good identification of collapsed lung, however, contouring variability was significantly larger on the tumor-atelectasis interface for all modalities compared to other interfaces. This is in agreement with the analysis by Steenbakkers et al [23]. MRI is currently used in routine diagnostic imaging to examine mediastinal, vascular and nerve invasion. In our study, MRI did not result in any significant reductions in delineation uncertainty, although some benefit was seen at the interfaces between GTV\_PT and mediastinum and GTV\_LN and vessels. Overall, MRI did not appear helpful for GTV\_LN interface detection in this small cohort. However, GTV\_LN volumes and interfaces were small. Even small contour differences therefore resulted in large errors relative to the median contour.

Our results indicate that some observers consistently drew larger/smaller volumes on all imaging modalities. The causes of this might be multifactorial. Observer training had been shown to reduce contouring variability [40, 41]. In this study, for each patient, graphical information on the location of the primary tumor and lymph nodes of interest was provided to minimize contouring errors due to uncertainty about tumor position and to focus the study on the detection of uncertainties arising from differences in imaging modalities only.

Technical factors such as slice thickness, resolution and signal-to-noise ratio (SNR) can also affect the differentiation of tumor from its surrounding structures. While VIBE offers

excellent soft tissue contrast, the low resolution of DW-MRI images and motion artifacts in the acquisition of VIBE and DW-MRI images may have potentially impacted contouring variability. Optimization of these factors may result in greater contrast-to-noise ratio (CNR) and may reduce delineation variability. Special attention should be given to artifacts/ distortions such as motion, susceptibility and chemical shift artifacts associated with different types of MRI, particularly during delineation of chest tumors.

## **V. CONCLUSIONS**

While MRI-based contouring showed overall larger variability than PET-CT, contouring variability depended on the interface type and was not significantly different between modalities despite of the limited observer experience with MRI. The GTV\_PT-atelectasis interface had the largest uncertainty for all imaging modalities. With more training and experience, anatomic MRI combined with functional MRI sequences might gain clinical importance for tumor delineation close to critical interfaces, such as for differentiating cancerous processes in atelectatic lung from inflammation where PET-CT has known limitations. Although MRI showed less variability for the GTV\_LN-vessel interface, larger contour variations and volumes for GTV\_LN in general indicate that MRI can currently not be recommended for GTV\_LN contouring with the MRI sequences used in this study. Multimodality imaging and combining different imaging characteristics might be the best approach to define the GTV most accurately.

#### Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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#### Summary

Target delineation in lung cancer radiotherapy is affected by large variability. MRI has excellent soft tissue visualization and was therefore compared to CT only and PET-CT for lung cancer target definition in a multiple observer delineation study. Contouring variability on MRI was usually less than CT, but greater than PET-CT without significant differences. Contouring variability was also observed dependent on the interface type with the largest variability identified for the tumor-atelectasis interface.



#### Figure 1.

An example of a large primary tumor GTV delineated by seven observers (different thin colored lines) and median contour (dark blue colored line) in different imaging modalities (for reference see also Figure S1 without contours). DW-MRI images and the corresponding ADC maps of a frame covered ~5.0 cm thickness. The surface mesh of local SD (SDSM) for the three modalities in an oblique view with the directions is shown. In total, three frames of DW-MRI and ADC maps were used to cover this tumor. The contoured volumes were 346, 282 and 328 cm<sup>3</sup> on CT, PET-CT and MRI, respectively.



#### Figure 2.

The interfaces of primary tumor GTV with lung (dark-blue), hilum (light blue), atelectasis (orange), chest-wall (green) and mediastinum (dark red) on median contour surfaces of PET-CT of an example patient in four representative axial slices (A–D), and in a 3D oblique (E) and its flipped views (F).



#### Figure 3.

CT and MRI based primary tumor (A) and lymph node (B) gross tumor volumes (GTVs) normalized by dividing average volume per patient by the corresponding PET-CT based volume. Boxes represent the inter-quartile range (IQR) extending from first quartile (Q1) to third quartile (Q3), respectively. Whiskers extend from Q1 and Q3 to end points that are typically defined as the most extreme data points within Q1-1.5\*IQR and Q3 +1.5\*IQR, respectively. The small open circles represent the individual data points. Fig. 3(A) shows that the normalized CT-based GTV\_PT volume was generally larger than the MRI-based volume, both volumes were larger than tumor volumes on PET-CT.



#### Figure 4.

Individual observers' contour volumes (mean±SD) normalized relative to the corresponding mean volume per patient and averaged over all patients for the gross tumor volumes (GTV) of primary tumor (PT) (A) and lymph node (LN) (B) per imaging modality. Some observers consistently delineated comparatively large or small volumes. Volumes delineated by radiologists (Observer#3 and Observer#5) were relatively small compared to radiation oncologists.

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#### Figure 5.

Overall uncertainty (RMS SD) of BLD values (mean±SD) for the gross tumor volumes (GTVs) of primary tumor (PT) interfaces (A) and lymph node (LN) interfaces (B). Uncertainties were largest for the PT- atelectasis interface for all modalities.

atient	Age (y)	Race	Sex	Histology	Stage	<b>Tumor Location</b>	Mean GTV_PT Volume on PET-CT (in ml)	SUV <sub>max</sub> of GTV_PT
A	57	AA	ц	SCC	IIIB	RUL	231.4	28.8
в	61	M	Μ	SCC	IIIB	TUL	50.4	24.3
C	62	W	ц	Adeno	ШA	RUL	95.7	11.8
D	57	M	Μ	SCC	ШA	RUL	3.6	18.1
ш	60	M	ц	Adeno	IIIB	TUL	117.4	32.9
ц	50	W	Μ	SCC	ША	RUL	107.8	11.1
IJ	56	AA	Μ	SCC	IIIB	RUL	284.9	63
Н	51	AA	Μ	SCC	IIB	RUL/RML	58.7	66
Ι	64	AA	Μ	Adeno	IIIB	TTT	78.2	17.1
ſ	53	AA	М	Undiff. Ca	ША	LN only		41

AA, African-American; Adeno, adenocarcinoma; Ca, carcinoma; F, female; GTV\_PT, gross target volume of primary tumor; LLL, left-lower lobe; LN, lymph nodes; LUL, left-upper lobe; M, male; RML, right-middle lobe; RUL, right-upper lobe; SCC, squamous cell carcinoma; Undiff, undifferentiated; W, white. All patients except C, D and J had atelectasis or collapsed lung.

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Table 1