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4

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6 Real-time CT image generation

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26 **Abstract**

27 *Purpose*

28 To show the feasibility of real-time CT image generation technique utilizing internal fiducial markers that facilitate
29 the evaluation of internal deformation.

30 *Methods*

31 In the proposed method, linear regression model that can derive internal deformation from the displacement of
32 fiducial markers is built for each voxel in the training process before the treatment session. Marker displacement
33 and internal deformation are derived from the four-dimensional computed tomography (4DCT) dataset. In the
34 treatment session, the three-dimensional deformation vector field is derived according to the marker displacement,
35 which is monitored by the real-time imaging system. The whole CT image can be synthesized by deforming the
36 reference CT image with a deformation vector field in real-time. To show the feasibility of the technique, image
37 synthesis accuracy, and tumor localization accuracy were evaluated using dataset generated by extended NURBS-
38 Based Cardiac-Torso (XCAT) phantom and clinical 4DCT datasets from six patients, containing ten CT datasets
39 each. In the validation with XCAT phantom, motion range of the tumor in training data and validation data were
40 about 10 and 15 mm, respectively, so as to simulate motion variation between 4DCT acquisition and treatment
41 session. In the validation with patient 4DCT dataset, eight CT datasets from the 4DCT dataset were used in the
42 training process. Two excluded inhale CT datasets can be regarded as the datasets with large deformations more
43 than training dataset. CT images were generated for each respiratory phase using the corresponding marker
44 displacement. Root mean squared error (RMSE), normalized RMSE (NRMSE) and structural similarity index
45 measure (SSIM) between the original CT images and the synthesized CT images were evaluated as the quantitative
46 indices of the accuracy of image synthesis. The accuracy of tumor localization was also evaluated.

47 *Results*

48 In the validation with XCAT phantom, the mean NRMSE, SSIM and three-dimensional tumor localization error
49 were $7.5\pm 1.1\%$, 0.95 ± 0.02 and 0.4 ± 0.3 mm, respectively. In the validation with patient 4DCT dataset, the mean
50 RMSE, NRMSE, SSIM and three-dimensional tumor localization error in six patients were 73.7 ± 19.6 HU,
51 $9.2\pm 2.6\%$, 0.88 ± 0.04 and 0.8 ± 0.6 mm, respectively. These results suggest that the accuracy of the proposed
52 technique is adequate when the respiratory motion is within the range of the training dataset. In the evaluation with
53 a marker displacement larger than that of the training dataset, the mean RMSE, NRMSE, and tumor localization
54 error were about 100 HU, 13%, and less than 2.0 mm, respectively, except for one case having large motion variation.
55 The performance of the proposed method was similar to those of previous studies. Processing time to generate the
56 volumetric image was less than 100 ms.

57 *Conclusion*

58 We have shown the feasibility of the real-time CT image generation technique for volumetric imaging.

59

60 **Keywords:** CT image generation, synthetic CT, volumetric imaging, motion management, fiducial markers, partial
61 least squares regression

62 **Introduction**

63 Respiratory motion is a major source of dosimetric error in radiation therapy. Respiratory-gated irradiation using
64 an external surrogate, such as abdominal motion, has been clinically applied for motion management.¹⁻⁵ In addition,
65 real-time tumor-tracking radiation therapy (RTRT)⁶, which monitors the internal tumor motion directly, has been
66 applied in lung^{7,8}, liver⁹ and pancreas. In RTRT, the three-dimensional position of internal fiducial markers, which
67 are inserted near the tumor, are monitored in real-time by means of orthogonal x-ray imaging during treatment.
68 Treatment beam is irradiated only when the fiducial marker is within the gating window, which is cubic region of
69 ± 2 mm in typical clinical setting. RTRT has been also clinically applied in proton therapy^{10,11}. Tracking irradiation
70 techniques that change the irradiation position/field according to the tumor position by using electromagnetic
71 transponder guided multileaf collimator (MLC)¹², a compact linear accelerator mounted on a robotic arm¹³ and the
72 gimbaled x-ray head¹⁴ have also been realized in photon therapy. However, for accurate irradiation, especially in
73 particle therapy and stereotactic body radiation therapy (SBRT), both the tumor position and volumetric information
74 of the surrounding organs and tissue are needed for real-time motion management.

75 Real-time volumetric imaging, which can reconstruct anatomical structures, could be applied for a novel motion
76 management technique. Volumetric imaging, based on the estimation of the coefficients of the principal components
77 of internal deformation from a planar x-ray image, has been studied.^{15,16} Other approaches based on deep learning
78 have also been reported.¹⁷⁻¹⁹ Previous studies demonstrated that clinically acceptable accuracy of image synthesis
79 and tumor localization can be obtained. However, imaging accuracy may depend on imaging angle and quality.¹⁹

80 In this study, we propose a novel real-time CT image generation technique for volumetric imaging by utilizing the
81 positional displacement of internal fiducial markers evaluated in real-time during treatment. In RTRT using an
82 orthogonal x-ray imaging system, the three-dimensional position of the fiducial markers can be obtained with a high
83 spatial accuracy of less than 1 mm and a high temporal resolution up to 30 times per second. Accuracy does not
84 depend as much on image quality of planar x-rays, as long as the marker can be recognized correctly.²⁰ In addition,
85 doses due to x-ray imaging can be suppressed by collimating the imaging area using only the regions of the fiducial

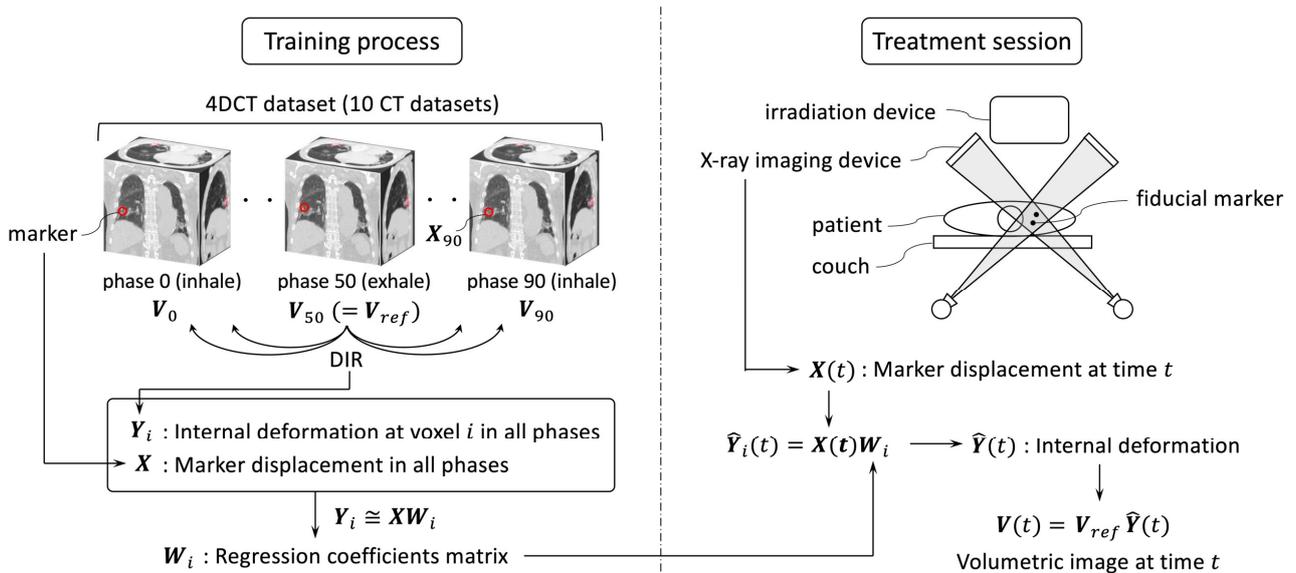
86 markers. The positional displacement of the internal fiducial markers, which are inserted near the tumor, should be
 87 an appropriate surrogate to evaluate internal deformation. Fewer studies focus on volumetric imaging based on
 88 internal fiducial markers. In the proposed method, linear regression coefficients to derive the three-dimensional
 89 internal deformation from the displacement of the fiducial markers are optimized in each voxel using four-
 90 dimensional computed tomography (4DCT) images as the training dataset. CT image can be generated in real-time
 91 by applying the deformation derived by the displacement of the fiducial markers to the reference CT image. The
 92 purpose of this study was to demonstrate the clinical feasibility of the proposed real-time CT image generation
 93 technique. The accuracy of image synthesis and tumor localization were evaluated using dataset generated by
 94 extended NURBS-Based Cardiac-Torso (XCAT²¹) phantom and clinical 4DCT datasets from six patients.

95 **Materials and Methods**

96 *Algorithm of real-time CT image generation*

97 Figure 1 shows the general scheme of the proposed real-time CT image generation technique, including the training
 98 process and the assumed use in the treatment session. In the training process (before the treatment), a linear
 99 regression coefficients matrix to evaluate the internal deformation from the displacement of the fiducial markers is
 100 generated for each voxel using a 4DCT dataset. The internal deformation \mathbf{Y}_i is derived from the deformable image
 101 registration (DIR) relative to the exhale phase CT, referred to as phase 50 in this study, as the reference volume
 102 \mathbf{V}_{ref} . \mathbf{Y}_i is a matrix with three columns and the number of rows for the matrix equals the number of phases of 4DCT,
 103 including the three-dimensional deformation in all respiratory phases at voxel i . The displacement of the fiducial
 104 markers \mathbf{X} is evaluated for each CT dataset relative to the reference. \mathbf{X} is a matrix whose columns and rows are $3n$
 105 and the number of phases, respectively, including the three-dimensional displacement of the n internal fiducial
 106 markers in all respiratory phases. Assuming a linear relationship between the internal deformation and the
 107 displacement of the fiducial markers, their relationship is described as $\mathbf{Y}_i \cong \mathbf{X}\mathbf{W}_i$, where \mathbf{W}_i is a matrix of linear
 108 regression coefficients whose columns and rows are 3 and $3n$, respectively. The linear relationship in the lung
 109 region, including the tumor, was considered to be reasonable since the fiducial markers were inserted around the
 110 tumor. \mathbf{W}_i can be derived mathematically by regarding \mathbf{X} and \mathbf{Y}_i as the explanatory and response variables,

111 respectively. When there is a high correlation between the explanatory variables, output values may fluctuate
 112 significantly due to multicollinearity in the ordinary multiple linear regression. In this study, partial least squares
 113 regression (PLSR)^{22,23}, which can address multicollinearity in regression analysis, was utilized to derive W_i .



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Fig. 1. General scheme of the proposed real-time CT image generation technique.

118 In the first step of the treatment session, a patient is aligned to the planning position, according to rigid anatomical
 119 landmarks such as the vertebra, by an image guidance technique, such as cone beam CT (CBCT). In this study, we
 120 assumed an orthogonal x-ray imaging system to obtain the three-dimensional position of the internal fiducial
 121 markers. Treatment system with floor/ceiling mounted orthogonal imaging device has been clinically applied in
 122 photon therapy⁶, and gantry mounted type has been realized in proton therapy¹⁰. Please refer the literatures for the
 123 detail of treatment system. Geometrical relationship between the iso-center and the reference position of the fiducial
 124 markers is determined in treatment planning process in advance. The three-dimensional positions of the fiducial
 125 markers are obtained in real-time, using orthogonal x-ray imaging during treatment. The three-dimensional
 126 positional displacement of the n fiducial markers $X(t)$ at time t is obtained as the difference between the current
 127 and reference position of the fiducial markers. The three-dimensional deformation vector $\hat{Y}_i(t)$ at voxel i is derived

128 as $\hat{\mathbf{Y}}_i(t) = \mathbf{X}(t)\mathbf{W}_i$, according to the marker displacement. The volumetric image $\mathbf{V}(t)$ can be synthesized by
129 deforming the reference volume \mathbf{V}_{ref} with the deformation vector field $\hat{\mathbf{Y}}(t)$, which is constructed from all voxels.

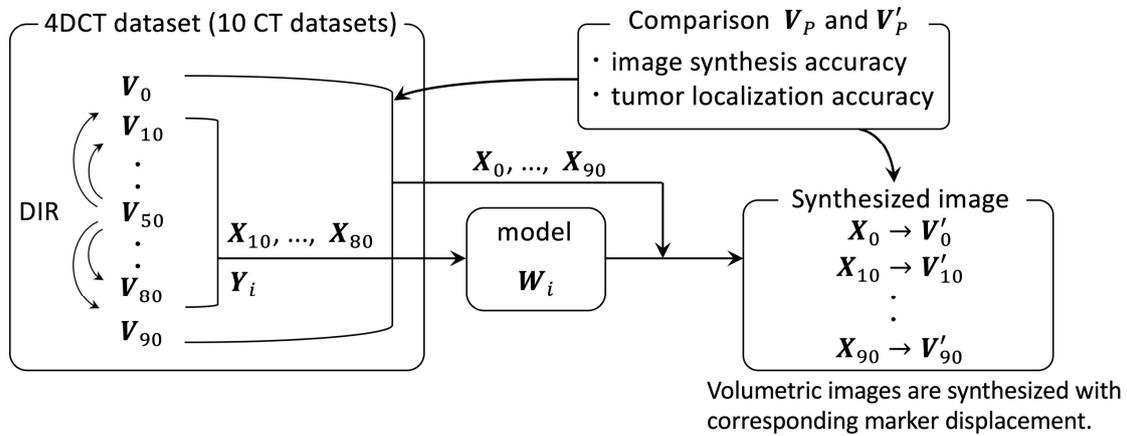
130 *Validation with XCAT phantom*

131 For the evaluations, a ground truth volumetric image was necessary. In this study, the evaluations were conducted
132 by using the dataset of volumetric images generated by extended NURBS-Based Cardiac-Torso (XCAT) phantom²¹
133 as with the previously reported studies^{24,25}. XCAT phantom can generate three-dimensional imaging data according
134 to the arbitrary anterior-posterior (AP) motion data and superior-inferior (SI) motion data for simulating the motion
135 of chest wall and diaphragm, respectively. In this study, in order to mimic actual patient motion, internal marker
136 motion data and external surrogate motion data that were acquired simultaneously by using RTRT system were
137 used. The external surrogate motion and SI direction motion of the fiducial marker were resampled into 5 Hz and
138 were used as input motion data for simulating chest wall and diaphragm motion, respectively, in XCAT phantom.
139 One motion dataset that can simulate the change of motion amplitude between training session and validation
140 session was used. Motion range of the tumor in training session and validation session was about 10 and 15 mm,
141 respectively. Tumor of which diameter was 20 mm and one fiducial marker were inserted in right lower lung region.
142 Image synthesis accuracy and tumor localization accuracy were evaluated as described later.

143 *Validation with patient 4DCT dataset*

144 The evaluations were also conducted with the patient 4DCT dataset. The schematic representation of the testing
145 protocol with patient 4DCT dataset for this study is shown in Figure 2. The 4DCT dataset included ten respiratory
146 phases. Eight CT images from phase 10 to phase 80 were used to obtain the matrix of linear regression coefficients
147 in the training process. The CT images were generated for each respiratory phase using the corresponding marker
148 displacement that was derived from 4DCT dataset and compared with the original CT datasets. The two excluded
149 inhale CT datasets, defining phases 0 and 90, corresponded to validation of the datasets that had large deformations
150 not included in the training process. Evaluation with the training CT datasets containing phases 10 to 80 was
151 regarded as the validation of model goodness of fit.

152 MATLAB (MathWorks Inc, USA) was used for data processing, including CT image import/export, PLSR, and
 153 DIR.



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Fig. 2 Schematic representation of the testing protocol with patient 4DCT dataset.

158 *Patient data for evaluation*

159 The 4DCT datasets from six patients, who underwent lung RTRT, were used for the analysis. Slice thickness, image
 160 size, and pixel spacing for all CT images were 2.5 mm, 512×512 pixels, and 0.98 mm/pixel, respectively. We set
 161 two criteria for data selection. The first criterion was that fiducial markers, with a three-dimensional motion range
 162 in ten respiratory phases of more than 5 mm, were included. The second criterion was that the difference in motion
 163 range between the ten respiratory phases and the eight respiratory phases without the two inhale phases was more
 164 than 1 mm. The motion ranges for each fiducial marker in the six patients are summarized in Table 1. Three or four
 165 fiducial markers have been included for each patient. Harada et al. reported²⁶ that the mean motion amplitude during
 166 radiation therapy was larger than that evaluated in a 4DCT dataset by about 2 mm. Four 4DCT datasets in this study
 167 had motion differences of more than 2 mm; hence, these evaluations adequately mimicked the clinical situation.

168 **Table 1¹.** Motion range for each internal fiducial marker in six patients.

Motion range in LR/SI/AP/3D [mm]

¹ RML, right middle lobe; LUL, left upper lobe; LLL, left lower lobe; RLL, right lower lobe.

Pat. #	Tumor location	Marker #	from phase 0 to phase 90	from phase 10 to phase 80	difference
#1	RML	1	3.1/7.5/5.7/9.9	2.1/5.5/5.7/8.2	1.0/2.0/0.0/1.7
		2	2.4/10.6/3.3/11.4	2.4/10.6/3.3/11.4	0.0/0.0/0.0/0.0
		3	2.6/9.3/4.6/10.7	1.5/6.5/4.6/ 8.1	1.1/2.8/0.0/2.6
#2	LUL	1	1.8/5.2/2.9/6.2	1.5/4.7/2.7/5.6	0.3/0.5/0.2/0.6
		2	2.6/6.6/3.1/7.7	1.7/5.7/2.5/6.4	0.9/0.9/0.6/1.3
		3	1.2/6.0/3.6/7.1	1.2/4.8/2.9/5.7	0.0/1.2/0.7/1.4
		4	2.4/5.5/5.5/8.1	1.1/5.5/3.8/6.8	1.3/0.0/1.7/1.3
#3	LLL	1	3.2/20.0/6.8/21.4	2.6/10.0/6.8/12.3	0.6/10.1/0.0/9.1
		2	2.2/18.5/4.7/19.3	2.2/10.7/4.7/11.9	0.0/7.8/0.0/7.3
		3	5.0/23.0/9.5/25.4	5.0/12.5/7.0/15.2	0.0/10.5/2.5/10.2
		4	7.2/25.6/5.7/27.2	7.0/22.1/5.1/23.7	0.2/3.5/0.6/3.5
#4	LUL	1	1.6/2.6/2.2/3.7	1.3/2.5/2.0/3.4	0.3/0.1/0.2/0.3
		2	1.5/5.8/2.4/6.4	1.2/4.6/2.4/5.3	0.3/1.2/0.0/1.1
		3	1.9/7.6/2.3/8.2	1.9/6.0/2.3/6.7	0.0/1.6/0.0/1.5
#5	RLL	1	1.8/12.6/3.4/13.1	1.8/7.9/2.0/8.4	0.0/4.7/1.4/4.7
		2	1.8/15.9/2.5/16.2	1.3/11.2/1.4/11.3	0.5/4.7/1.1/4.9
		3	1.4/18.0/3.1/18.4	0.7/12.0/3.0/12.3	0.7/6.0/0.1/6.1
		4	0.5/18.2/1.2/18.2	0.5/17.2/1.2/17.3	0.0/1.0/0.0/0.9
#6	RML	1	4.2/3.2/4.0/6.6	3.6/3.2/3.5/6.0	0.6/0.0/0.5/0.6
		2	1.9/6.0/2.9/6.9	1.9/6.0/2.3/6.7	0.0/0.0/0.6/0.2
		3	2.9/3.2/5.2/6.7	2.2/3.2/2.5/4.6	0.7/0.0/2.7/2.1
		4	4.8/2.5/1.4/5.6	4.2/2.5/1.4/5.0	0.6/0.0/0.0/0.6

169

170 Three-dimensional positions of the fiducial markers in the CT images were determined using projection images
171 created in the anterior-posterior (AP) and left-right (LR) directions. First, maximum intensity projection (MIP)
172 images were created. Then, the images were converted to binary images with a threshold CT value of 2000 HU to
173 segment the marker. The marker position was designated as the center of gravity of the rectangular-shaped region
174 of interest enclosing the segmented fiducial marker.

175 *Accuracy of image synthesis*

176 In this study, RMSE and NRMSE between the ground truth CT images and the synthesized CT images were
 177 evaluated as the quantitative indices of the accuracy for the image synthesis. RMSE and NRMSE were evaluated
 178 as:

$$179 \quad \text{RMSE} = \sqrt{\frac{1}{n} \sum_{i=1}^n (I_i - I_i^*)^2},$$

$$180 \quad \text{NRMSE} = \sqrt{\frac{\sum_{i=1}^n (I_i - I_i^*)^2}{\sum_{i=1}^n I_i^{*2}}},$$

181 where I_i^* is the actual CT value at voxel i in the ground truth image, I_i is the CT value at voxel i in the synthesized
 182 image, and n is voxel number. In addition, structural similarity index measure (SSIM) which is commonly used in
 183 pattern matching as a measure of the similarity of image structures was evaluated as:

$$184 \quad \text{SSIM} = \frac{(2\mu_x\mu_y + C_1)(2\sigma_{xy} + C_2)}{(\mu_x^2 + \mu_y^2 + C_1)(\sigma_x^2 + \sigma_y^2 + C_2)},$$

185 where μ_x and μ_y are the average CT values in ground truth image and synthesized image respectively, σ_x and σ_y
 186 are variance of CT value in ground truth image and synthesized image respectively, and σ_{xy} is a covariance of
 187 ground truth image and synthesized image. C_1 and C_2 were defined as $C_1 = (0.01 \times L)^2$ and $C_2 = (0.03 \times L)^2$,
 188 respectively. L was dynamic range of the image and was defined as the maximum CT value except for the fiducial
 189 marker in the ground truth image. RMSE, NRMSE and SSIM were evaluated in the three-dimensional region
 190 constructed from the body half including the tumor and the slices encompassing tumor motion in the superior-
 191 inferior (SI) direction. The performance of the proposed technique was limited by the accuracy of the DIR. For the
 192 benchmark of the image synthesis accuracy, the reference image, referred to as phase 50 in this study, was deformed
 193 to the ground truth images. RMSE, NRMSE and SSIM were evaluated by comparing the ground truth image and
 194 the registered image. To quantitatively confirm the goodness of fit in the linear regression, percent variance
 195 explained (PVE), which corresponds to the determination coefficient in the linear regression, was examined in the

196 PLSR. In XCAT phantom, NRMSE and SSIM were evaluated since the dataset generated by XCAT phantom
197 consisted of attenuation coefficient, not of CT value.

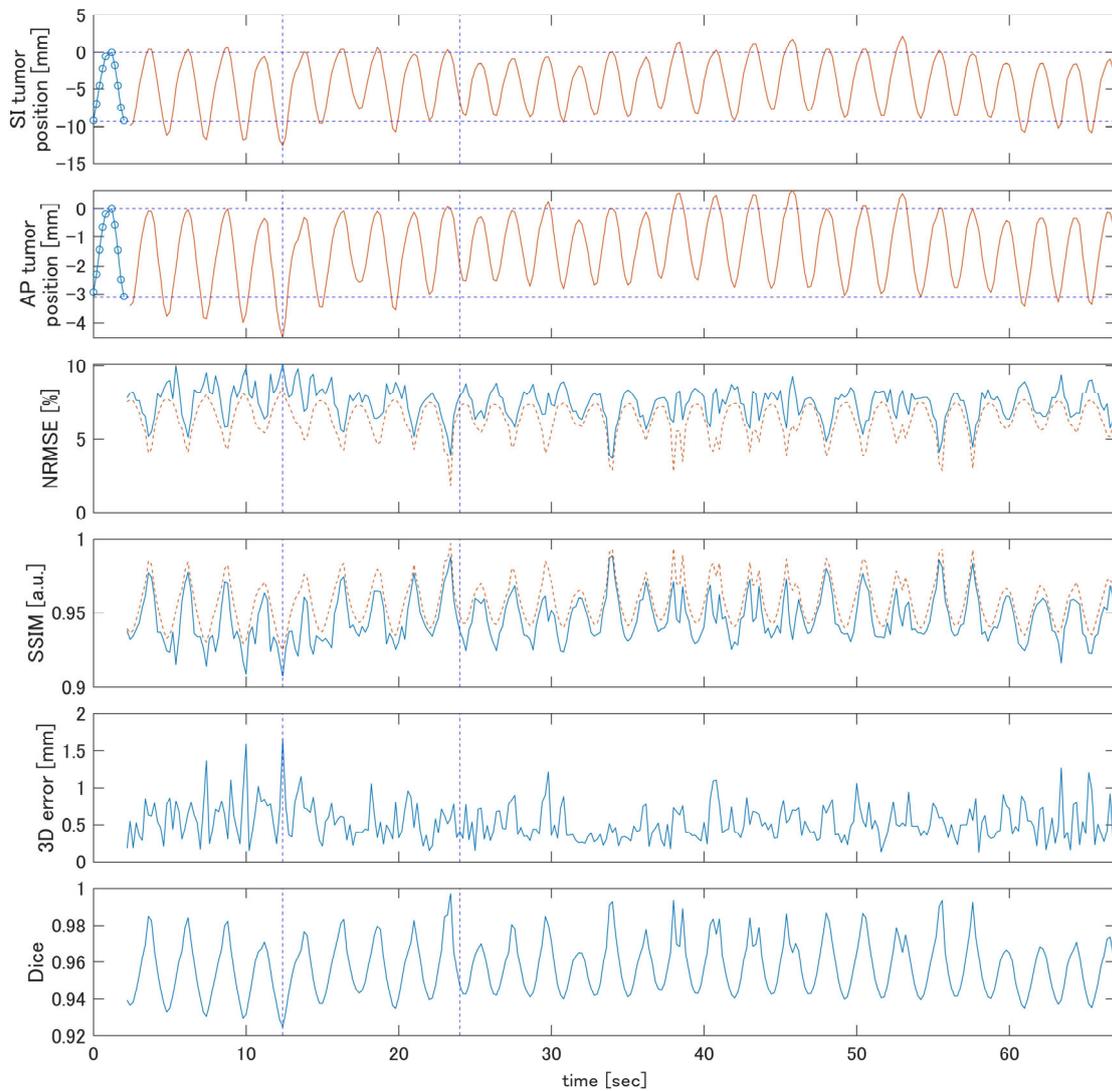
198 *Accuracy of tumor localization*

199 Tumor was manually segmented with referring the images binarized with CT value which was evaluated from the
200 tumor in the ground truth image as threshold. After delineation of the tumor region in both ground truth image and
201 the synthetic image, the distance of center-of-mass between two tumor regions was evaluated as tumor localization
202 accuracy. Dice coefficient between two delineated tumor regions was also evaluated in order to confirm the
203 similarity of the tumor shape. Since the uncertainty of tumor delineation in patients #2 and 4 whose tumor in lung
204 peripheral region was in contact with the pleura would be large, evaluation of tumor localization accuracy was
205 limited to four patients, #1, 3, 5, and 6, whose tumor was located in the central lung region.

206 **Results**

207 *Validation with XCAT phantom*

208 Figure 3 shows NRMSE, SSIM, three-dimensional tumor localization error and Dice coefficient along with the
209 tumor motion. All metrics showed better performance when the displacement of tumor position from the reference
210 position was decreased. Relatively large error was observed when the tumor motion exceeded the range of training
211 data. Average of NRMSE, SSIM and three-dimensional tumor localization error were $7.5\pm 1.1\%$ (ranging from 2.2
212 to 10.1%), 0.95 ± 0.02 (ranging from 0.91 to 0.99) and 0.4 ± 0.3 mm (ranging from 0.0 to 1.3 mm), respectively. Trend
213 of NRMSE and SSIM of the synthesized CT images were similar to those of registered images by DIR which can
214 be regarded as benchmark. Examples of ground truth, synthesized, and absolute difference images in the worst and
215 typical NRMSE are shown in Figure 4. Although the difference around the tumor and diaphragm were large in the
216 worst case compared with the typical cases, internal anatomical structures were reasonably reconstructed with
217 NRMSE less than 10%.

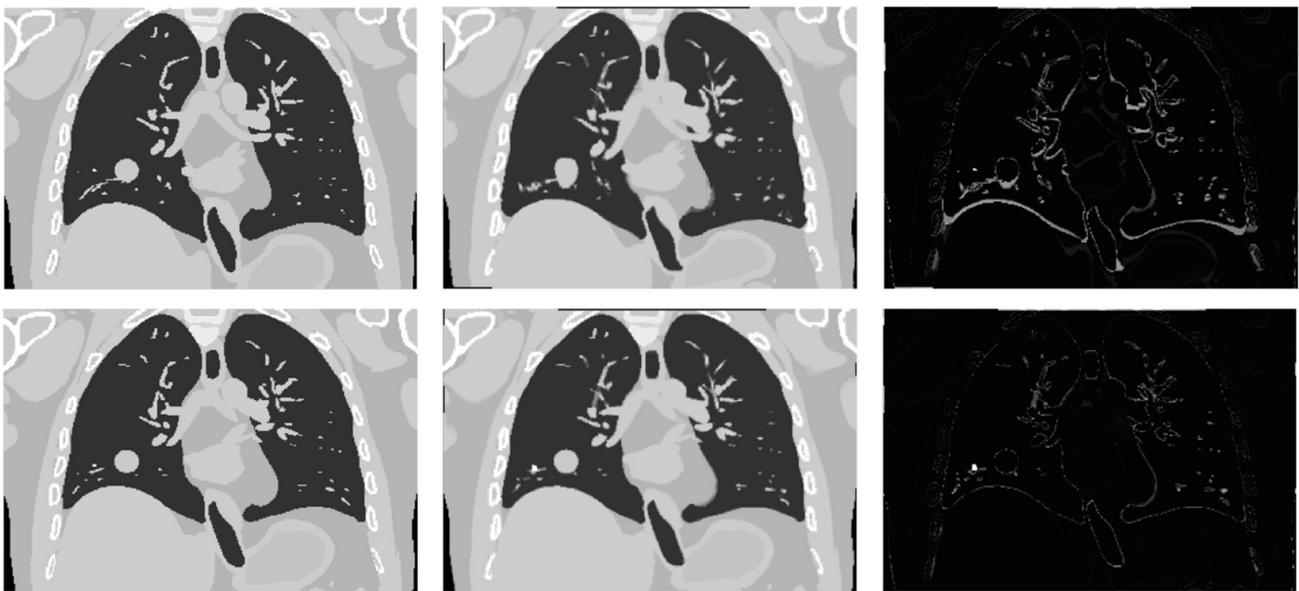


218

219 **Fig. 3** Top two rows: SI and AP motion of the tumor, respectively. Ten circled points in the first part of the trajectory
 220 were corresponding to the training data. The sixth dataset at the exhale phase was used as the reference dataset. The
 221 rest data was used for the validation. Horizontal dashed lines show the motion range of the training data. Third row
 222 from the top: NRMSE in the validation data. Solid line and dashed line represent the NRMSE evaluated with the
 223 synthesized CT images and the registered CT images by DIR, respectively. Fourth row from the top: SSIM in the

224 validation data. Solid line and dashed line represent the SSIM evaluated with the synthesized CT images and the
 225 registered CT images by DIR, respectively. Fifth row from the top: three-dimensional tumor localization error.
 226 Bottom row: Dice coefficient between the tumor regions in synthesized images and ground truth images. Vertical
 227 dashed lines around 12 sec and 24 sec are corresponding to the data positions having worst and typical NRMSE.
 228 The example images in each dataset are shown in Figure 4.

229



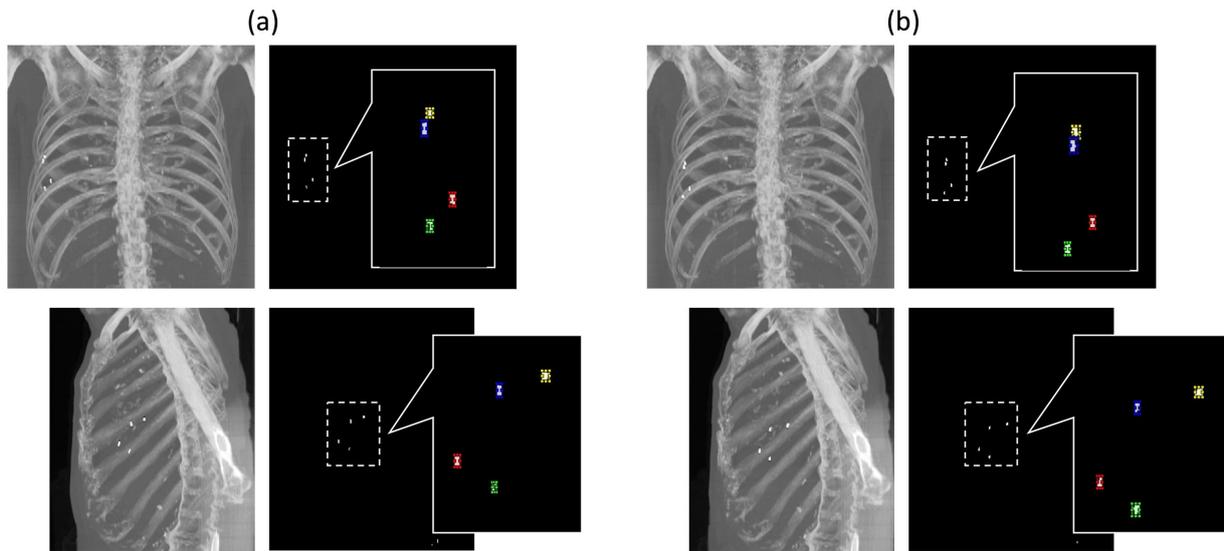
230

231 **Fig. 4** Top row: from left column, sagittal view of the ground truth image, synthesized image and the absolute
 232 difference image between the ground truth image and synthesize image in case of the worst RMSE of 10.1%. Bottom
 233 row: images in case of the RMSE of 7.5%. The tumor is a round object in right lower lung region. Fiducial marker
 234 is shown at the right side of the tumor.

235 *Accuracy of image synthesis in patient dataset*

236 Examples of MIP images and binary images for the marker segmentation of patient #1 are shown in Figure 5. The
 237 rectangle enclosing the segmented marker area was larger than the actual marker size, especially in the SI direction,
 238 due to motion artifact. The center of the rectangle was defined as the marker position. In this study, the mean size

239 of the rectangle in the SI direction was about 5.0 mm. Considering the marker diameter of 1.5 mm, the range of
 240 respiratory motion, and CT image acquisition time of several hundred milliseconds, it was thought that most of the
 241 marker positions were correctly determined.



242

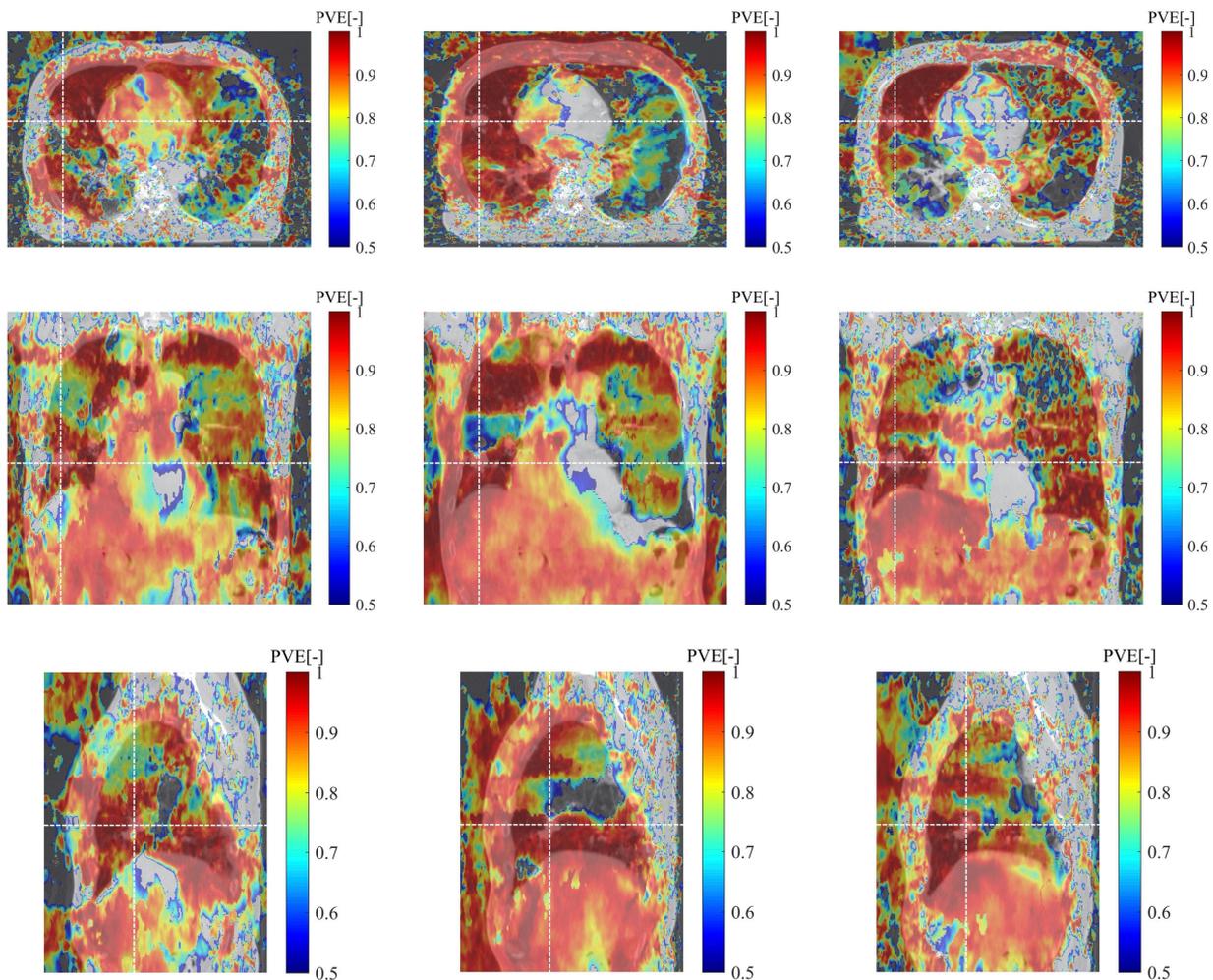
243 **Fig. 5** (a) Maximum intensity projection image and binary image for marker segmentation generated in the anterior-
 244 posterior (top row) and left-right (bottom row) directions in phase 50. The colored rectangles represent the marker
 245 enclosing area and its center was used to define the three-dimensional marker position. Images generated in phase
 246 0 are shown in (b). The motion of the fiducial markers between phase 0 and 50 due to respiration were confirmed.
 247

248 Examples of the PVE distribution for each deformation in the LR, AP, and SI directions evaluated in patient #1 are
 249 shown in Figure 6. Relatively high PVEs were obtained around the tumor and the fiducial markers. The median,
 250 25th, and 75th percentile of PVEs for each patient are summarized in Table 2. Focusing on the median values, high
 251 PVEs of more than 0.8 for deformations in all directions were obtained in most cases. These results suggest that
 252 there was reasonable linearity between the displacement of fiducial markers and the internal deformation in the
 253 evaluation region.

254 **Table 2.** Median (25th - 75th percentile) of percent variance explained (PVE) representing the strength of
 255 relationship between the displacement of the markers and the deformation for each patient.

	#1	#2	#3	#4	#5	#6
LR	0.82 (0.60 – 0.93)	0.88 (0.61 – 0.97)	0.86 (0.66 – 95)	0.78 (0.49 – 0.92)	0.80 (0.58 – 0.90)	0.80 (0.53 – 0.94)
AP	0.88 (0.64 – 0.96)	0.97 (0.86 – 0.99)	0.88 (0.72 – 95)	0.87 (0.63 – 0.94)	0.76 (0.49 – 0.90)	0.91 (0.65 – 0.97)
SI	0.80 (0.58 – 0.92)	0.95 (0.77 – 0.99)	0.84 (0.50 – 0.95)	0.84 (0.53 – 0.94)	0.87 (0.65 – 0.95)	0.78 (0.52 – 0.94)

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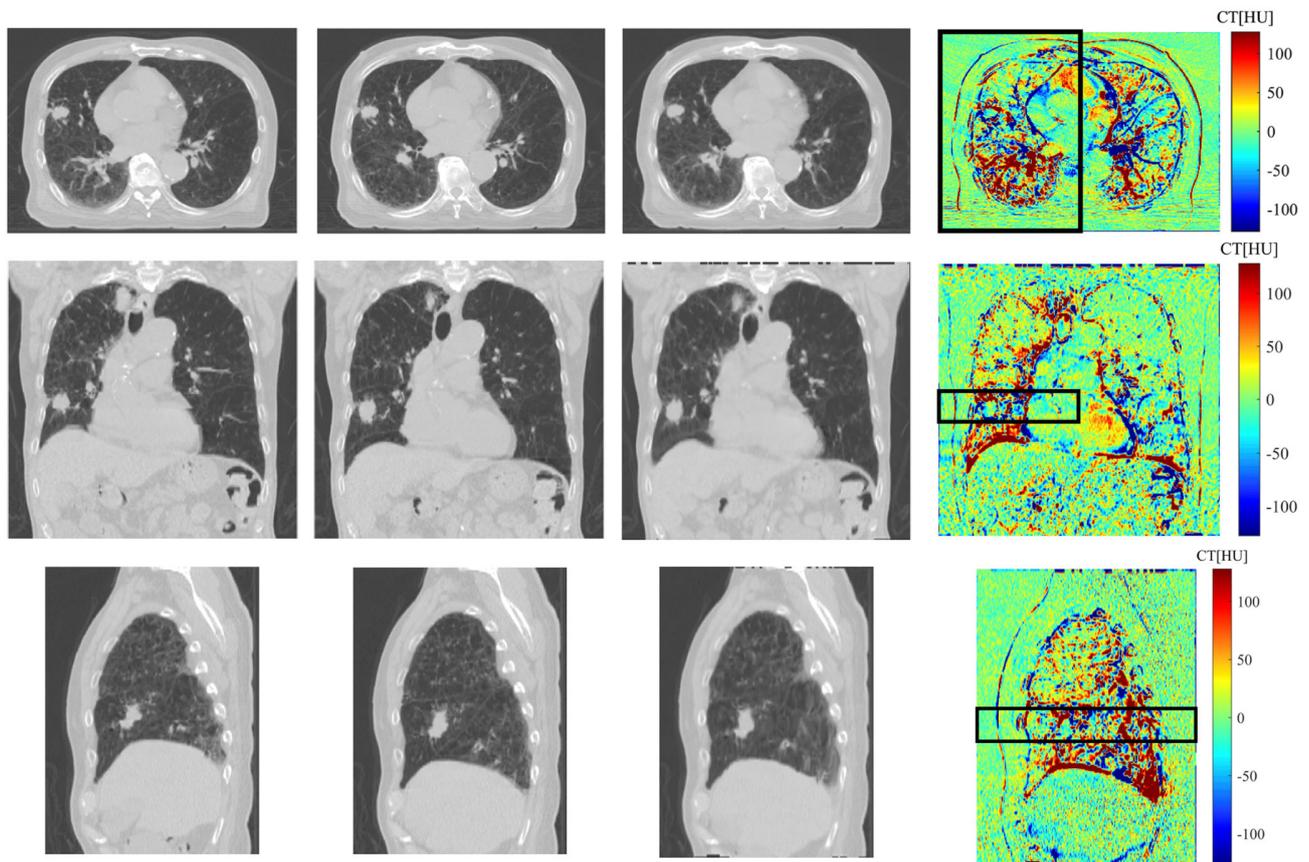
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Fig. 6 The distribution of percent variance explained (PVE) obtained in the partial least squares regression (PLSR) between the displacement of the fiducial markers and the internal deformation in the left-right (left column), anterior-posterior (center column), and superior-inferior (right column) directions. Axial, coronal, and sagittal planes including the target center are shown in the top, middle, and bottom rows.

263 Examples of reference, ground truth, synthesized, and difference images for phase 0 in patient #1 are shown in
 264 Figure 7. The internal anatomical structure, including the tumor shape, was well reconstructed in the synthesized
 265 image. Since cardiac motion cannot be modeled by the proposed method, differences in CT values around the heart
 266 were increased. In this example, differences in CT values in the back side region tended to increase since the
 267 linearity in phase 0 was probably decreased. Note that CT dataset of phase 0 was not included in the training dataset.
 268 To quantify the accuracy of the image synthesis, box and whisker plots of the absolute differences in CT values,
 269 RMSE, NRMSE and SSIM, were evaluated. Differences in CT values may be small in regions where deformations
 270 were relatively small, such as the upper lung region. Hence, to avoid underestimations, evaluation areas were limited
 271 by three-dimensional regions, including the body half represented by a black solid line shown in the difference
 272 images in Figure 7.

273

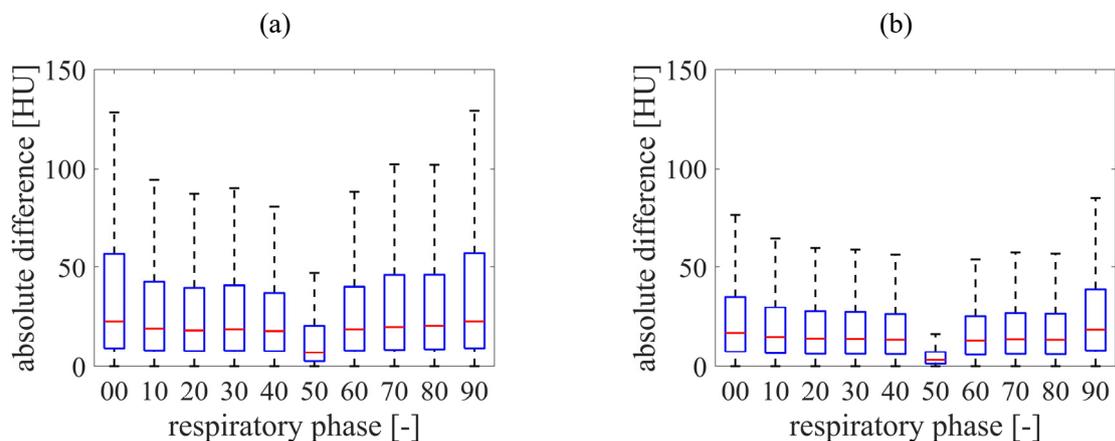


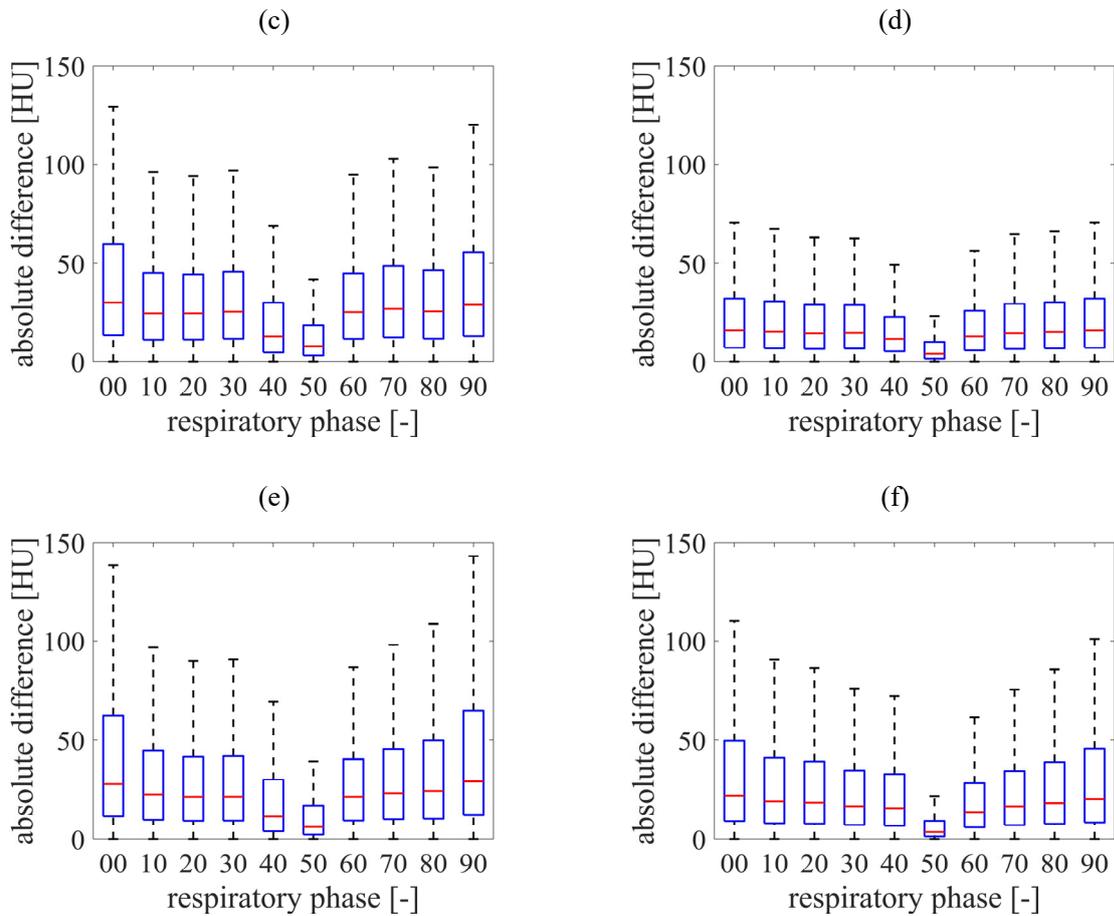
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275 **Fig. 7** From left column, reference image of phase 50, ground truth image of phase 0, synthesized image targeting
 276 phase 0 and difference image between the ground truth image and synthesized image. Axial, coronal and sagittal
 277 plane are shown from top row. Black solid rectangular in the difference image represent the evaluation region of
 278 PVE, RMSE, NRMSE and SSIM.

279

280 To show the distribution of the absolute differences in CT values between the ground truth image and the
 281 synthesized image, box plots for each patient and each respiratory phase are shown in Figure 8. The bottoms and
 282 tops of the boxes represent the 25% point (Q1) and 75% point (Q3) of the absolute difference in the CT values,
 283 respectively. The lower and upper whiskers are $Q1 - 1.5 \times (Q3 - Q1)$ and $Q3 + 1.5 \times (Q3 - Q1)$. Large differences were
 284 observed around the fiducial markers and bones having high CT values. The ratio of voxels with a large discrepancy
 285 (more than upper whisker) to all voxels in the evaluation area was about 10%. The samples having large differences
 286 were regarded as outliers and not shown in the plot. In all patients, absolute differences tended to increase in the
 287 inhale respiratory phase and the distribution expanded gradually. The median was less than 25 HU in most cases.





288 **Fig. 8** Boxplots of the absolute difference of CT value between the ground truth image and the synthesized image
 289 for each patient. Listed from patient #1 to #6 in order from (a) to (f).

290

291 To quantify the accuracy of image synthesis, RMSE and NRMSE are summarized in Tables 3 and 4, respectively.
 292 For comparison to the benchmark, RMSE and NRMSE evaluated with the registered image by DIR are also
 293 summarized. As shown in Table 3, RMSEs from phase 10 to phase 80, corresponding to validation with the training
 294 dataset, were comparable to the benchmark performance in all patients. Since the linear regression by PLSR can
 295 addresses multicollinearity in multiple explanatory variables, the volumetric images synthesized from the input
 296 within the range of the training dataset were expected to be stable. In the evaluation of phases 0 and 90, which had
 297 marker displacement larger than that of the training dataset, the mean RMSEs were about 100 HU. The trends in
 298 NRMSEs were similar to the trends in RMSEs. Hence, there was less dependence of the image synthesis accuracy

299 on the distribution of CT values. SSIM are summarized in Table 5. As with the evaluations of RMSE and NRMSE,
 300 SSIMs evaluated from phase 10 to phase 80 were comparable to the benchmark performance in all patients, although
 301 SSIMs in phase 0 and 90 were decreased a little. These results suggested that the both image intensity and structure
 302 of the synthesized images could be comparative those of DIR, although the accuracy could be reduced when the
 303 marker displacement exceeds the range of training dataset.

304 The processing time to generate the volumetric images using the proposed method was less than 100 ms using a
 305 conventional CPU (Intel(R) i9-9820X 3.30GHz). Thus, the proposed technique can be applied to real-time processes
 306 during treatment.

307 **Table 3.** The root mean squared error (RMSE) [HU] of image synthesis for each patient. For reference, RMSEs of
 308 deformable image registration (DIR) are shown at the bottom of each cell.

phase	#1	#2	#3	#4	#5	#6	Mean \pm SD
0	111.1	85.4	135.7	62.8	110.0	121.5	104.4 \pm 22.2
	80.2	68.8	83.3	57.1	63.5	76.8	71.6 \pm 8.6
10	77.9	70.3	75.4	62.5	63.4	78.3	71.3 \pm 6.0
	75.0	69.1	73.2	59.7	55.8	78.1	68.5 \pm 7.5
20	68.1	68.5	69.9	62.4	58.6	74.3	67.0 \pm 7.5
	60.7	66.1	63.5	65.0	55.3	66.0	62.8 \pm 4.8
30	72.5	64.9	67.1	61.7	63.9	81.0	68.5 \pm 6.0
	57.4	63.1	61.4	61.0	48.0	76.1	61.2 \pm 7.7
40	60.4	63.1	58.6	59.7	49.0	72.3	60.5 \pm 6.4
	49.1	60.8	54.1	53.8	31.8	70.4	53.3 \pm 10.9
50	50.3	57.1	61.4	56.0	41.5	48.9	52.5 \pm 6.0
	-	-	-	-	-	-	-
60	72.9	65.1	65.4	65.5	48.4	69.5	64.5 \pm 7.1
	55.8	63.9	58.7	63.0	40.9	58.2	56.8 \pm 7.1
70	100.5	67.3	71.4	65.1	65.1	74.5	74.0 \pm 11.4
	101.3	64.6	67.0	65.0	54.0	69.9	70.3 \pm 13.6
80	89.4	66.4	72.8	61.4	68.7	87.2	74.3 \pm 9.7
	82.1	65.3	69.4	60.9	64.5	78.1	70.0 \pm 7.1
90	113.2	95.0	116.6	63.2	105.5	106.2	100.0 \pm 6.5
	81.4	67.7	77.9	60.3	66.2	91.6	74.2 \pm 9.8

Mean	81.6 ± 20.2	70.3 ± 10.7	79.4 ± 24.2	62.0 ± 2.6	67.4 ± 21.8	81.4 ± 19.1	73.7 ± 19.6
± SD	64.3 ± 26.2	58.9 ± 19.8	60.8 ± 22.0	54.6 ± 18.5	48.0 ± 19.0	66.5 ± 23.7	58.9 ± 22.6

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Table 4. The normalized root mean squared error (NRMSE) [%] of image synthesis for each patient. For

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reference, NRMSEs of deformable image registration (DIR) are shown at the bottom of each cell.

phase	#1	#2	#3	#4	#5	#6	Mean ± SD
0	15.2	11.3	15.2	7.6	14.6	15.2	13.2 ± 2.7
	11.0	9.1	9.4	6.9	8.4	9.6	9.1 ± 1.1
10	10.6	9.3	8.5	7.5	8.2	9.8	9.0 ± 0.9
	10.2	9.1	8.2	7.2	7.2	9.8	8.6 ± 1.1
20	9.2	9.0	7.8	7.5	7.5	9.2	8.4 ± 0.7
	8.2	8.7	7.1	7.8	7.1	8.2	7.8 ± 0.5
30	9.8	8.5	7.5	7.4	8.1	9.9	8.5 ± 0.9
	7.8	8.3	6.9	7.3	6.1	9.3	7.6 ± 0.9
40	8.2	8.3	6.5	7.1	6.1	8.8	7.5 ± 0.9
	6.6	8.0	6.0	6.4	4.0	8.6	6.6 ± 1.4
50	6.8	7.5	6.8	6.7	5.2	6.0	6.5 ± 0.7
	-	-	-	-	-	-	-
60	9.7	8.5	7.3	7.8	6.0	8.5	8.0 ± 1.1
	7.5	8.3	6.5	7.5	5.1	7.2	7.0 ± 0.9
70	13.4	8.8	7.9	7.8	8.2	9.1	9.2 ± 1.8
	13.5	8.4	7.5	7.8	6.8	8.6	8.8 ± 2.1
80	12.0	8.7	8.1	7.4	9.0	10.7	9.3 ± 1.4
	11.0	8.5	7.7	7.3	8.5	9.5	8.8 ± 1.1
90	15.5	12.6	13.1	7.6	13.9	13.0	12.6 ± 2.2
	11.1	9.0	8.7	7.2	8.7	11.2	9.3 ± 1.3
Mean ±	11.0 ± 2.8	9.2 ± 1.4	8.9 ± 2.7	7.4 ± 0.3	8.7 ± 3.0	10.0 ± 2.4	9.2 ± 2.6
SD	8.7 ± 3.5	7.7 ± 2.6	6.8 ± 2.5	6.5 ± 2.2	6.2 ± 2.5	8.2 ± 2.9	7.4 ± 2.9

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Table 5. The structural similarity index measure (SSIM) for each patient. For reference, SSIMs of deformable image registration (DIR) are shown at the bottom of each cell.

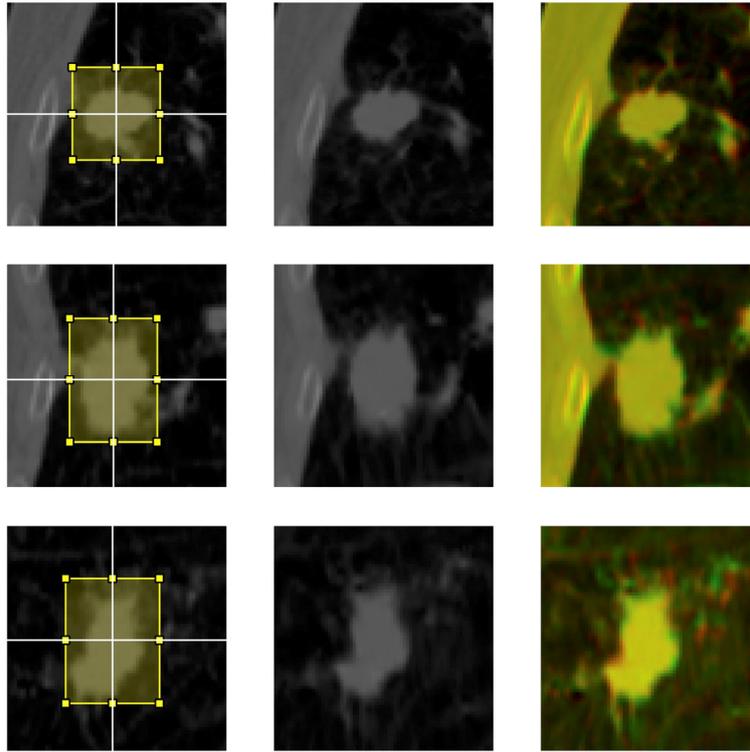
phase	#1	#2	#3	#4	#5	#6	Mean ± SD
0	0.78	0.88	0.70	0.93	0.79	0.81	0.82 ± 0.08
	0.85	0.92	0.82	0.94	0.88	0.88	0.88 ± 0.04
10	0.85	0.92	0.82	0.93	0.88	0.88	0.88 ± 0.04
	0.85	0.92	0.83	0.93	0.90	0.88	0.89 ± 0.04

20	0.87	0.92	0.83	0.93	0.89	0.89	0.89 ± 0.04
	0.88	0.93	0.85	0.94	0.90	0.90	0.90 ± 0.03
30	0.86	0.93	0.82	0.93	0.89	0.89	0.89 ± 0.04
	0.89	0.93	0.84	0.94	0.91	0.91	0.90 ± 0.03
40	0.86	0.93	0.82	0.93	0.89	0.89	0.89 ± 0.04
	0.89	0.93	0.84	0.94	0.91	0.91	0.90 ± 0.03
50	0.93	0.98	0.93	0.97	0.96	0.98	0.96 ± 0.02
	-	-	-	-	-	-	-
60	0.86	0.93	0.83	0.94	0.90	0.93	0.90 ± 0.05
	0.89	0.94	0.84	0.95	0.92	0.95	0.91 ± 0.04
70	0.81	0.93	0.81	0.93	0.88	0.90	0.88 ± 0.05
	0.83	0.93	0.83	0.93	0.90	0.91	0.89 ± 0.05
80	0.82	0.93	0.82	0.93	0.87	0.87	0.87 ± 0.05
	0.83	0.93	0.83	0.93	0.88	0.90	0.88 ± 0.05
90	0.78	0.86	0.73	0.92	0.80	0.82	0.82 ± 0.07
	0.85	0.92	0.82	0.93	0.88	0.86	0.88 ± 0.04
Mean	0.84 ± 0.05	0.92 ± 0.03	0.82 ± 0.07	0.94 ± 0.01	0.88 ± 0.05	0.89 ± 0.05	0.88 ± 0.04
± SD	0.87 ± 0.03	0.93 ± 0.01	0.84 ± 0.04	0.94 ± 0.01	0.90 ± 0.02	0.90 ± 0.02	0.90 ± 0.02

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319 *Accuracy of tumor localization in patient dataset*

320 An example of tumor image in phase 0 of patient #1 is shown in Figure 9. The tumor shape was well reconstructed
321 in the synthesized image. The accuracy of tumor localization and Dice coefficient for each patient are summarized
322 in Table 6. In all evaluated patients, three-dimensional tumor localization error was less than 1.0 mm from phase
323 10 to 80 with Dice coefficient more than about 0.9, which suggested that the tumor shape was reasonably reproduced.
324 Although patient #3 exhibited a large tumor localization error, the three-dimensional tumor localization error in
325 phase 0 and 90 of the other patients was less than 2.0 mm.



326
 327 **Fig. 9** Left column: The ground truth images around the tumor in phase 0 of patient #1. The yellow rectangular
 328 region represents the tumor region. Center column: The synthesized image cropped at the same position as the
 329 ground truth image. Right column: The overlay of the ground truth and synthesized images. The green and red
 330 regions indicate the mismatch of image intensity. The axial, coronal, and sagittal planes are shown from top to
 331 bottom.

332

333 **Table 6.** Tumor localization error [mm] in the left-right (LR), superior-inferior (SI) and anterior-posterior (AP)
 334 direction. Dice coefficient was also summarized as a metric of similarity of tumor shape.

Phase		#1	#3	#5	#6	Mean \pm SD
0	LR	0.5	-0.3	0.8	-1.2	0.0 ± 0.9
	SI	1.5	2.4	0.8	-0.9	1.0 ± 1.4
	AP	-0.5	-2.7	0.5	1.1	-0.4 ± 1.7
	3D	1.7	3.6	1.2	1.9	2.1 ± 1.1
	Dice	0.85	0.73	0.78	0.88	0.81 ± 0.07
10	LR	0.0	-0.5	0.1	-0.7	-0.3 ± 0.4
	SI	-0.6	-0.5	0.6	-0.7	-0.3 ± 0.6
	AP	-0.2	0.3	0.2	0.6	0.2 ± 0.3
	3D	0.6	0.7	0.6	1.2	0.8 ± 0.3
	Dice	0.91	0.91	0.83	0.91	0.89 ± 0.04
20	LR	-0.1	0.0	-0.1	-0.5	-0.2 ± 0.2
	SI	0.0	0.7	0.5	-1.2	0.0 ± 0.8
	AP	0.5	0.2	0.0	0.4	0.3 ± 0.2

	3D	0.5	0.7	0.5	1.4	0.8 ± 0.4
	Dice	0.91	0.93	0.86	0.88	0.90 ± 0.03
30	LR	0.0	-1.0	-0.1	-0.2	-0.3 ± 0.5
	SI	0.0	0.5	-0.1	-0.8	-0.1 ± 0.5
	AP	0.3	0.0	0.6	0.1	0.2 ± 0.3
	3D	0.3	1.1	0.6	0.8	0.7 ± 0.3
	Dice	0.90	0.91	0.86	0.91	0.89 ± 0.02
40	LR	-0.1	-0.1	-0.1	-0.1	-0.1 ± 0.0
	SI	-0.1	-0.4	0.0	-0.6	-0.3 ± 0.3
	AP	0.1	-0.2	0.5	0.2	0.1 ± 0.3
	3D	0.2	0.4	0.5	0.6	0.4 ± 0.2
	Dice	0.90	0.95	0.92	0.89	0.92 ± 0.03
50	LR	0.1	-0.1	0.0	0.0	0.0 ± 0.1
	SI	0.1	0.5	0.2	-0.2	0.1 ± 0.3
	AP	0.0	0.4	0.5	0.1	0.3 ± 0.2
	3D	0.1	0.6	0.6	0.2	0.4 ± 0.3
	Dice	0.94	0.95	0.93	0.97	0.95 ± 0.02
60	LR	-0.2	-0.2	0.1	0.2	0.0 ± 0.2
	SI	0.0	-0.1	0.0	-0.2	-0.1 ± 0.1
	AP	0.0	-0.1	0.3	0.0	0.1 ± 0.2
	3D	0.2	0.2	0.4	0.3	0.3 ± 0.1
	Dice	0.90	0.95	0.91	0.94	0.93 ± 0.03
70	LR	0.2	0.2	0.3	-0.2	0.1 ± 0.2
	SI	-0.5	-0.1	0.8	0.0	0.1 ± 0.5
	AP	0.3	0.0	-0.1	0.4	0.1 ± 0.3
	3D	0.6	0.2	0.9	0.5	0.5 ± 0.3
	Dice	0.86	0.93	0.85	0.93	0.89 ± 0.04
80	LR	0.0	0.0	0.1	0.5	0.2 ± 0.2
	SI	0.3	0.1	0.9	0.0	0.3 ± 0.4
	AP	0.1	-0.2	-0.2	-0.3	-0.2 ± 0.2
	3D	0.3	0.3	1.0	0.6	0.5 ± 0.3
	Dice	0.89	0.94	0.83	0.92	0.90 ± 0.05
90	LR	0.6	0.3	0.5	0.3	0.4 ± 0.1
	SI	1.7	3.8	0.5	-0.6	1.4 ± 1.9
	AP	-0.6	-1.8	0.0	-0.3	-0.7 ± 0.8
	3D	1.9	4.2	0.7	0.7	1.9 ± 1.7
	Dice	0.84	0.76	0.81	0.89	0.83 ± 0.06
Mean ± SD	LR	0.1 ± 0.3	-0.1 ± 0.4	0.2 ± 0.3	-0.2 ± 0.5	0.0 ± 0.2
	SI	0.2 ± 0.8	0.7 ± 1.4	0.4 ± 0.4	-0.5 ± 0.4	0.2 ± 0.5
	AP	0.0 ± 0.3	-0.4 ± 1.0	0.2 ± 0.4	0.2 ± 0.4	0.0 ± 0.3
	3D	0.6 ± 0.6	1.2 ± 1.5	0.9 ± 0.4	0.8 ± 0.5	0.8 ± 0.6
	Dice	0.89 ± 0.03	0.90 ± 0.08	0.86 ± 0.05	0.91 ± 0.03	0.89 ± 0.04

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337 **Discussion**

338 We presented a detailed evaluation of image synthesis accuracy and tumor localization accuracy for the proposed
339 real-time CT image generation technique. The accuracies are similar to previous studies. Mishra et al. proposed
340 volumetric imaging based on principal component analysis (PCA).²⁵ They evaluated NRMSE, assuming the
341 variation of motion magnitude of the chest wall and diaphragm, using XCAT phantom. In case of large variations
342 of motion amplitude, NRMSE increased up to about 15%. Li et al. also demonstrated real-time volumetric imaging
343 based on PCA²⁴ and the mean NRMSE was $6.9 \pm 2.4\%$ when the respiratory motion amplitude changed between
344 the 4DCT acquisition and the treatment session. NRMSEs evaluated in this study were $9.2 \pm 2.6\%$ and $7.5 \pm 1.1\%$ in
345 patient 4DCT dataset and XCAT phantom, respectively. Although the testing protocol, data for validation, and
346 algorithm of DIR were different, the imaging accuracy of the proposed technique is expected to be similar to those
347 of previous research.

348 The three-dimensional tumor localization accuracy in ten respiratory phases for the patient 4DCT dataset was 0.8
349 ± 0.6 mm and 0.4 ± 0.3 mm for the XCAT phantom in this study. These were equivalent to those of previous studies,
350 including the study of Li et al.¹⁹, which demonstrated a three-dimensional accuracy of 0.8 ± 0.5 mm, and the study
351 of Mishra et al.²⁰, which demonstrated an accuracy of 1.0 ± 0.9 mm. The accuracy in phases 0 and 90 of the patient
352 4DCT dataset was less than 2 mm, except for patient #3, and is considered clinically acceptable. Three-dimensional
353 tumor localization accuracy in phases 0 and 90 of patient #3 was about 4 mm. In patient #3, variations in the
354 difference of motion range were relatively large compared with the other cases. Linearity around the tumor may
355 have been decreased in phases 0 and 90 due to the large motion variation. A more robust technique like deep learning,
356 which can address nonlinearity, may be applicable.

357 Motion artifact in 4DCT is a major source of error since the proposed volumetric imaging utilizes the CT datasets
358 to derive marker displacement and internal deformation. CT reconstruction algorithms, which can reduce the
359 artifacts, will be more effective^{27,28} in obtaining accurate deformations. Accuracy of image synthesis may be
360 decreased in regions far from the fiducial markers since linearity between marker displacement and the internal
361 deformation may decrease. Inferring models, such as deep learning, can be combined with the proposed model to
362 address nonlinearity. In this study, patient setup error or inter-fractional anatomical variations, such as tumor

363 shrinkage or weight change, were not simulated in the evaluation, since a series of 4DCT dataset was used for
364 testing. As for initial patient setup, the current image guidance techniques, such as CBCT, can correct patient
365 position with an accuracy of less than 1 mm.^{29,30} However, the current volumetric imaging techniques, including
366 the previous studies, could not handle large internal anatomical variations between 4DCT acquisition and treatment
367 sessions. These variations may cause large errors in image synthesis and tumor localization. To adapt to these
368 variations, 4D-CBCT, acquired just before the treatment, could be utilized for updating the evaluation model.

369 In this study, we assumed an orthogonal x-ray imaging system to obtain the three-dimensional position of the
370 internal fiducial markers. The fiducial markers such as gold sphere of which diameter is 1.5 or 2.0 mm can be
371 recognized in x-ray images by means of pattern matching technique in most cases in lung or liver. Note that there
372 could be a risk of miss detection of the markers in one or both x-ray images if the fiducial markers such as thin
373 coiled shape are used³¹. In the current RTRT, the fiducial markers should be recognized in both x-ray images for
374 three-dimensional calculation. On the other hand, the treatment systems that have dual imaging functions during
375 treatment are not widely used yet. Li et al. proposed deriving the three-dimensional position from a single planar x-
376 ray image.³² A study on volumetric imaging utilizing the fiducial markers with a single x-ray imaging device, which
377 can be widely applied in most clinical systems, is the next step of investigation for the proposed technique.

378 Regarding clinical significance, the proposed CT image generation technique can be applied to real-time beam
379 gating/tracking as a motion management technique based on tumor position and anatomical structure of the
380 surrounding organs, especially for lungs in SBRT and particle therapy. For instance, since the volumetric images
381 have CT values similar to treatment planning CTs, treatment beam gating based on the evaluation of water
382 equivalent thickness for particle therapy³³ could be realized. Synthesized volumetric images could be applied for
383 both real-time dose calculations during treatment³⁴ and retrospective dose accumulations. In addition, imaging doses
384 due to continuous x-ray imaging can be minimized by collimating the imaging region using only the marker area.
385 In all volumetric imaging techniques, the imaging accuracy will be decreased when actual motion is different from
386 the training dataset. Displacement of the fiducial markers could also be used for interlock-function. For instance,
387 when the marker displacement exceeds the threshold value, the treatment beam can be forced off. The volumetric

388 imaging technique using the fiducial markers can be one of the options due to its effectiveness, although the
389 technique requires marker insertion and an additional imaging dose during treatment.

390 **Conclusion**

391 In this study, we proposed a real-time CT image generation technique utilizing the positional displacement of
392 internal fiducial markers evaluated in real-time during the treatment. The XCAT phantom dataset and patient 4DCT
393 dataset, including ten CT datasets for each respiratory phase, were used for the evaluation of image synthesis
394 accuracy and tumor localization accuracy. In the validation with XCAT phantom, the mean NRMSE, SSIM and
395 three-dimensional tumor localization error were $7.5\pm 1.1\%$, 0.95 ± 0.02 and 0.4 ± 0.3 mm, respectively. In the
396 evaluation with patient 4DCT dataset, eight CT datasets, excluding two CT datasets of the inhale respiratory phase,
397 were used in the training process. CT images were generated for each of the ten respiratory phases using the
398 corresponding marker displacements and compared with the original CT dataset. RMSEs, NRMSEs and SSIMS
399 from phase 10 to 80, corresponding to the training dataset, were comparable to those of DIR. The three-dimensional
400 tumor localization error was less than 1 mm. In the evaluation of phases 0 and 90, which had marker displacement
401 larger than that of the training dataset, the mean RMSEs and NRMSEs were about 100 HU and 13%, respectively,
402 and were comparable to previous studies. Three-dimensional tumor localization errors in phases 0 and 90 were less
403 than 2.0 mm, except for one case having large motion variation. These results demonstrate the feasibility of the
404 proposed real-time CT image generation technique.

405 **References**

406

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514 Conflict of Interest Statement

515 The authors have no relevant conflicts of interest to disclose.