Motion Correction of $^{90}$Y Dose Maps with PET/MR Imaging

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Introduction

Yttrium-$^{90}$ radioembolization is a therapeutic procedure that delivers local radiation to hepatic tumors [1]. Clinically, patients undergo a $^{90}$Y Bremsstrahlung SPECT scan to determine if there was any extrahepatic deposition and, most importantly, to predict if the tumor is likely to respond to therapy based on the level of the delivered dose [2]. In addition, it is also possible to image and quantify the delivered $^{90}$Y dose by PET, which was shown to be superior to Bremsstrahlung SPECT in terms of spatial resolution and quantification accuracy [3]. Clinical SPECT and PET scanners are currently integrated with CT systems to enable localization as well as attenuation correction of the detected emission activity signal from the CT anatomical transmission signal [4]. However, clinical PET/CT studies have shown that the relative difference in dose between responding and non-responding lesions may be as low as 25%, thus demonstrating the importance of precise anatomical localization and high quantitative accuracy when assessing dose deposition [5]. Localization of the $^{90}$Y signal distribution in the liver can be challenging with PET/CT, mainly due to the poor soft tissue discrimination and limited motion tracking capabilities offered by CT.

However, the recent advent of integrated PET/MR systems in clinic, supporting the simultaneous acquisition of PET and MR data, has enabled the automatic and highly accurate spatiotemporal co-registration of metabolic PET with anatomical and functional MR images [6]. The novel hybrid PET/MR technology may thus offer significant clinical benefits in $^{90}$Y dose imaging assessments over PET/CT. First, MRI is associated with considerably better soft tissue constant therefore permitting the more accurate drawing of ROIs on the MR image when evaluating $^{90}$Y PET regional assessments (Fig. 1). Second, the superior soft tissue resolution and absence of radiation exposure of MRI allows for more accurate tracking of the respiratory motion for improved PET motion correction. Thus, PET/MR can considerably enhance quantification in $^{90}$Y dose imaging and therefore potentially improve therapeutic efficiency in clinic. Indeed, recent PET/MR studies have indicated a stronger relationship between tumor response and delivered dose even in the absence of motion correction [7]. In this study we are targeting the optimization of clinical $^{90}$Y PET/MR imaging with a particular focus on MR-based motion correction of the $^{90}$Y dose maps using the Biograph mMR integrated PET/MR system [8].

Fig. 1. MRI (1A), PET (1B), and fused PET/MRI (1C) images of a subject who underwent $^{90}$Y radioembolization. Arrows point to the lesion in the MR image (COR-HASTE) and the PET image.
Motion tracking and correction strategy for simultaneous $^{90}$Y-PET/MR imaging

The acquisition of anatomical MR signal of high spatial and temporal resolution allows for high temporal sampling rates of detailed 3D respiratory 3D cartesian motion vector field (MVF) estimates. In addition, the simultaneous acquisition of $^{90}$Y PET data permits their synchronization with the MR-based respiratory motion phase for the accurate respiratory gating of the PET data. Finally, the gated PET data and the MVF are imported into a 4D PET motion-compensated image reconstruction (MCIR) algorithm to directly generate the motion-corrected $^{90}$Y PET dose maps.

In this study, we exploit MR-based motion correction capabilities of the Biograph mMR system to assess the improvement in the quantitative accuracy of the $^{90}$Y dose distribution assessments by reducing the respiratory motion blurring effect in the final PET reconstructed images. The lesions are often in the top region of the liver, which could move up to 2 cm due to respiration [9]. Therefore, our main goals are to:

1) Develop an optimal data acquisition and reconstruction scheme specially tuned for $^{90}$Y imaging post radioembolization on the Siemens Biograph mMR; and

2) Evaluate the Biograph mMR motion correction algorithm (software version syngo MR E11p).

Previously, we conducted a preliminary evaluation of the MCIR algorithm performance on $^{90}$Y phantom studies [10, 11]. Currently, we expand our validation on patient data to optimize the motion-compensated reconstruction parameters in the clinic.

Our current protocol at Mount Sinai is outlined in Figure 2. The total scan time ranges between 30 and 35 minutes. Currently we run a prototype motion tracking sequence (Siemens BodyCOMPASS) for the entire duration of the PET scan. The sequence permits the generation of a set of high resolution 3D MR gated images, namely a 4D MR image, each corresponding to a different phase of the respiratory cycle, from end-expiration to end-inspiration. Subsequently, standard image registration methods are used to calculate from the gated MR images the 3D non-rigid motion transformation maps, which constitute the estimated motion model. In addition, the same MR data can be utilized to track the trace of the respiratory motion throughout the PET acquisition. This trace can be later employed to sort the synchronized PET data into the same set of respiratory gates. After the completion of the MR tracking sequence, we acquire additional MRI data with sequences designed for high-resolution anatomical static imaging to facilitate the accurate MR-based region-of-interest (ROI) localization in the $^{90}$Y PET dose maps. Current MRI sequences in the exam are: 1) Axial HASTE, 2) Coronal HASTE, 3) 3D Dixon, 4) Axial T1w. We find HASTE images to be best for that purpose; however, contrast-enhanced MRI is used by other groups and its use should be investigated in the future. It is important to note that for $^{90}$Y imaging it would be ideal to find only one sequence to be used for ROI definition in order to minimize scan time as much as possible. As a consequence, would then be feasible to dedicate the entire PET scan for motion tracking if needed.

Figure 3 shows sample sagittal images showing the various phases. The sequence as well as the data sorting algorithm seems to perform well in resolving motion.

Examples of 3 different MR gated images corresponding to respective phases of the respiratory motion cycle, as generated with MR motion tracking sequence included in Biograph mMR syngo MR E11p software.
Enhancing ¹⁰⁹Y dose maps quantification with motion-compensated PET/MR imaging

Figure 4A illustrates a clear visual improvement in resolution and contrast of ¹⁰⁹Y dose maps after application of motion correction within the PET reconstruction. Motion corrected ¹⁰⁹Y PET images (right) are characterized by higher signal contrast recovery compared to the respective images without motion correction, i.e. static images (left). Moreover, the motion-corrected ¹⁰⁹Y images are associated with superior signal-to-noise ratio (SNR) compared to the gated ¹⁰⁹Y image (middle). The line plot in Figure 4B quantitatively confirms the improved contrast recovery for ¹⁰⁹Y dose maps when motion correction is applied. The degree of ¹⁰⁹Y contrast enhancement in the liver would be expected to reach maximum score levels for lesions located in the top of the liver, at the liver-lung interface. This is attributed to the strongest resolution degradation effects often observed in the liver-lung interface due to respiratory motion-induced contamination of the liver ¹⁰⁹Y uptake signal with the considerably smaller background signal from the lung. Indeed, the alignment of the ¹⁰⁹Y dose with the MR anatomic map before and after motion correction in Figure 5 illustrates the automatic correction of the position of the ¹⁰⁹Y deposited dose distribution within the liver after motion correction. This is of high importance in clinical practice, as occasionally a percentage of ¹⁰⁹Y activity may be observed in lungs due to air embolization [12].

Furthermore, in Figure 6 more clinical cases are presented where reconstructed ¹⁰⁹Y dose maps have been benefited from MR-based PET motion correction. The contrast recovery enhancement of motion-corrected versus static PET images is visually evident in focal ¹⁰⁹Y uptake regions.

Moreover, in some clinical cases, no attachment to the target was observed for the delivered ¹⁰⁹Y dose thus resulting in diffused ¹⁰⁹Y distribution as shown in Figure 7.

Nevertheless, the Biograph mMR syngo MR E11p motion correction algorithm did not induce any artifacts or false positives.

Clinical prospects in motion-compensated ¹⁰⁹Y-PET/MR imaging

Our preliminary findings in a few patients show that MR based motion correction for ¹⁰⁹Y could improve the quantitative accuracy of the data. As mentioned above, the literature indicates a difference between responding and non-responding lesions of 25%, and thus the margin for error is quite small. The effect of motion, especially at the top of the liver, could be higher than that margin and thus its use could be significant. To accurately measure the improvement, a cohort of about 20-30 subjects is desirable to show the potential benefits of motion correction. There are some attenuation correction issues including lack of a lung segment (i.e. LAC = 0) in some of our cases which require further evaluation. We have been using the motion correction sequence using the default parameters and this might require some optimization. Moreover, optimization, streamlining, and integration of motion correction into routine reconstruction are needed. Finally, the best number of gates and the navigator signal from the belt should be further investigated.
Clinical 90Y PET images without (static) and with motion correction for 2 clinical cases characterized by absence of specific binding of 90Y to the liver tissue.

References


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