Tumor-site specific geometric distortions in high field integrated magnetic resonance linear accelerator radiotherapy

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1. Introduction

The goal of radiotherapy (RT) is to deliver a high radiation dose to the tumor while sparing surrounding organs at risk (OAR). This is possible with advanced techniques such as intensity modulated RT, volumetric modulated arc therapy, and stereotactic body RT. Image guided RT (IGRT) is used to ensure precise dose delivery \cite{1} and is generally achieved by using cone-beam CT at each treatment fraction, but these images suffer from poor soft-tissue contrast. Magnetic resonance imaging (MRI) has superior soft-tissue contrast and enables visualization of the tumor and OAR, while eliminating ionizing radiation exposure from imaging. MRI based IGRT is made possible with integrated MRI linear accelerators (MR-linac) \cite{2,3}.

The MR-linac enables daily in-room adaption of RT treatment plans based on an anatomy-of-the-day strategy acquired with the integrated MRI \cite{4,5}. Fast MRI sequences enable real time imaging during beam-on which allows intra-fractional motion monitoring of the tumor or OAR, and potentially gated treatment delivery \cite{5,6}. Therefore, IGRT not only requires clear visualization of relevant anatomical structures but also a valid geometrical representation of imaged structures to ensure precise dose delivery. Geometric distortion of images is however a well-known problem in MRI \cite{7,8}, and is a major concern when MRI is to be used for guidance of RT.

Geometric accuracy can be impairedb both by the MRI system and the patient. The important MRI system related sources of distortion are non-linearity of the spatial encoding gradients (gradient non-linearity, GNL) and inhomogeneity of the main magnetic field (B_0) \cite{8}. Patient-induced geometric distortions arise due to differences in the magnetic susceptibilities of anatomical structures (e.g. air cavities and tissue boundaries) distorting the local B_0 field \cite{9}. Geometric distortion in MR-linacs has been studied by others \cite{9–15}, i.e. in the study by Tijssen et al. where a method for cumulative total geometric distortion is presented \cite{10}, but larger sample sizes and anatomical tumor-site specific investigation is still warranted. The aim of this study was to investigate the total geometric distortion in a larger sample of patients grouped into anatomical tumor-sites. The total geometric distortion was assessed for clinical scans used in an MRI guided daily adaptive workflow with a 1.5 T MR-linac system. Four different tumor-sites (adrenal gland, liver, pancreas, and prostate) have been evaluated.

2. Materials and methods

2.1. Patients

For this study 37 patients undergoing treatment on the MR-linac were included prospectively. All patients were included in feasibility protocols \cite{16,17} and have given their signed consent to participate. Tumor-sites were: adrenal gland in 7 patients, liver in 4 patients, pancreas in 6 patients, and prostate in 20 patients.
2.2. MR-linac

All imaging acquisitions were performed on the 1.5 T MR-linac (Unity, Elekta AB, Stockholm, Sweden). A complete user-oriented MR commissioning was performed initially [12] on the MR-linac, including test of overall B0-homogeneity, linac gantry rotation dependent distortions, distortions while beam-on and potential eddy currents effects. These effects were found to be negligible.

2.3. B0-mapping

B0-mapping MRI scans were acquired on the MR-linac. Patients were scanned in treatment position during beam-on at their first treatment fraction. For few cases the B0 mapping scan was acquired immediately after treatment delivery if motion monitoring images were acquired during the treatment delivery. A 3D gradient echo (GRE) dual-echo MRI sequence was used (TE1/TE2: 4.6/9.2 ms, TR: 11.4 ms, flip angle: 30°).

| Table 1 | Geometric distortions for four tumor-sites. B0 related distortions and total geometrical distortion (consisting of B0 and gradient non-linearity (GNL) related distortions) are reported. Median distortion and maximum (defined in section 2.6 Statistics) are reported. The sign of the B0 related distortion reflects the direction of the distortion in the frequency-encoding direction (anterior-posterior direction on images in Fig. 1a–d). The B0 related distortions are reported for a clinical 3D T2W MRI sequence (specifications can be seen in section 2.3 B0-mapping). Statistical significant difference between the pooled abdominal tumor-sites (adrenal gland, liver, and prostate) and the pelvic tumor-site (prostate) is denoted by * next to the calculated p-values. |
| --- | --- | --- | --- | --- | --- | --- | --- |
| | GTV | GTV + 30 mm | GTV | GTV + 30 mm |
| | Median [mm] | Max [mm] | Median [mm] | Max [mm] | Median [mm] | Max [mm] | Median [mm] | Max [mm] |
| **Adrenal gland** | | | | | | | | |
| 1 | 0.01 | 0.08 | 0.01 | -0.06 | 0.09 | 0.23 | 0.09 | 0.44 |
| 2 | -0.01 | 0.22 | 0.02 | 0.10 | 0.05 | 0.26 | 0.09 | 0.28 |
| 3 | 0.00 | 0.24 | -0.01 | 0.21 | 0.08 | 0.27 | 0.10 | 0.32 |
| 4 | 0.08 | 0.13 | 0.07 | 0.12 | 0.09 | 0.21 | 0.12 | 0.28 |
| 5 | -0.02 | 0.12 | -0.01 | 0.10 | 0.13 | 0.27 | 0.11 | 0.33 |
| 6 | 0.04 | 0.07 | 0.03 | 0.07 | 0.12 | 0.31 | 0.11 | 0.28 |
| 7 | -0.02 | 0.17 | 0.01 | 0.10 | 0.26 | 0.48 | 0.28 | 0.75 |
| **Mean** | 0.02 | 0.15 | 0.02 | 0.09 | 0.12 | 0.29 | 0.13 | 0.38 |
| **Liver** | | | | | | | | |
| 1 | 0.08 | 0.13 | 0.07 | 0.16 | 0.21 | 0.26 | 0.20 | 0.52 |
| 2 | 0.04 | 0.08 | 0.05 | 0.08 | 0.57 | 0.67 | 0.54 | 1.02 |
| 3 | 0.02 | 0.10 | 0.03 | 0.10 | 0.09 | 0.11 | 0.08 | 0.31 |
| 4 | 0.02 | 0.15 | 0.01 | 0.13 | 0.44 | 0.96 | 0.33 | 1.27 |
| **Mean** | 0.04 | 0.12 | 0.04 | 0.12 | 0.33 | 0.50 | 0.29 | 0.78 |
| **Pancreas** | | | | | | | | |
| 1 | -0.01 | -0.05 | 0.01 | 0.07 | 0.05 | 0.12 | 0.13 | 0.28 |
| 2 | -0.03 | -0.08 | -0.02 | -0.10 | 0.06 | 0.14 | 0.13 | 0.26 |
| 3 | -0.01 | -0.05 | -0.02 | -0.07 | 0.20 | 0.28 | 0.17 | 0.31 |
| 4 | -0.03 | 0.17 | -0.01 | 0.14 | 0.20 | 0.30 | 0.18 | 0.37 |
| 5 | -0.01 | -0.08 | 0.00 | 0.07 | 0.20 | 0.28 | 0.16 | 0.27 |
| 6 | -0.03 | -0.07 | -0.01 | 0.09 | 0.13 | 0.24 | 0.14 | 0.28 |
| **Mean** | -0.02 | -0.03 | -0.01 | 0.03 | 0.14 | 0.23 | 0.15 | 0.29 |
| **Prostate** | | | | | | | | |
| 1 | -0.05 | -0.10 | -0.04 | -0.14 | 0.13 | 0.27 | 0.12 | 0.26 |
| 2 | -0.06 | -0.10 | -0.04 | -0.09 | 0.22 | 0.34 | 0.11 | 0.26 |
| 3 | -0.03 | -0.07 | -0.02 | -0.08 | 0.18 | 0.28 | 0.10 | 0.26 |
| 4 | -0.01 | -0.05 | -0.01 | -0.06 | 0.20 | 0.28 | 0.10 | 0.24 |
| 5 | -0.01 | -0.04 | 0.01 | 0.04 | 0.18 | 0.26 | 0.08 | 0.21 |
| 6 | -0.01 | -0.07 | -0.01 | 0.07 | 0.18 | 0.25 | 0.10 | 0.24 |
| 7 | -0.03 | -0.08 | -0.02 | -0.07 | 0.24 | 0.30 | 0.10 | 0.25 |
| 8 | -0.04 | -0.07 | -0.02 | -0.08 | 0.16 | 0.32 | 0.10 | 0.25 |
| 9 | -0.03 | -0.09 | -0.01 | -0.06 | 0.21 | 0.32 | 0.09 | 0.25 |
| 10 | -0.05 | -0.09 | -0.03 | -0.07 | 0.19 | 0.25 | 0.09 | 0.25 |
| 11 | -0.02 | -0.05 | -0.01 | -0.06 | 0.18 | 0.28 | 0.10 | 0.25 |
| 12 | -0.03 | -0.08 | -0.01 | -0.07 | 0.18 | 0.28 | 0.09 | 0.24 |
| 13 | -0.04 | -0.09 | -0.04 | -0.08 | 0.20 | 0.31 | 0.10 | 0.27 |
| 14 | 0.00 | -0.05 | 0.00 | -0.05 | 0.19 | 0.29 | 0.10 | 0.24 |
| 15 | -0.03 | -0.07 | -0.01 | -0.06 | 0.20 | 0.30 | 0.09 | 0.25 |
| 16 | -0.04 | -0.08 | -0.02 | -0.07 | 0.10 | 0.21 | 0.13 | 0.27 |
| 17 | -0.03 | -0.08 | -0.01 | -0.06 | 0.17 | 0.26 | 0.09 | 0.25 |
| 18 | -0.03 | -0.07 | -0.01 | -0.09 | 0.15 | 0.27 | 0.09 | 0.24 |
| 19 | -0.03 | -0.08 | -0.02 | -0.08 | 0.17 | 0.26 | 0.09 | 0.24 |
| 20 | -0.07 | -0.12 | -0.03 | -0.11 | 0.19 | 0.30 | 0.10 | 0.27 |
| **Mean** | -0.03 | -0.07 | -0.02 | -0.06 | 0.18 | 0.29 | 0.10 | 0.25 |
| **P-value** | < 0.001* | < 0.001* | < 0.001* | < 0.001* | 0.20 | 0.22 | 0.0013* | < 0.001* |
with echo times selected to ensure in-phase water-fat-signal to avoid signal cancellation. Magnitude and phase images were acquired with a matrix size of $349 \times 351 \times 229$, readout BW (rBW) of $740$ Hz/pixel, and 2 averages resulting in a reconstructed resolution of $0.75 \times 0.75 \times 2$ mm$^3$, and reconstructed matrix of $560 \times 560 \times 229$. Sensitivity encoding (SENSE) was used with reduction factors of 1.5 (frequency encoding (FE)) and 2 (phase encoding (PE)). Percent sampling of the k-space was 78%. Acquisition duration was 406 s. For all acquisitions FE was anterior-posterior (AP) direction, in-plane PE was left-right (LR) direction, and out-of-plane PE was head-feet (HF) direction.

Phase images were unwrapped in 3D using a Laplacian unwrapping algorithm in MATLAB [18,19] and an axial through-tumor slice was used for calculating $B_0$ maps. $B_0$ maps were calculated from the unwrapped phase images with in-house developed software in MATLAB using the dual-echo method [20] where the $B_0$ variation ($\Delta B_0$) is found from the phase differences ($\Delta \phi$): $\Delta B_0 = \Delta \phi / (2 \gamma \Delta TE)$, where $\gamma$ is the gyromagnetic ratio. The $B_0$ maps were converted to spatial distortion maps using the rBW (693 Hz/pixel) and the pixel size ($1.2 \times 1.2 \times 2$ mm$^3$) of a clinical 3D T2W spin-echo (SE) sequence for the abdominal/pelvic region.

2.4. GNL mapping

To map the residual GNL related distortion (after 3D system correction) a vendor provided phantom was used (Elekta AB, Stockholm, Sweden). The phantom consisted of 7 slices and each slice contained 274 circular markers located $25 \times 25$ mm apart. GNL was calculated in 2D (central axial plane) using the gradient reversal method [3] (TE: 3.4 ms, TR: 6.7 ms, flip angle: 15°). Magnitude images were acquired with a matrix size of $372 \times 374 \times 200$ and a rBW of 461 Hz/pixel resulting in a reconstructed matrix size of $512 \times 512 \times 400$, and a reconstructed resolution of $1.09 \times 1.09 \times 2$ mm$^3$. The gradient reversal method enabled separation of distortions due to GNL from $B_0$ induced distortions and a 2D GNL distortion map was created for the central axial slice of the MR-linac which corresponded to the position of the tumor-sites investigated. The GNL distortion map was considered constant and equal in all patients.

2.5. Total distortion

A total distortion map was created for each patient adding the GNL distortion and $B_0$ distortion contributions. In the FE direction the $B_0$ related distortion was added to the GNL related distortion. In the PE direction only contribution from GNL existed. Total distortion was found as the magnitude of the added distortion vectors in the FE and PE direction. The median and maximum distortions were reported for the gross tumor volume (GTV) and an expanded region (GTV + 30 mm margin). To remove effects of noise the maximum distortion was reported as the 95-percentile (or 5-percentile if negative) distortion.

2.6. Statistics

Differences in median and maximum distortion between the pooled group of abdominal tumor-sites (adrenal gland, liver, pancreas: 17 patients) and the pelvic tumor-site (prostate: 20 patients) was tested using a Wilcoxon rank-sum test (with $\alpha = 5\%$).

3. Results

The maximum $B_0$ related distortions inside the GTVs were < 0.24 mm for adrenal gland (range 0.07 – 0.24 mm in seven patients), < 0.15 mm for liver (range 0.08 – 0.15 mm in four patients), < 0.17 mm for pancreases (range −0.08 – 0.17 mm in six patients), and < |−0.12| mm for prostate (range −0.04 – −0.12 mm in 20 patients) (Table 1 and Fig. 1a–d). All results including results for the median values as well as the expanded region can be seen in Table 1. Negative distortions corresponded to a distortion in the anterior direction and positive distortions corresponded to a distortion in the posterior direction. Significant difference (p-values can be seen in Table 1) was seen between the abdominal tumor-sites and the pelvic tumor-site for both median and maximum $B_0$ related distortion and for both the GTV and expanded region. Difference was due to the direction of the distortion rather than magnitude.

The influence of the systematic GNL distortion depended on the distance of the tumor from the iso-center (Fig. 1e). Maximum total distortion was $< 0.48$ mm for adrenal gland, $< 0.96$ mm for liver, $< 0.30$ mm for pancreas, and $< 0.32$ mm for prostate for the GTVs. For the expanded region the largest distortion was 1.27 mm (liver, patient 4). No significant difference was seen when comparing median and
maximum values for total distortion in GTVs of abdominal tumor-sites and pelvic tumor-sites. When expanding the regions (GTV > 30 mm) significant difference was seen for median and maximum total distortion when comparing abdominal tumor-sites and pelvic tumor-sites.

4. Discussion

The contribution from the B0 related distortion was generally low, as the highest contribution among the 37 patients was 0.24 mm (adrenal gland, patient 3). Patient-induced distortions in the pelvic region have previously been reported [2,22] for different field strengths and receiver bandwidths, which makes direct comparison difficult. Stanescu et al. [9], Gide-Hurst et al. [21] and Tyagi et al. [22] reported values in the same magnitude for tumors/anatomies in the pelvis and prostate. Tijssen et al. reported distortions below 1.5 mm in the abdominal region [10]. As field strength, rBW and pixel size have direct influence on the B0 related spatial distortion, these parameters need careful consideration in MR guided RT [23].

The median total distortion was not significantly different for the pelvic tumor-site (prostate) compared to the abdominal tumor-sites (adrenal gland, liver, and pancreas). Common for prostate patients the GTV was located within a radius of 50 mm from the MRI iso-center and the contribution from GNL distortion was low (Fig. 1e). The pancreatic tumors were also located near the iso-center, whereas adrenal gland tumors and liver metastasis were located further from the iso-center, which was reflected in a higher GNL related distortion and thereby total distortion. The GNL related distortions were in agreement with results by Kemppainen et al. [24]. They reported maximum GNL distortion below 2 mm in most patients, where the tumor-site was close to the iso-center. The largest GNL related distortions in our study was found in two patients with liver metastases (liver, patient 3 and 4, Table 1), at comparable distance to the iso-center. This was in agreement with general deterioration of linearity of the imaging gradient when moving from the iso-center. Liver metastases located farther from iso-center could be affected by GNL related distortions > 2 mm in the MR-linac (Fig. 1e). The GNL contribution was implemented as 2D in-plane in this study. This potentially underestimated the GNL, however, since targets in patients were aligned with the central axial plane the estimation of distortion due to GNL was expected to be sufficiently accurate.

The level of total geometric distortion could be clinically relevant for some of the patients presented, as distortions > 1 mm were seen. Intra-patient variation over time needs to be investigated for the patient-specific B0 maps to clarify the influence of e.g. patient positioning and air, which could also clarify the usability of patient-specific B0 maps. The intra-patient variation is a crucial indication for whether B0 maps should be acquired for each treatment fraction to be relevant, or if a single map at the first treatment is sufficient. To address the consequences of the distortion level for the treatment the dosimetric effect should also be evaluated [25].

In conclusion, the maximum total geometric distortion seen in the GTV was 0.96 mm (liver, patient 4) with no significant difference between abdominal and pelvic patients. The maximum contribution of B0 related distortion in the GTV was 0.24 mm (adrenal gland, patient 3) and the B0 related distortion was significantly different in pelvic (prostate) and abdominal tumor-sites (adrenal gland, liver, pancreas). The main source of distortion was due to gradient non-linearity (GNL), and the total distortion level increased as the distance to the MRI iso-center increased. Distortions due to GNL should ideally be taken into consideration if the tumor-site is far off-center in a MR based adaptive RT workflow.

Declaration of Competing Interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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