

[A hybrid reconstruction algorithm for fast and accurate 4D cone-beam](http://dx.doi.org/10.1118/1.4881326) [CT imaging](http://dx.doi.org/10.1118/1.4881326)[a\)](#page-10-0)

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(Received 21 January 2014; revised 18 April 2014; accepted for publication 15 May 2014; published 9 June 2014)

Purpose: 4D cone beam CT (4D-CBCT) has been utilized in radiation therapy to provide 4D image guidance in lung and upper abdomen area. However, clinical application of 4D-CBCT is currently limited due to the long scan time and low image quality. The purpose of this paper is to develop a new 4D-CBCT reconstruction method that restores volumetric images based on the 1-min scan data acquired with a standard 3D-CBCT protocol.

Methods: The model optimizes a deformation vector field that deforms a patient-specific planning CT (p-CT), so that the calculated 4D-CBCT projections match measurements. A forward-backward splitting (FBS) method is invented to solve the optimization problem. It splits the original problem into two well-studied subproblems, i.e., image reconstruction and deformable image registration. By iteratively solving the two subproblems, FBS gradually yields correct deformation information, while maintaining high image quality. The whole workflow is implemented on a graphic-processing-unit to improve efficiency. Comprehensive evaluations have been conducted on a moving phantom and three real patient cases regarding the accuracy and quality of the reconstructed images, as well as the algorithm robustness and efficiency.

Results: The proposed algorithm reconstructs 4D-CBCT images from highly under-sampled projection data acquired with 1-min scans. Regarding the anatomical structure location accuracy, 0.204 mm average differences and 0.484 mm maximum difference are found for the phantom case, and the maximum differences of 0.3–0.5 mm for patients 1–3 are observed. As for the image quality, intensity errors below 5 and 20 HU compared to the planning CT are achieved for the phantom and the patient cases, respectively. Signal-noise-ratio values are improved by 12.74 and 5.12 times compared to results from FDK algorithm using the 1-min data and 4-min data, respectively. The computation time of the algorithm on a NVIDIA GTX590 card is 1–1.5 min per phase.

Conclusions: High-quality 4D-CBCT imaging based on the clinically standard 1-min 3D CBCT scanning protocol is feasible via the proposed hybrid reconstruction algorithm. *© 2014 American Association of Physicists in Medicine*. [\[http://dx.doi.org/10.1118/1.4881326\]](http://dx.doi.org/10.1118/1.4881326)

Key words: 4D cone beam CT, reconstruction, GPU

1. INTRODUCTION

Respiratory motion is a major problem in lung and abdomen radiotherapy. While advanced 4D treatment planning and delivery techniques have been developed to deliver a highly conformal dose to the moving target, their efficacies are compromised by static in-room imaging guidance, e.g., 3D conebeam CT (CBCT), 1,2 1,2 1,2 1,2 which is registered to the 4D-CT image for patient setup. 4D image guidance is therefore needed for precise tumor targeting, 3 particularly in the context of the increasingly used stereotactic body radio-surgery, $2,4$ $2,4$ where the high fractional dose makes it less forgiving to targeting error. Moreover, adaptive radiation therapy (ART) (Ref. [5\)](#page-10-6) is currently an active research topic, which addresses interfractional variations of patients' respiratory motion pattern $6-8$ $6-8$ and tumor shrinkage $9,10$ $9,10$ by adaptively developing new treatment plans based on the up-to-date anatomy. It calls for highquality in-room 4D images with accurate Hounsfield Units (HU) to facilitate accurate deformable registration and dose calculation for plan adaptations.

In recent years, respiratory-phase resolved CBCT, or 4D-CBCT technique has been developed. $4,11-16$ $4,11-16$ However, current 4D-CBCT technique cannot fully satisfy the aforementioned clinical needs. Specifically, (1) Current 4D-CBCT needs a long scan protocol (typically 4–6 min, referred as 4-min scan hereafter) to provide enough number of projections per breathing phase for reconstruction. This inherently impedes the clinical workflow and elevates the imaging dose, greatly limiting its clinical use; (2) the overall image quality (e.g., signal-to-noise ratio, SNR) of 4D-CBCT is still inferior to that of a standard 3D-CBCT (referred as 1-min scan hereafter), let alone the planning CT; (3) 4D-CBCT usually adopts a full-fan mode to maximize the angular sampling. Nonetheless, the associated small field-of-view (FOV) (e.g., 25 cm in diameter) usually cannot cover the whole thorax/abdomen, leading to truncation problem, $17,18$ $17,18$ and the truncated 4D-CBCT images cannot serve the purpose of dose calculations due to missing anatomies outside the FOV; and (4) the HU of 4D-CBCT images is inaccurate due to scatter contaminations. The factors mentioned here are usually coupled together. Improving one factor may inevitably worsen another. For example, any attempt to reduce scanning time could speed up the workflow and save imaging dose, but would further introduce viewaliasing/streaking artifacts[.19](#page-10-15) Another example is adopting a half-fan mode to increase the FOV. Yet, because only half of the patient is covered at each view angle, the angular sampling rate is effectively reduced. In light of all of these facts, a fast, high-quality, low dose 4D-CBCT imaging technique is in great demand. $20-42$ $20-42$

One unique feature of radiotherapy is that a patients' own planning CT (p-CT) is always available from the treatment simulation stage. In light of the availability of this *prior* information, 2D-3D matching-based reconstruction models have been proposed in a wide spectrum of imaging tasks, such as tomosynthesis, real-time volumetric imaging, and 3D/4D CBCT. $27, 36, 37, 39, 43-45$ $27, 36, 37, 39, 43-45$ $27, 36, 37, 39, 43-45$ $27, 36, 37, 39, 43-45$ $27, 36, 37, 39, 43-45$ $27, 36, 37, 39, 43-45$ $27, 36, 37, 39, 43-45$ $27, 36, 37, 39, 43-45$ $27, 36, 37, 39, 43-45$ These methods optimize a motion vector field (MVF) to deform the p-CT, such that its forward projections match the measurements. Brock *et al.* proposed to match the forward projections by adjusting the MVF control points, so that the forward projections have minimal mismatch with the measurements. They evaluated the algorithm via simulation and reported that it is sensitive to contrast mismatch errors between the measurements and the calcu-lated projections.^{[27](#page-10-17)} Ren *et al.* proposed to solve a MVF energy function using a nonlinear conjugate gradient (NCG) method. They investigated various scan protocols, suggesting that ∼57 projections over 360◦ are required to achieve acceptable accuracy. 36 To mitigate the local minima problem associated with the NCG framework, Wang and Gu introduced a better initial image using a compressed sensing (CS) based iterative reconstruction algorithm. They tested it with one simulation case and one patient case of a 2-min scan.^{[37](#page-11-2)} Recently, a multiple-step algorithm combing PCA modeling, optical flow based registration and reconstruction was proposed. 39 It was validated with simulation studies and one patient data study and was compared with the state-of-the-art CS-based methods, such as TV (Ref. [46\)](#page-11-6) and PICCS.^{[21,](#page-10-18) [22](#page-10-19)} The results show that it offers better overall compromise in the depiction of moving and stationary anatomical structures.

In spite of the success, there is still room for improvement. It is desirable to reconstruct 4D-CBCT based on projections acquired in a short scan time, e.g., 1-min as in the current clinical protocol for 3D-CBCT. In addition to the limited number of projections, practical issues such as intensity difference due to scatter, truncation, and noise also make this a challenging problem. The local-minima problem in 2D-3D registration is another issue. While standard gradient-based algorithm can be employed to solve the optimization problem, more comprehensive evaluations of reconstruction results in patient cases are needed to demonstrate the clinical utility of algorithms of this kind, particularly with respect to both the intensity accuracy and geometry accuracy. Finally, computation efficiency should be improved to render the algorithm in a clinically applicable form.

In this paper, we will present an alternative approach to the existing algorithms, $36, 37, 39$ $36, 37, 39$ $36, 37, 39$ $36, 37, 39$ $36, 37, 39$ called forward-backward splitting (FBS) algorithm, to solve the 2D-3D registration-based 4D-CBCT reconstruction problem. It splits the model into two well-studied subproblems, i.e., image reconstruction and deformable image registration (DIR). By alternatively solving these two problems in an iterative manner, FBS gradually yield 4D-CBCT images with satisfactory even with very sparsely acquired projections from a 1-min CBCT scan (e.g., ∼10–20 projections over a 200◦ arc for each phase). To achieve a clinically acceptable efficiency, all components along the whole reconstruction workflow have been implemented on graphic processing unit (GPU). The GPU implementation has substantially reduced the reconstruction time to 1–1.5 min per phase. As a comparison, hours of computation time are usually observed for those CPU based 2D-3D matching algorithms.³⁹ Both phantom and multiple real patient data sets are used to test the efficacy of the proposed algorithm. For each case, comprehensive evaluations are conducted regarding the image quality, algorithm robustness, and computational efficiency.

2. METHODS AND MATERIALS

2.A. The hybrid 4D-CBCT reconstruction algorithm

2.A.1. Reconstruction model

The p-CT $f_p(x)$, $x = (x, y, z) \in \mathbb{R}^3$ in this study is chosen to be one phase of the 4D-CT image or a breath-hold CT image, in which there is minimal motion artifacts. We first reconstruct an average 3D-CBCT image using all projections in a 4D-CBCT scan via the conventional Feldkamp-Davis-Kress (FDK) (Ref. [47\)](#page-11-7) algorithm. The p-CT is then rigidly aligned with it. Let us denote 4D-CBCT image at phase *i* as $f_i(x)$, which can be obtained by deforming $f_p(x)$ with an MVF $v_i(\mathbf{x})$, i.e., $f_i(\mathbf{x}) = f_p(\mathbf{x} + v_i(\mathbf{x}))$. In our study, it is our objective to restore $v_i(x)$ by solving the following optimization problem:

$$
v_i = \arg \min_{v_i} E[v_i]
$$

=
$$
\arg \min_{v_i} \frac{1}{2} ||P_i f_p(\mathbf{x} + v_i(\mathbf{x})) - c(g_i)||_2^2 + \frac{\lambda}{2} ||\nabla v_i||_2^2,
$$

(1)

where $\|\cdot\|_2$ represents l_2 norm and ∇ is the gradient operator. P_i is the x-ray projection matrix for the phase *i*, and g_i is the corresponding measured projections. The first term in the objective function is to ensure the fidelity between the reconstructed results and the projection measurements. *c* represents a correction step to address the intensity inconsistency between the measured x-ray projections *gi* and the calculated forward projections *Pifp* due to, e.g., x-ray scatter contaminations in the projection measurements. The second term is to enforce the MVF smoothness. λ is a penalty weight. Once Eq. [\(1\)](#page-2-0) is solved, $f_i(x)$ can be obtained by deforming $f_p(x)$ via $v_i(x)$.

2.A.2. Forward-backward splitting algorithm

Let us first consider the optimality condition of the problem. By taking the derivative of Eq. [\(1\),](#page-2-0) we arrive at

$$
\nabla f_p P_i^T [P_i f_p - c(g_i)] - \lambda \Delta v_i = 0, \qquad (2)
$$

where f_p is the simplified representation of $f_p(\mathbf{x} + v_i(\mathbf{x}))$, P_i^T denotes the matrix transpose of P_i , and Δ is the Laplacian operator. By introducing an auxiliary function $s_i(\mathbf{x})$ and a term $\nabla f_p[f_p - s_i]$, we can get

$$
\nabla f_p P_i^T [P_i f_p - c(g_i)] - \lambda \Delta v_i - \nabla f_p [f_p - s_i]
$$

+
$$
\nabla f_p [f_p - s_i] = 0.
$$
 (3)

Since we have the freedom to choose the function $s_i(\mathbf{x})$, we would like to combine the first and the third term, and the second and the fourth term together, respectively, and split Eq. (3) into

$$
\nabla f_p P_i^T [P_i f_p - c(g_i)] - \nabla f_p [f_p - s_i] = 0,
$$
\n(4)

$$
\nabla f_p[f_p - s_i] - \lambda \Delta v_i = 0. \tag{5}
$$

For Eq. [\(4\),](#page-2-2) assuming $\nabla f_p \neq 0$, we can get

$$
s_i = f_p - P_i^T [P_i f_p - c(g_i)].
$$
\n(6)

Interestingly, Eq. [\(6\)](#page-2-3) is a typical gradient descent update step for a CBCT reconstruction problem, and meanwhile, Eq. [\(5\)](#page-2-4) is essentially the Euler-Lagrange equation of $E_1(v_i)$ $= || f_p(\mathbf{x} + v_i(\mathbf{x})) - s_i ||_2^2 + ||\lambda \nabla v_i||_2^2$, an energy function of the DIR problem between the moving image $f_p(x)$ and the static image $s_i(x)$. Based on these observations, we propose a FBS algorithm to solve Eq. (1) , by alternatively performing CBCT reconstruction as in Eq. [\(6\)](#page-2-3) and DIR corresponding to Eq. [\(5\).](#page-2-4)

Specifically, FBS can be summarized as three steps (S1)– (S3) shown in Table [I.](#page-2-5) In the reconstruction step of (S1), we

TABLE I. FBS algorithm solving the problem in Eq. [\(1\).](#page-2-0)

$s_i = f_i^{(k)} - F(Pf_i^{(k)} - c(g_i)),$	(S1)
$v_i = \arg \min_{v_i} \frac{1}{2} s_i - f_p(x + v_i(x)) _2^2 + \frac{\lambda}{2} \nabla v_i _2^2$,	(S2)
$f_i^{(k+1)}(x) = f_p(x + v_i(x)),$	(S3)

have heuristically substituted P_i^T by FDK (Ref. [47\)](#page-11-7) operator *F*, which has been found to improve the convergence of the reconstruction problem. $48, 49$ $48, 49$ Since s_i is obtained purely based on reconstruction, it contains the correct anatomy location information decoded from the CBCT projections, although it is contaminated by streak artifacts due to limited number of projections and inaccurate HU values due to scattering. In the subsequent steps S2 and S3, *si* serves as a guidance to deform the prior image *fp*. Steps S1–S3 are iteratively performed, gradually leading to a fusion of both correct anatomy location/motion information from s_i and accurate intensity values from f_p .

2.A.3. Implementation

Before FBS, all 4D-CBCT projections are sorted into ten respiratory phases by performing a projection-based sorting via local principle component analysis $(LPCA).⁵⁰$ $(LPCA).⁵⁰$ $(LPCA).⁵⁰$ In principle, 4D-CT image at an arbitrary phase can serve as the p-CT *fp*. However, in order to minimize the motion blur in the p-CT, 0% (maximum inhale) or 50% (maximum exhale) phase of the 4D-CT is recommended. An average 3D-CBCT is reconstructed using the FDK algorithm and a mutual informationbased rigid registration⁵¹ is performed to align the p-CT with the average 3D-CBCT image. In (S1), to realize the *c* operator, linear intensity equalization is implemented as in Ref. [45.](#page-11-5) To address the residual intensity inconsistency between CT and CBCT images, which is indeed an open problem existing in all 2D-3D matching-based methods, 27 Demons with simultaneous intensity correction (DISC) (Ref. 52) is adopted for the DIR step in (S2) and (S3), instead of a conventional intensity-based registration algorithm, such as Demons algorithm. 53 We have used a standard mul-tiscale registration strategy in DISC.^{[52](#page-11-12)} At each scale, the iteration stops when the MVF does not change anymore. The workflow of FBS is shown in details in Table [II.](#page-3-0)

To yield a satisfactory efficiency, key components of FBS, including FDK reconstruction *F*, [54](#page-11-14) forward projection calculation $P_{1}^{55,56}$ $P_{1}^{55,56}$ $P_{1}^{55,56}$ $P_{1}^{55,56}$ and DISC,^{[52](#page-11-12)} have been implemented under CUDA programming environment on a GPU platform with an NVIDIA GTX590 card.

2.B. Evaluation

2.B.1. Experimental data

The experiments include one moving phantom case and three patient cases. For all cases, the resolution of a CBCT projection is 512×384 with pixel size of 0.784×0.784 mm². In each case, 4D-CBCT projection images are acquired. The respiratory signal is obtained by analyzing those projections

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TABLE II. FBS algorithm for hybrid 4D-CBCT reconstruction.

using a LPCA method.⁵⁰ Projections are then distributed to each breathing phase according to the obtained signal.

In the phantom case, a small ball is attached to a stick, which is then set on a QUASARTM respiratory motion platform. The platform motion follows a standard sinusoidal curve along the superior–inferior (SI) direction with an amplitude of \pm 7.5 mm. In order to mimic patient interfractional breathing variations, we have set the breathing period to 4 s for the 4D-CT simulation to obtain the p-CT and 6 s for the 1-min 4D-CBCT scan. The 4-s period is a typical value for a patient with normal breathing. The 6-s breathing period is selected to create a challenging yet realistic case to test our algorithm. In fact, for this case with a long breathing period, there are only 9–10 projections for each respiratory phase during the 1-min scan time. The purpose of this phantom study is to evaluate the accuracy of the target (moving ball) location, as we have the predefined tumor trajectory along the SI direction as the ground-truth.

Three patients (patients 1–3) were scanned with a 4D-CBCT protocol under a full fan mode with a slow gantry rotation in 5.8, 4.5, and 5.7 min over a 200 $^{\circ}$ arc.^{[15](#page-10-20)} Their breathing periods are 2.87 ± 0.65 , 2.78 ± 0.26 , and 4.45 ± 0.83 s, respectively. According to the breathing signal, we extract 121, 97, and 77 projections for each phase. These data sets are denoted as 4-min data here after. Furthermore, we select among the 4-min data 21, 20, and 13 projections per phase for the three patients, corresponding to 0.97, 0.93, and 0.96 min of scanning time over the $200°$ arc, to simulate scans of 1 min scan time (denoted as 1-min data). In these patient cases, we consider the 4D-CBCT images reconstructed by the FDK algorithm using all the 4-min data as ground truth.

2.B.2. Evaluations

For the moving ball phantom, besides the visual inspection, we determine its incremental shift in the SI direction between two adjacent phases by template matching, where the shift is indicated by the maximum cross correlation (CC) value between the shifted 4D-CBCT image at one phase and the image at the next phase. The ball location at each phase is then determined and compared with the ground truth ball trajectory set at the motion platform. Average and maximal location differences will be calculated. A region of interest (ROI) is also selected to calculate the intensity differences.

For patients' cases, in addition to directly compare the images visually, we have conducted quantitative evaluations on both geometry and image quality accuracy. For geometry accuracy evaluation, the *reference* is the 4D-CBCT images reconstructed by the FDK algorithm using the 4-min data, denoted as *FDK-4D* (4-min). *Test* images are (1) *Hybrid-4D:* 4D-CBCT images reconstructed by the proposed hybrid method using the 1-min data, (2) *FDK-4D:* 4D-CBCT images reconstructed by the FDK algorithm using the 1-min data, (3) *3D-CBCT:* CBCT image reconstructed by the FDK algorithm using all projections, and (4) *p-CT:* the image used as the input of the proposed hybrid method. A ROI containing the tumor has been selected. We have used the derivative of the ROI images to highlight the structure edges in this test in order to minimize the impacts of intensity differences between the reference and some of the test images. We have used two metrics

FIG. 1. Results of the moving phantom experiment. Display window: [−1000, 250] HU. (Upper-left) the transverse and sagittal images of (a) 3D-CBCT image, (b) FDK based 4D-CBCT image at 0% phase, (c) FDK based 4D-CBCT image at 50% phase, (d) p-CT image, (e) and (f) 4D-CBCT image at 0% and 50% phases reconstructed by the proposed hybrid algorithm, respectively. Arrows indicate the mean HU values inside the ball. In each group the two rows are transverse view and sagittal view, respectively. (g) Sagittal views at all phases of the results from the proposed hybrid algorithm. (h) Comparison of the motion trajectories derived from the results of the proposed algorithm and from ground truth.

FIG. 2. Intermediate results of the moving ball experiment (all ten phases): (a-1)–(a-3): the difference images $F(Pf_i^{(k)} - c(g_i))$ after 1, 2, and 3 iterations. Display window: [−500, 500] HU; (b-1)–(b-3): *si* image after 1, 2, and 3 iterations. Display window: [−1000, 250] HU. In each subfigure, the upper and the bottom rows show the transverse and the sagittal view, respectively.

that are robust to the intensity discrepancies to a certain degree. The first metric is the spatial shift corresponding to the maximum CC (denoted by *S*). Specifically, we shift the *test* image pixel by pixel along the SI direction until the maximum CC value between the test and the *reference* images is achieved. The smaller this shift *S* is, the better the alignment between the two images is, indicating a good reconstruction with correct anatomical structure locations. The purpose of using this quantity is to test the anatomical structure location accuracy with respect to the reference. The second one is feature similarity index $(FSIM)$, $57,58$ $57,58$ a novel human-perception based metric, with a higher value reflecting a better similarity between the *test* and *reference* images. FSIM ranges in [0, 1], with 1 being the best. To evaluate images from intensity accuracy aspect, intensity differences and SNR are calculated. In this study, a ROI has been selected, which is located at the center of the tumor. p-CT is considered as the reference im-

SNR = μ/σ is used, where μ and σ represent the mean and the standard deviation of the ROI image. **3. RESULTS 3.A. Phantom case**

> Figure [1](#page-3-1) summarizes the moving phantom results. While the average 3D-CBCT image [Fig. $1(a)$] shows significant blurring and the FDK-based 4D-CBCT images suffer from streaking artifacts [Figs. $1(b)$ and $1(c)$], the proposed hybrid method yields 4D-CBCT images with superior quality

> age, which has the highest image quality. The test images are (1) *Hybrid-4D*, (2) *FDK-4D*, (3) *3D-CBCT*, and (4) *FDK-4D* (4-min). Absolute image intensity difference averaged in the ROI between each of the test images and the reference image is calculated. For SNR calculation, no reference is needed.

FIG. 3. Experimental results for Patient 1. Display window: [−1000, 250] HU. (Upper) Transverse, coronal, and sagittal views of (a) FDK-based 3D-CBCT image, (b) FDK-based 4D-CBCT image at 50% phase using 4-min data, (c) FDK-based 4D-CBCT image at 50% phase using 1-min data, (d) p-CT image, (e) 4D-CBCT image at 50% phase reconstructed by the proposed algorithm using 1-min data. (f)–(h) Sagittal ROI 4D-CBCT images corresponding to (b), (c), and (e), respectively. ROI is indicated by the box in (a). Arrows in transverse views highlight structures with relatively large motions. Horizontal dashed lines are plotted to help comparing the anatomy locations.

[Figs. $1(e)-1(g)$]. Compared with the p-CT, the average 3D-CBCT and the FDK based 4D-CBCT images attain intensity difference of ∼–270 and ∼185 HU; the images from the proposed hybrid method yield a negligible HU difference, since it is deformed from the p-CT. Satisfactory agreements have been observed in terms of motion trajectory. Comparing the reconstructed ball trajectory with ground truth, our result achieves 0.204 mm average and 0.484 mm maximum differences [Fig. $1(h)$].

To investigate the iteration process of our algorithm in detail, we present the intermediate results of the moving phantom at all ten phases in Fig. [2.](#page-4-0) From the difference images $F(Pf_i^{(k)} - c(g_i))$ for $k = 1, 2, 3$ shown in Fig. [2\(](#page-4-0)a-1)–(a-3), we can see that the difference decreases quickly. After 3 iterations, the resulting $f_i^{(3)}$ has been deformed to a status such that its projections match well with the measurements $c(g_i)$, indicated by the small difference at this stage. From the corresponding s_i images shown in Fig. [2\(](#page-4-0)b-1)–(b-3), we observe that these images always give the correct anatomical information derived from the projection measurements. When registering $f_i^{(k)}$ with them in the DIR step, the correct MVF can be generated. It is this fact that leads to the successful estimations of the MVF and hence the reconstruction of 4D-CBCT images.

3.B. Patients case

3.B.1. Visual inspections

The results of the patient 1 are shown in Figs. [3](#page-5-0) and [4.](#page-6-0) From Fig. [3,](#page-5-0) we have the following observations: (1) FDK based average 3D-CBCT image [Fig. $3(a)$] shows minimal noise but blurred anatomies; (2) FDK based 4D-CBCT images from the 4-min data [Fig. $3(b)$] can greatly reduce motion artifacts, but the image quality is inferior to the average 3D-CBCT because of amplified streaks; (3) FDK based 4D-CBCT images from the 1-min data [Fig. $3(c)$] shows amplified noise and severe undersampling streaking artifacts; (4) Truncation exists in all the FDK-based images [Figs. $3(a)$ – $3(c)$], which will lead to inaccurate dose calculations in radiation therapy due to the missing tissue information outside the FOV; (5) With the 1-min projection data and a p-CT image [Fig. $3(d)$], the proposed method yields 4D-CBCT images with much better quality for tumor delineation, i.e., accurate anatomy location and accurate HU value [Fig. $3(e)$]. The images are also free of truncation artifacts, facilitating dose calculations. Detailed comparisons of image quality inside ROI are also shown in Figs. $3(f)-3(h)$.

Figure [4](#page-6-0) shows intermediate results during the 4D-CBCT reconstruction for the 0% phase by using the 4D-CT image of 0% phase [Fig. $3(d)$] as p-CT. Similar observations to those in Fig. [2](#page-4-0) are made. From the difference images $F(Pf_i^{(k)} - c(g_i))$ shown in Fig. [4\(a\),](#page-6-0) it can be seen that the contents of $f_i^{(k)}$ image are gradually deformed to the correct locations during the iterations, and the mismatch between $Pf_i^{(k)}$ and the measured $c(g_i)$ is reduced gradually. Figure [4\(b\)](#page-6-0) uses $||F(Pf_i^{(k)} - c(g_i))||_2$ calculated after 1, 2, 3, 10, 50, 100, and 400 iterations to quantify this fact. It can be seen that the first several iterations dramatically reduce the mismatch, and this decrease becomes gradual in later iterations. Note that this plot only illustrates the gradual reductions of the mismatches between the reconstructed results and the measurements, rather than demonstrating the algorithm convergence.

FIG. 4. Intermediate results during reconstructing the 0% phase of 4D-CBCT using the 0% phase of p-CT (Patient 1). (a) The difference images $F(Pf_i^{(k)} - c(g_i))$ after 1, 2, 3, and 50 iterations. Display window: [-700, 700] HU. (b) $||F(Pf_i^{(k)} - c(g_i))||_2$ as a function of iteration steps, with the x-axis in a log scale. (c) The s_i images after 1, 2, 3, and 50 iterations. (d) Zoomed-in ROIs to compare the s_i image after 50 iterations and the FDK-based 4D-CBCT (4-min scan). ROIs are indicated by the box. The three columns are FDK image, zoomed-in view of s_i , and the zoomed-in view of FDK image. In subfigures (a), (c), and (d), the upper and the bottom rows show the transverse and the sagittal view, respectively.

From Fig. $4(c)$, we can see that the quality of s_i images is also improved during the iteration. For example, the streaking artifacts and the evident boundary around the original FOV are eliminated greatly after 50 iterations. We have also shown in Fig. [4\(d\)](#page-6-0) a zoomed-in ROI to demonstrate that *si* provides correct anatomy information compared to that in the groundtruth, despite the intensity variations. These facts help guiding the deformations of the p-CT toward a correct location.

We have observed the same performance of our algorithm in patients 2 and 3 shown in Figs. [5](#page-6-1) and [6,](#page-7-0) respectively. In general, the 1-min FDK reconstructed images are not suitable for the tumor delineation due to the obvious streaks, especially for patient 3 where only 13 projections are available for each phase [Fig. $6(g)$] because of the long breathing period. In contrast, the proposed hybrid method is able to reconstruct images with better quality for both cases.

3.B.2. Quantitative evaluations

Figure [7](#page-7-1) summarizes the quantitative evaluation results regarding the geometry accuracy for all the three patient cases. We have the following observations: (1) *Hybrid-4D* method

FIG. 5. Experimental results of Patient 2. Subfigure arrangement is the same as that in Fig. [3](#page-5-0) except that the upper part shows 0% phase.

FIG. 6. Experimental results of Patient 3. Subfigure arrangement is the same as that in Fig. [3.](#page-5-0)

has the smallest shift *S* and the highest FSIM scores in almost all cases, indicating the high quality of our reconstruction results. This is also consistent with the visual inspections performed above. (2) *FDK-based 4D-CBCT* using the 1-min scan data also has small *S* values for all cases. It indicates that these images have correct anatomy locations, although the images are full of streaking artifacts; however, for patients 2 and 3, the FSIM scores are lower than those of our results, implying that visualization-based tasks, e.g., soft-tissue/tumor delineations are significantly affected by the artifacts. (3) *3D-CBCT* image

generally has large *S* values and low FSIM scores. There are a few cases where the 3D-CBCT has slightly better numbers in these two metrics, e.g., lower *S* values for patient 2 at 40% and 50% phases and patient 3 at 50% phase, and higher FSIM scores for patient 2 at 70% phase and patient 3 at 30% and 40% phases. Since there is only one static average 3D-CBCT for a patient, when the structure location in it happens to be close to that in one breathing phase, the corresponding *S* value may be very low. For the FSIM scores, not only does it compare structures, it also considers intensity difference. Because

FIG. 7. Comparisons of S (left) and FSIM (right) of different methods. The results of patients 1–3 are shown in (a)–(c), respectively. For S value, absence of a bar corresponds to zero.

FIG. 8. Comparisons of different methods in terms of (a) intensity difference and (b) signal-noise-ratio based on the results of patient 2.

our reference images are reconstructed by the FDK algorithm, the FSIM score is actually biased toward those images reconstructed by this algorithm due to the similar HU values. It is this fact that leads to higher FSIM scores to the average 3D-CBCT image sometimes. (4) *p-CT* always has the largest *S* value and the lowest FSIM scores. This is due to both mismatched structural information and different image intensities between the p-CT and the reference images. It implies that only using rigid registration cannot guarantee accurate softtissue alignment. Yet, using *p-CT* as an input, the proposed method is able to generate high quality outputs.

Quantitative evaluations regarding the intensity accuracy have also been conducted. An example result is shown in Fig. [8](#page-8-0) for patient 2 and similar results have been observed in other patient cases. Our *Hybrid-4D* method has the smallest intensity difference (from 14.76 to 18.21 HU for different phases, 16.70 HU on average) and the highest SNR (from 60.94 to 83.35 for different phases, 69.55 on average) in all phases. In contrast, all three FDK based reconstructions have large intensity differences [254.96, 288.68, and 286.43 HU on average for *FDK-4D*, *3D-CBCT*, and *FDK-4D* (4-min)]. As for SNR, *FDK-4D* using the 1-min scan data has the lowest ones (5.46 on average), due to the streaking artifacts and amplified noise with limited data. For *FDK-4D* (4-min), SNR is improved (13.58 on average) yet still lower than the regular *3D-CBCT* (30.65 on average). The underlying reason for such observations is that using the *Hybrid-4D* method, high image quality is inherited from the p-CT, while FDK-based reconstruction results have inferior quality due to scatter (causing large intensity differences) and the limited data induced streaking artifacts and noise (causing low SNRs).

3.B.3. Impacts of scanning time

The above experiments were focused on 1-min scan time. We would also like to understand how the scan time impacts on the results in our method. As such, we have extracted multiple subsets of projection data of patient 1 to simulate cases with different scan time. To evaluate the results, we consider the one reconstructed by the hybrid method using all the 4 min scan data as ground truth. *S* and FSIM are calculated with respect to it. Figure [9](#page-8-1) shows the results. Along the x-axis, the scanning time is increased from 0.39 min (∼7 projections per phase) to 2.90 min (∼61 projections per phase). Vertical black line indicates the case with 1-min scan time studied previously. It is interesting to see that neither metric is significantly affected by the scan time over a large range, until the time is reduced to ∼0.53 min (∼9 projections per phase). The existence of a margin from 1 min scan time to ∼0.53 min scan time indicates the robustness of our algorithm against the projection number reduction. The range of this margin is certainly dependent on the patient breathing period. For those patients with a long breathing period, e.g., patient 3, there are less number of projections per phase for a given scan time. It is therefore harder to reconstruct images for them, and the margin is expected to be smaller.

3.B.4. Impacts of deformation size

Another aspect of the algorithm robustness is the dependence on deformation size, since it is very challenging to address cases with large deformations. To test our algorithm in this regard, we have also reconstructed all the ten phases of 4D-CBCT for patients 1–3 using different p-CTs (0% or 50% phase of 4D-CT). No observable difference is found in the results compared with those shown in Figs. [3–](#page-5-0)[6.](#page-7-0) As an example, Fig. [10](#page-9-0) shows the results of patient 1, when reconstructing the 50% phase of 4D-CBCT using the 0% phase of p-CT. Due to the large deformation (∼17 mm of diaphragm motion), this case is more challenging than that shown in Fig. [4.](#page-6-0) In such a case, FBS algorithm still performs very robust. As indicated

FIG. 9. S and FSIM as the functions of scan time (patient 1, phase 0%). The vertical black lines indicate the scanning time in Fig. [3](#page-5-0) (0.97 min).

FIG. 10. Intermediate results during reconstructing the 50% phase of 4D-CBCT using the 0% phase of p-CT (Patient 1). Figure layout is the same as in Fig. [4.](#page-6-0)

by Figs. $10(a)$ and $10(b)$, $f_i^{(k)}$ image is deformed to the correct locations, and the mismatch between $Pf_i^{(k)}$ and the measurement is reduced gradually. Figure $10(c)$ shows that through two iterations, diaphragm in *si* image has already been seen in the correct location, and small structures are gradually refined in the following iterations. This guides the deformable registration to generate a good final image.

The maximum motion amplitude in our patient cases is ∼23 mm at diaphragm. While good results have been achieved using p-CT at one phase (e.g., 0% phase) to reconstruct the 4D-CBCT at another phase that is far away (e.g., 50% phase), it does not exclude the possibility that registration may fail when the respiration deformation is extremely large. We currently do not have patient data with larger deformations to test this. Yet, this concern is associated with all registration-based algorithms. To prevent such a problem, a feasible solution is to use the p-CT at a certain phase to reconstruct the 4DCBCT at the same or close phase, to ensure that the deformation is within a reasonable range.

3.C. Efficiency

The entire reconstruction process has been implemented on a GPU platform with a NVIDIA GTX590 card. Two resolution levels are used in our studies in all cases. The corresponding reconstruction time and the iteration number for each multiresolution level are listed in Table [III.](#page-9-1) It can be seen that the computational time depends on the complexity of the cases tested. Currently, for the typical clinical patient

TABLE III. Iteration number and computational time per phase.

Test cases		Moving phantom Patient 1 Patient 2 Patient 3			
Iterations	Level 0		77	72	78
	Level 1	3	53	57	53
Time (s)		2.16	83.90	81.38	67.33

cases shown here, the runtime is about 1–1.5 min for each phase.

4. DISCUSSIONS AND CONCLUSION

In this paper, we have developed a 4D-CBCT reconstruction method to facilitate fast and accurate 4D-CBCT imaging for lung and abdomen radiotherapy. The method has been thoroughly evaluated in terms of efficacy and efficiency in both phantom and real patients' cases. By using this method, we have obtained 4D-CBCT images with correct motion information and good image quality inherited from p-CT, namely, high SNR, correct intensities, and free of truncation artifacts. We have also investigated algorithm robustness with respect to scan time and deformation size. High computational efficiency (1–1.5 min per phase) has been observed due to GPU implementations.

Despite the success in those cases studied here, we would like to point out that there is no theoretical proof regarding the convergence of the algorithm. In addition, the local-minima issue associated with this and all other 2D-3D registrationbased reconstruction method may lead to geometrical inaccuracy in the results to a certain extent. There have been active researches to solve this problem in deformable image registration context, such as using multiple resolution levels. These methods may be employed here to maximally mitigate the problem. Using p-CT at the same phase to the 4D-CBCT to be reconstructed will also reduce the deformation amplitude and therefore help preventing this problem.

We would like to further comment on the reconstruction image quality. For the moving phantom case, the results show a certain amount of blurring at the superior and inferior boundaries [Fig. $1(g)$]. This can be ascribed to the blurring in the p-CT at the same area [Fig. $1(d)$]. Because of the smoothness of the MVF, this blurring originated in the p-CT will be transfer to the resulting images. Practical solution to this problem will be using high quality p-CT images, e.g., those obtained with fast CT gantry rotations. Another interesting observation is that, even though the hybrid method yields the best "similarity" with respect to the reference images, the FSIM scores are still relatively low [Fig. [7\]](#page-7-1). This is because FSIM method considers both structural and intensity similarity between the test and the reference images. Since our reference images are reconstructed by FDK algorithm, which have different HU values from the p-CT, the FSIM score underestimated the image quality to a certain extent.

While high computational efficiency has been achieved in this work, further efficiency boost is still necessary for routine clinical applications. For example, it may be possible to use less strict stopping criteria to terminate the iteration early, as it is observed in Fig. $4(b)$ that the later iterations only improve the deformation marginally. Yet, this will inevitably impact on the accuracy of deformation. Since clinically acceptable tolerance of deformation accuracy definitely depends on specific applications, the termination stopping criteria will tie to the applications as well, which will be investigated in our future studies. Another direction to accelerate the computations is to take advantages of the advanced hardware, such as multi-GPU implementation.^{[59](#page-11-19)}

ACKNOWLEDGMENTS

This work is supported in part by NIH (1R01CA154747- 01), Varian Medical Systems, Inc., and the National Natural Science Foundation of China (Nos. 30970866 and 81301940). The authors would also like to thank Yan Jiang Graves and Quentin Gautier for the assistance in CUDA implementation.

- a)Presented at the 55th Annual Meeting of the American Association of Physicists in Medicine, Indianapolis, Indiana, August 4–8, 2013.
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